

SEQUENCE LISTING

<110> Berlin-Chemie AG

<120> Pharmaceutical combinations

<130> 489.131183/01

<160> 207

<170> PatentIn version 3.5

<210> 1

<211> 119

<212> PRT

<213> Human

<400> 1

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

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

Gly Arg Ile Lys Ser Lys Thr Asp Gly Gly Thr Thr Asp Tyr Ala Ala
50 55 60

Pro Val Gln Gly Arg Phe Thr Ile Ser Arg Asp Asp Ser Lys Asn Thr
65 70 75 80

Leu Tyr Leu Gln Met Asn Ser Leu Lys Thr Glu Asp Thr Ala Val Tyr
85 90 95

Tyr Cys Thr Ile Phe Gly Val Val Ser Phe Asp Tyr Trp Gly Gln Gly
100 105 110

Thr Leu Val Thr Val Ser Ser
115

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<213> Human

<400> 2

Asp Val Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
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Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Asp Tyr
20 25 30

Leu Ser Trp Tyr Gln Gln Arg Pro Gly Lys Ala Pro Asn Leu Leu Ile
35 40 45

Tyr Ala Ala Ser Asn Leu Lys Thr Gly Val Pro Ser Arg Phe Ser Gly
50 55 60

Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Thr Leu Gln Pro
65 70 75 80

Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Arg Ser Pro Trp
85 90 95

Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg
100 105

<210> 3
<211> 357
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<400> 3

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ccagggaagg ggctggagtg ggttggccgt attaaaagca aaactgatgg tgggacaaca 180
gactacgctg caccctgca aggcagattc accatctcaa gagatgattc aaaaaacacg 240

ctgtatctgc aaatgaacag cctgaaaacc gaggacacag ccgtgtatta ctgtacgatt 300

tttggagtgg ttagctttga ctactggggc cagggaaccc tggtcaccgt ctcctca 357

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gggaaagccc ctaacctcct gatctatgct gcatccaatt taaagactgg ggtcccatca 180

aggttcagtg gcagtggatc tgggacagat ttactctca ccatcagcac tctgcaacct 240

gaagattttg caacgtacta ctgtcaacag agttacaggt ccccgaggac gttcggccaa 300

gggaccaagg tggaaatcaa acga 324

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<212> PRT

<213> Human

<400> 5

Asn Ala Trp Met Ser

1 5

<210> 6

<211> 19

<212> PRT

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Arg Ile Lys Ser Lys Thr Asp Gly Gly Thr Thr Asp Tyr Ala Ala Pro

1 5 10 15

Val Gln Gly

<210> 7
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<400> 7

Phe Gly Val Val Ser Phe Asp Tyr
1 5

<210> 8
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Arg Ala Ser Gln Ser Ile Ser Asp Tyr Leu Ser
1 5 10

<210> 9
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Ala Ala Ser Asn Leu Lys Thr
1 5

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Gln Gln Ser Tyr Arg Ser Pro Trp Thr
1 5

<210> 11
<211> 104
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<400> 11

Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Lys Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Asn Ala
20 25 30

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

Gly Arg Ile Lys Ser Lys Thr Asp Gly Gly Thr Thr Asp Tyr Ala Ala
50 55 60

Pro Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asp Ser Lys Asn Thr
65 70 75 80

Leu Tyr Leu Gln Met Asn Ser Leu Lys Thr Glu Asp Thr Ala Val Tyr
85 90 95

Tyr Cys Thr Thr Thr Thr Val Thr
100

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<400> 12

Tyr Phe Asp Tyr Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
1 5 10 15

<210> 13
<211> 93
<212> PRT
<213> Human

<400> 13

Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
1 5 10 15

Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Ser Tyr
20 25 30

Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
35 40 45

Tyr Ala Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly
50 55 60

Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
65 70 75 80

Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Ser
85 90

<210> 14
<211> 13
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<400> 14

Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg
1 5 10

<210> 15
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<212> PRT
<213> Human

<400> 15

Met Arg Thr Gly Trp Ala Thr Pro Arg Arg Pro Ala Gly Leu Leu Met
1 5 10 15

Leu Leu Phe Trp Phe Phe Asp Leu Ala Glu Pro Ser Gly Arg Ala Ala
20 25 30

Asn Asp Pro Phe Thr Ile Val His Gly Asn Thr Gly Lys Cys Ile Lys
35 40 45

Pro Val Tyr Gly Trp Ile Val Ala Asp Asp Cys Asp Glu Thr Glu Asp
50 55 60

Lys Leu Trp Lys Trp Val Ser Gln His Arg Leu Phe His Leu His Ser
65 70 75 80

Gln Lys Cys Leu Gly Leu Asp Ile Thr Lys Ser Val Asn Glu Leu Arg
85 90 95

Met Phe Ser Cys Asp Ser Ser Ala Met Leu Trp Trp Lys Cys Glu His
100 105 110

His Ser Leu Tyr Gly Ala Ala Arg Tyr Arg Leu Ala Leu Lys Asp Gly
115 120 125

His Gly Thr Ala Ile Ser Asn Ala Ser Asp Val Trp Lys Lys Gly Gly
130 135 140

Ser Glu Glu Ser Leu Cys Asp Gln Pro Tyr His Glu Ile Tyr Thr Arg
145 150 155 160

Asp Gly Asn Ser Tyr Gly Arg Pro Cys Glu Phe Pro Phe Leu Ile Asp
165 170 175

Gly Thr Trp His His Asp Cys Ile Leu Asp Glu Asp His Ser Gly Pro
180 185 190

Trp Cys Ala Thr Thr Leu Asn Tyr Glu Tyr Asp Arg Lys Trp Gly Ile
195 200 205

Cys Leu Lys Pro Glu Asn Gly Cys Glu Asp Asn Trp Glu Lys Asn Glu
210 215 220

Gln Phe Gly Ser Cys Tyr Gln Phe Asn Thr Gln Thr Ala Leu Ser Trp
225 230 235 240

Lys Glu Ala Tyr Val Ser Cys Gln Asn Gln Gly Ala Asp Leu Leu Ser

245

250

255

Ile Asn Ser Ala Ala Glu Leu Thr Tyr Leu Lys Glu Lys Glu Gly Ile
 260 265 270

Ala Lys Ile Phe Trp Ile Gly Leu Asn Gln Leu Tyr Ser Ala Arg Gly
 275 280 285

Trp Glu Trp Ser Asp His Lys Pro Leu Asn Phe Leu Asn Trp Asp Pro
 290 295 300

Asp Arg Pro Ser Ala Pro Thr Ile Gly Gly Ser Ser Cys Ala Arg Met
 305 310 315 320

Asp Ala Glu Ser Gly Leu Trp Gln Ser Phe Ser Cys Glu Ala Gln Leu
 325 330 335

Pro Tyr Val Cys Arg Lys Pro Leu Asn Asn Thr Val Glu Leu Thr Asp
 340 345 350

Val Trp Thr Tyr Ser Asp Thr Arg Cys Asp Ala Gly Trp Leu Pro Asn
 355 360 365

Asn Gly Phe Cys Tyr Leu Leu Val Asn Glu Ser Asn Ser Trp Asp Lys
 370 375 380

Ala His Ala Lys Cys Lys Ala Phe Ser Ser Asp Leu Ile Ser Ile His
 385 390 395 400

Ser Leu Ala Asp Val Glu Val Val Val Thr Lys Leu His Asn Glu Asp
 405 410 415

Ile Lys Glu Glu Val Trp Ile Gly Leu Lys Asn Ile Asn Ile Pro Thr
 420 425 430

Leu Phe Gln Trp Ser Asp Gly Thr Glu Val Thr Leu Thr Tyr Trp Asp
 435 440 445

Glu Asn Glu Pro Asn Val Pro Tyr Asn Lys Thr Pro Asn Cys Val Ser
450 455 460

Tyr Leu Gly Glu Leu Gly Gln Trp Lys Val Gln Ser Cys Glu Glu Lys
465 470 475 480

Leu Lys Tyr Val Cys Lys Arg Lys Gly Glu Lys Leu Asn Asp Ala Ser
485 490 495

Ser Asp Lys Met Cys Pro Pro Asp Glu Gly Trp Lys Arg His Gly Glu
500 505 510

Thr Cys Tyr Lys Ile Tyr Glu Asp Glu Val Pro Phe Gly Thr Asn Cys
515 520 525

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

Met Lys Lys Tyr Asp Lys Ser Leu Arg Lys Tyr Phe Trp Thr Gly Leu
545 550 555 560

Arg Asp Val Asp Ser Cys Gly Glu Tyr Asn Trp Ala Thr Val Gly Gly
565 570 575

Arg Arg Arg Ala Val Thr Phe Ser Asn Trp Asn Phe Leu Glu Pro Ala
580 585 590

Ser Pro Gly Gly Cys Val Ala Met Ser Thr Gly Lys Ser Val Gly Lys
595 600 605

Trp Glu Val Lys Asp Cys Arg Ser Phe Lys Ala Leu Ser Ile Cys Lys
610 615 620

Lys Met Ser Gly Pro Leu Gly Pro Glu Glu Ala Ser Pro Lys Pro Asp
625 630 635 640

Asp Pro Cys Pro Glu Gly Trp Gln Ser Phe Pro Ala Ser Leu Ser Cys

645

650

655

Tyr Lys Val Phe His Ala Glu Arg Ile Val Arg Lys Arg Asn Trp Glu
660 665 670

Glu Ala Glu Arg Phe Cys Gln Ala Leu Gly Ala His Leu Ser Ser Phe
675 680 685

Ser His Val Asp Glu Ile Lys Glu Phe Leu His Phe Leu Thr Asp Gln
690 695 700

Phe Ser Gly Gln His Trp Leu Trp Ile Gly Leu Asn Lys Arg Ser Pro
705 710 715 720

Asp Leu Gln Gly Ser Trp Gln Trp Ser Asp Arg Thr Pro Val Ser Thr
725 730 735

Ile Ile Met Pro Asn Glu Phe Gln Gln Asp Tyr Asp Ile Arg Asp Cys
740 745 750

Ala Ala Val Lys Val Phe His Arg Pro Trp Arg Arg Gly Trp His Phe
755 760 765

Tyr Asp Asp Arg Glu Phe Ile Tyr Leu Arg Pro Phe Ala Cys Asp Thr
770 775 780

Lys Leu Glu Trp Val Cys Gln Ile Pro Lys Gly Arg Thr Pro Lys Thr
785 790 795 800

Pro Asp Trp Tyr Asn Pro Asp Arg Ala Gly Ile His Gly Pro Pro Leu
805 810 815

Ile Ile Glu Gly Ser Glu Tyr Trp Phe Val Ala Asp Leu His Leu Asn
820 825 830

Tyr Glu Glu Ala Val Leu Tyr Cys Ala Ser Asn His Ser Phe Leu Ala
835 840 845

Thr Ile Thr Ser Phe Val Gly Leu Lys Ala Ile Lys Asn Lys Ile Ala
850 855 860

Asn Ile Ser Gly Asp Gly Gln Lys Trp Trp Ile Arg Ile Ser Glu Trp
865 870 875 880

Pro Ile Asp Asp His Phe Thr Tyr Ser Arg Tyr Pro Trp His Arg Phe
885 890 895

Pro Val Thr Phe Gly Glu Glu Cys Leu Tyr Met Ser Ala Lys Thr Trp
900 905 910

Leu Ile Asp Leu Gly Lys Pro Thr Asp Cys Ser Thr Lys Leu Pro Phe
915 920 925

Ile Cys Glu Lys Tyr Asn Val Ser Ser Leu Glu Lys Tyr Ser Pro Asp
930 935 940

Ser Ala Ala Lys Val Gln Cys Ser Glu Gln Trp Ile Pro Phe Gln Asn
945 950 955 960

Lys Cys Phe Leu Lys Ile Lys Pro Val Ser Leu Thr Phe Ser Gln Ala
965 970 975

Ser Asp Thr Cys His Ser Tyr Gly Gly Thr Leu Pro Ser Val Leu Ser
980 985 990

Gln Ile Glu Gln Asp Phe Ile Thr Ser Leu Leu Pro Asp Met Glu Ala
995 1000 1005

Thr Leu Trp Ile Gly Leu Arg Trp Thr Ala Tyr Glu Lys Ile Asn
1010 1015 1020

Lys Trp Thr Asp Asn Arg Glu Leu Thr Tyr Ser Asn Phe His Pro
1025 1030 1035

Leu Leu Val Ser Gly Arg Leu Arg Ile Pro Glu Asn Phe Phe Glu

1040

1045

1050

Glu Glu Ser Arg Tyr His Cys Ala Leu Ile Leu Asn Leu Gln Lys
1055 1060 1065

Ser Pro Phe Thr Gly Thr Trp Asn Phe Thr Ser Cys Ser Glu Arg
1070 1075 1080

His Phe Val Ser Leu Cys Gln Lys Tyr Ser Glu Val Lys Ser Arg
1085 1090 1095

Gln Thr Leu Gln Asn Ala Ser Glu Thr Val Lys Tyr Leu Asn Asn
1100 1105 1110

Leu Tyr Lys Ile Ile Pro Lys Thr Leu Thr Trp His Ser Ala Lys
1115 1120 1125

Arg Glu Cys Leu Lys Ser Asn Met Gln Leu Val Ser Ile Thr Asp
1130 1135 1140

Pro Tyr Gln Gln Ala Phe Leu Ser Val Gln Ala Leu Leu His Asn
1145 1150 1155

Ser Ser Leu Trp Ile Gly Leu Phe Ser Gln Asp Asp Glu Leu Asn
1160 1165 1170

Phe Gly Trp Ser Asp Gly Lys Arg Leu His Phe Ser Arg Trp Ala
1175 1180 1185

Glu Thr Asn Gly Gln Leu Glu Asp Cys Val Val Leu Asp Thr Asp
1190 1195 1200

Gly Phe Trp Lys Thr Val Asp Cys Asn Asp Asn Gln Pro Gly Ala
1205 1210 1215

Ile Cys Tyr Tyr Ser Gly Asn Glu Thr Glu Lys Glu Val Lys Pro
1220 1225 1230

Val Asp Ser Val Lys Cys Pro Ser Pro Val Leu Asn Thr Pro Trp
1235 1240 1245

Ile Pro Phe Gln Asn Cys Cys Tyr Asn Phe Ile Ile Thr Lys Asn
1250 1255 1260

Arg His Met Ala Thr Thr Gln Asp Glu Val His Thr Lys Cys Gln
1265 1270 1275

Lys Leu Asn Pro Lys Ser His Ile Leu Ser Ile Arg Asp Glu Lys
1280 1285 1290

Glu Asn Asn Phe Val Leu Glu Gln Leu Leu Tyr Phe Asn Tyr Met
1295 1300 1305

Ala Ser Trp Val Met Leu Gly Ile Thr Tyr Arg Asn Lys Ser Leu
1310 1315 1320

Met Trp Phe Asp Lys Thr Pro Leu Ser Tyr Thr His Trp Arg Ala
1325 1330 1335

Gly Arg Pro Thr Ile Lys Asn Glu Lys Phe Leu Ala Gly Leu Ser
1340 1345 1350

Thr Asp Gly Phe Trp Asp Ile Gln Thr Phe Lys Val Ile Glu Glu
1355 1360 1365

Ala Val Tyr Phe His Gln His Ser Ile Leu Ala Cys Lys Ile Glu
1370 1375 1380

Met Val Asp Tyr Lys Glu Glu Tyr Asn Thr Thr Leu Pro Gln Phe
1385 1390 1395

Met Pro Tyr Glu Asp Gly Ile Tyr Ser Val Ile Gln Lys Lys Val
1400 1405 1410

Thr Trp Tyr Glu Ala Leu Asn Met Cys Ser Gln Ser Gly Gly His

1415

1420

1425

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

Ile Val Lys Arg Asp Gly Phe Pro Leu Trp Val Gly Leu Ser Ser
 1445 1450 1455

His Asp Gly Ser Glu Ser Ser Phe Glu Trp Ser Asp Gly Ser Thr
 1460 1465 1470

Phe Asp Tyr Ile Pro Trp Lys Gly Gln Thr Ser Pro Gly Asn Cys
 1475 1480 1485

Val Leu Leu Asp Pro Lys Gly Thr Trp Lys His Glu Lys Cys Asn
 1490 1495 1500

Ser Val Lys Asp Gly Ala Ile Cys Tyr Lys Pro Thr Lys Ser Lys
 1505 1510 1515

Lys Leu Ser Arg Leu Thr Tyr Ser Ser Arg Cys Pro Ala Ala Lys
 1520 1525 1530

Glu Asn Gly Ser Arg Trp Ile Gln Tyr Lys Gly His Cys Tyr Lys
 1535 1540 1545

Ser Asp Gln Ala Leu His Ser Phe Ser Glu Ala Lys Lys Leu Cys
 1550 1555 1560

Ser Lys His Asp His Ser Ala Thr Ile Val Ser Ile Lys Asp Glu
 1565 1570 1575

Asp Glu Asn Lys Phe Val Ser Arg Leu Met Arg Glu Asn Asn Asn
 1580 1585 1590

Ile Thr Met Arg Val Trp Leu Gly Leu Ser Gln His Ser Val Asp
 1595 1600 1605

Gln Ser Trp Ser Trp Leu Asp Gly Ser Glu Val Thr Phe Val Lys
1610 1615 1620

Trp Glu Asn Lys Ser Lys Ser Gly Val Gly Arg Cys Ser Met Leu
1625 1630 1635

Ile Ala Ser Asn Glu Thr Trp Lys Lys Val Glu Cys Glu His Gly
1640 1645 1650

Phe Gly Arg Val Val Cys Lys Val Pro Leu Gly Pro Asp Tyr Thr
1655 1660 1665

Ala Ile Ala Ile Ile Val Ala Thr Leu Ser Ile Leu Val Leu Met
1670 1675 1680

Gly Gly Leu Ile Trp Phe Leu Phe Gln Arg His Arg Leu His Leu
1685 1690 1695

Ala Gly Phe Ser Ser Val Arg Tyr Ala Gln Gly Val Asn Glu Asp
1700 1705 1710

Glu Ile Met Leu Pro Ser Phe His Asp
1715 1720

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<211> 30
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<213> Human

<400> 16

Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Lys Pro Gly Gly
1 5 10 15

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

<210> 17
<211> 14

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<213> Human

<400> 17

Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val Gly
1 5 10

<210> 18
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<212> PRT
<213> Human

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Arg Phe Thr Ile Ser Arg Asp Asp Ser Lys Asn Thr Leu Tyr Leu Gln
1 5 10 15

Met Asn Ser Leu Lys Thr Glu Asp Thr Ala Val Tyr Tyr Cys Thr Ile
20 25 30

<210> 19
<211> 11
<212> PRT
<213> Human

<400> 19

Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
1 5 10

<210> 20
<211> 23
<212> PRT
<213> Human

<400> 20

Asp Val Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
1 5 10 15

Asp Arg Val Thr Ile Thr Cys
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<210> 21
<211> 15
<212> PRT
<213> Human

<400> 21

Trp	Tyr	Gln	Gln	Arg	Pro	Gly	Lys	Ala	Pro	Asn	Leu	Leu	Ile	Tyr
1				5					10					15

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<212> PRT
<213> Human

<400> 22

Gly	Val	Pro	Ser	Arg	Phe	Ser	Gly	Ser	Gly	Ser	Gly	Thr	Asp	Phe	Thr
1				5					10					15	

Leu	Thr	Ile	Ser	Thr	Leu	Gln	Pro	Glu	Asp	Phe	Ala	Thr	Tyr	Tyr	Cys
			20					25					30		

<210> 23
<211> 11
<212> PRT
<213> Human

<400> 23

Phe	Gly	Gln	Gly	Thr	Lys	Val	Glu	Ile	Lys	Arg
1				5					10	

<210> 24
<211> 10
<212> PRT
<213> Human

<400> 24

Trp	Glu	Val	Lys	Asp	Cys	Arg	Ser	Phe	Lys
1				5					10

<210> 25
<211> 12

<212> PRT
<213> Human

<400> 25

Pro Ala Ser Leu Ser Cys Tyr Lys Val Phe His Ala
1 5 10

<210> 26
<211> 20
<212> PRT
<213> Human

<400> 26

Pro Trp Arg Arg Gly Trp His Phe Tyr Asp Asp Arg Glu Phe Ile Tyr
1 5 10 15

Leu Arg Pro Phe
20

<210> 27
<211> 19
<212> PRT
<213> Human

<400> 27

Asp Asp His Phe Thr Tyr Ser Arg Tyr Pro Trp His Arg Phe Pro Val
1 5 10 15

Thr Phe Gly

<210> 28
<211> 12
<212> PRT
<213> Human

<400> 28

Arg Glu Leu Thr Tyr Ser Asn Phe His Pro Leu Leu
1 5 10

<210> 29
<211> 17
<212> PRT
<213> Human

<400> 29

Phe Thr Ser Cys Ser Glu Arg His Phe Val Ser Leu Cys Gln Lys Tyr
1 5 10 15

Ser

<210> 30
<211> 12
<212> PRT
<213> Human

<400> 30

Thr Val Lys Tyr Leu Asn Asn Leu Tyr Lys Ile Ile
1 5 10

<210> 31
<211> 11
<212> PRT
<213> Human

<400> 31

Glu Ala Val Tyr Phe His Gln His Ser Ile Leu
1 5 10

<210> 32
<211> 11
<212> PRT
<213> Human

<400> 32

Lys Lys Leu Ser Arg Leu Thr Tyr Ser Ser Cys
1 5 10

<210> 33
<211> 20

<212> PRT
<213> Human

<400> 33

Asn	Gly	Ser	Arg	Trp	Ile	Gln	Tyr	Lys	Gly	His	Cys	Tyr	Lys	Ser	Asp
1				5					10					15	

Gln	Ala	Leu	His
			20

<210> 34
<211> 25
<212> PRT
<213> Human

<400> 34

Ile	Ser	Glu	Trp	Pro	Ile	Asp	Asp	His	Phe	Thr	Tyr	Ser	Arg	Tyr	Pro
1				5					10					15	

Trp	His	Arg	Phe	Pro	Val	Thr	Phe	Gly
			20					25

<210> 35
<211> 20
<212> PRT
<213> Human

<400> 35

Gln	Thr	Leu	Gln	Asn	Ala	Ser	Glu	Thr	Val	Lys	Tyr	Leu	Asn	Asn	Leu
1				5					10					15	

Tyr	Lys	Ile	Ile
			20

<210> 36
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<213> Human

<400> 36

Asp Asp His Phe Thr Tyr Ser Arg Tyr Pro
1 5 10

<210> 37
<211> 15
<212> PRT
<213> Human

<400> 37

Phe Thr Ser Cys Ser Glu Arg His Phe Val Ser Leu Cys Gln Lys
1 5 10 15

<210> 38
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<212> PRT
<213> Artificial Sequence

<220>
<223> Rituximab VH sequence

<400> 38

Gln Val Gln Leu Gln Gln Pro Gly Ala Glu Leu Val Lys Pro Gly Ala
1 5 10 15

Ser Val Lys Met Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Ser Tyr
20 25 30

Asn Met His Trp Val Lys Gln Thr Pro Gly Arg Gly Leu Glu Trp Ile
35 40 45

Gly Ala Ile Tyr Pro Gly Asn Gly Asp Thr Ser Tyr Asn Gln Lys Phe
50 55 60

Lys Gly Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Ser Thr Ala Tyr
65 70 75 80

Met Gln Leu Ser Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys
85 90 95

Ala Arg Ser Thr Tyr Tyr Gly Gly Asp Trp Tyr Phe Asn Val Trp Gly

100

105

110

Ala Gly Thr Thr Val Thr Val Ser Ala Ala Ser Thr Lys Gly Pro Ser
 115 120 125

Val Phe Pro Leu Ala Pro Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala
 130 135 140

Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val
 145 150 155 160

Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala
 165 170 175

Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val
 180 185 190

Pro Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val Asn His
 195 200 205

Lys Pro Ser Asn Thr Lys Val Asp Lys Lys Ala Glu Pro Lys Ser Cys
 210 215 220

Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly
 225 230 235 240

Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met
 245 250 255

Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His
 260 265 270

Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val
 275 280 285

His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr
 290 295 300

Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly
305 310 315 320

Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile
325 330 335

Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val
340 345 350

Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val Ser
355 360 365

Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu
370 375 380

Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro
385 390 395 400

Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val
405 410 415

Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met
420 425 430

His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser
435 440 445

Pro Gly Lys
450

<210> 39
<211> 213
<212> PRT
<213> Artificial Sequence

<220>
<223> Rituximab VL sequence

<400> 39

Gln Ile Val Leu Ser Gln Ser Pro Ala Ile Leu Ser Ala Ser Pro Gly
1 5 10 15

Glu Lys Val Thr Met Thr Cys Arg Ala Ser Ser Ser Val Ser Tyr Ile
20 25 30

His Trp Phe Gln Gln Lys Pro Gly Ser Ser Pro Lys Pro Trp Ile Tyr
35 40 45

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

Gly Ser Gly Thr Ser Tyr Ser Leu Thr Ile Ser Arg Val Glu Ala Glu
65 70 75 80

Asp Ala Ala Thr Tyr Tyr Cys Gln Gln Trp Thr Ser Asn Pro Pro Thr
85 90 95

Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Arg Thr Val Ala Ala Pro
100 105 110

Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu Lys Ser Gly Thr
115 120 125

Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro Arg Glu Ala Lys
130 135 140

Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser Gly Asn Ser Gln Glu
145 150 155 160

Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr Tyr Ser Leu Ser Ser
165 170 175

Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys His Lys Val Tyr Ala
180 185 190

Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro Val Thr Lys Ser Phe

195

200

205

Asn Arg Gly Glu Cys
210

<210> 40
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<220>
<223> Rituximab VH_CDR1

<400> 40

Ser Tyr Asn Met His
1 5

<210> 41
<211> 17
<212> PRT
<213> Artificial Sequence

<220>
<223> Rituximab VH_CDR2

<400> 41

Ala Ile Tyr Pro Gly Asn Gly Asp Thr Ser Tyr Asn Gln Lys Phe Lys
1 5 10 15

Gly

<210> 42
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Rituximab VH_CDR3

<400> 42

Ser Thr Tyr Tyr Gly Gly Asp Trp Tyr Phe Asn Val

1 5 10

<210> 43
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Rituximab VL_CDR1

<400> 43

Arg Ala Ser Ser Ser Val Ser Tyr Ile His
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35 40 45

Pro Val Tyr Gly Trp Ile Val Ala Asp Asp Cys Asp Glu Thr Glu Asp
50 55 60

Lys Leu Trp Lys Trp Val Ser Gln His Arg Leu Phe His Leu His Ser
65 70 75 80

Gln Lys Cys Leu Gly Leu Asp Ile Thr Lys Ser Val Asn Glu Leu Arg
85 90 95

Met Phe Ser Cys Asp Ser Ser Ala Met Leu Trp Trp Lys Cys Glu His
100 105 110

His Ser Leu Tyr Gly Ala Ala Arg Tyr Arg Leu Ala Leu Lys Asp Gly
115 120 125

His Gly Thr Ala Ile Ser Asn Ala Ser Asp Val Trp Lys Lys Gly Gly
130 135 140

Ser Glu Glu Ser Leu Cys Asp Gln Pro Tyr His Glu Ile Tyr Thr Arg
145 150 155 160

Asp Gly Asn Ser Tyr Gly Arg Pro Cys Glu Phe Pro Phe Leu Ile Asp
165 170 175

Gly Thr Trp His His Asp Cys Ile Leu Asp Glu Asp His Ser Gly Pro
180 185 190

Trp Cys Ala Thr Thr Leu Asn Tyr Glu Tyr Asp Arg Lys Trp Gly Ile
195 200 205

Cys Leu Lys Pro Glu Asn Gly Cys Glu Asp Asn Trp Glu Lys Asn Glu
210 215 220

Gln Phe Gly Ser Cys Tyr Gln Phe Asn Thr Gln Thr Ala Leu Ser Trp
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Lys Glu Ala Tyr Val Ser Cys Gln Asn Gln Gly Ala Asp Leu Leu Ser
245 250 255

Ile Asn Ser Ala Ala Glu Leu Thr Tyr Leu Lys Glu Lys Glu Gly Ile
260 265 270

Ala Lys Ile Phe Trp Ile Gly Leu Asn Gln Leu Tyr Ser Ala Arg Gly
275 280 285

Trp Glu Trp Ser Asp His Lys Pro Leu Asn Phe Leu Asn Trp Asp Pro
290 295 300

Asp Arg Pro Ser Ala Pro Thr Ile Gly Gly Ser Ser Cys Ala Arg Met
305 310 315 320

Asp Ala Glu Ser Gly Leu Trp Gln Ser Phe Ser Cys Glu Ala Gln Leu
325 330 335

Pro Tyr Val Cys Arg Lys Pro Leu Asn Asn Thr Val Glu Leu Thr Asp
340 345 350

Val Trp Thr Tyr Ser Asp Thr Arg Cys Asp Ala Gly Trp Leu Pro Asn
355 360 365

Asn Gly Phe Cys Tyr Leu Leu Val Asn Glu Ser Asn Ser Trp Asp Lys
370 375 380

Ala His Ala Lys Cys Lys Ala Phe Ser Ser Asp Leu Ile Ser Ile His
385 390 395 400

Ser Leu Ala Asp Val Glu Val Val Val Thr Lys Leu His Asn Glu Asp
405 410 415

Ile Lys Glu Glu Val Trp Ile Gly Leu Lys Asn Ile Asn Ile Pro Thr
420 425 430

Leu Phe Gln Trp Ser Asp Gly Thr Glu Val Thr Leu Thr Tyr Trp Asp
435 440 445

Glu Asn Glu Pro Asn Val Pro Tyr Asn Lys Thr Pro Asn Cys Val Ser
450 455 460

Tyr Leu Gly Glu Leu Gly Gln Trp Lys Val Gln Ser Cys Glu Glu Lys
465 470 475 480

Leu Lys Tyr Val Cys Lys Arg Lys Gly Glu Lys Leu Asn Asp Ala Ser
485 490 495

Ser Asp Lys Met Cys Pro Pro Asp Glu Gly Trp Lys Arg His Gly Glu
500 505 510

Thr Cys Tyr Lys Ile Tyr Glu Asp Glu Val Pro Phe Gly Thr Asn Cys
515 520 525

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

Met Lys Lys Tyr Asp Lys Ser Leu Arg Lys Tyr Phe Trp Thr Gly Leu
545 550 555 560

Arg Asp Val Asp Ser Cys Gly Glu Tyr Asn Trp Ala Thr Val Gly Gly
565 570 575

Arg Arg Arg Ala Val Thr Phe Ser Asn Trp Asn Phe Leu Glu Pro Ala
580 585 590

Ser Pro Gly Gly Cys Val Ala Met Ser Thr Gly Lys Ser Val Gly Lys
595 600 605

Trp Glu Val Lys Asp Cys Arg Ser Phe Lys Ala Leu Ser Ile Cys Lys
610 615 620

Lys Met Ser Gly Pro Leu Gly Pro Glu Glu Ala Ser Pro Lys Pro Asp
625 630 635 640

Asp Pro Cys Pro Glu Gly Trp Gln Ser Phe Pro Ala Ser Leu Ser Cys
645 650 655

Tyr Lys Val Phe His Ala Glu Arg Ile Val Arg Lys Arg Asn Trp Glu
660 665 670

Glu Ala Glu Arg Phe Cys Gln Ala Leu Gly Ala His Leu Ser Ser Phe
675 680 685

Ser His Val Asp Glu Ile Lys Glu Phe Leu His Phe Leu Thr Asp Gln
690 695 700

Phe Ser Gly Gln His Trp Leu Trp Ile Gly Leu Asn Lys Arg Ser Pro
705 710 715 720

Asp Leu Gln Gly Ser Trp Gln Trp Ser Asp Arg Thr Pro Val Ser Thr
725 730 735

Ile Ile Met Pro Asn Glu Phe Gln Gln Asp Tyr Asp Ile Arg Asp Cys
740 745 750

Ala Ala Val Lys Val Phe His Arg Pro Trp Arg Arg Gly Trp His Phe
755 760 765

Tyr Asp Asp Arg Glu Phe Ile Tyr Leu Arg Pro Phe Ala Cys Asp Thr
770 775 780

Lys Leu Glu Trp Val Cys Gln Ile Pro Lys Gly Arg Thr Pro Lys Thr
785 790 795 800

Pro Asp Trp Tyr Asn Pro Asp Arg Ala Gly Ile His Gly Pro Pro Leu
805 810 815

Ile Ile Glu Gly Ser Glu Tyr Trp Phe Val Ala Asp Leu His Leu Asn
820 825 830

Tyr Glu Glu Ala Val Leu Tyr Cys Ala Ser Asn His Ser Phe Leu Ala
835 840 845

Thr Ile Thr Ser Phe Val Gly Leu Lys Ala Ile Lys Asn Lys Ile Ala
850 855 860

Asn Ile Ser Gly Asp Gly Gln Lys Trp Trp Ile Arg Ile Ser Glu Trp
865 870 875 880

Pro Ile Asp Asp His Phe Thr Tyr Ser Arg Tyr Pro Trp His Arg Phe
885 890 895

Pro Val Thr Phe Gly Glu Glu Cys Leu Tyr Met Ser Ala Lys Thr Trp
900 905 910

Leu Ile Asp Leu Gly Lys Pro Thr Asp Cys Ser Thr Lys Leu Pro Phe
915 920 925

Ile Cys Glu Lys Tyr Asn Val Ser Ser Leu Glu Lys Tyr Ser Pro Asp
930 935 940

Ser Ala Ala Lys Val Gln Cys Ser Glu Gln Trp Ile Pro Phe Gln Asn
945 950 955 960

Lys Cys Phe Leu Lys Ile Lys Pro Val Ser Leu Thr Phe Ser Gln Ala
965 970 975

Ser Asp Thr Cys His Ser Tyr Gly Gly Thr Leu Pro Ser Val Leu Ser
980 985 990

Gln Ile Glu Gln Asp Phe Ile Thr Ser Leu Leu Pro Asp Met Glu Ala
995 1000 1005

Thr Leu Trp Ile Gly Leu Arg Trp Thr Ala Tyr Glu Lys Ile Asn
1010 1015 1020

Lys Trp Thr Asp Asn Arg Glu Leu Thr Tyr Ser Asn Phe His Pro
1025 1030 1035

Leu Leu Val Ser Gly Arg Leu Arg Ile Pro Glu Asn Phe Phe Glu
1040 1045 1050

Glu Glu Ser Arg Tyr His Cys Ala Leu Ile Leu Asn Leu Gln Lys
1055 1060 1065

Ser Pro Phe Thr Gly Thr Trp Asn Phe Thr Ser Cys Ser Glu Arg
1070 1075 1080

His Phe Val Ser Leu Cys Gln Lys Tyr Ser Glu Val Lys Ser Arg
1085 1090 1095

Gln Thr Leu Gln Asn Ala Ser Glu Thr Val Lys Tyr Leu Asn Asn
1100 1105 1110

Leu Tyr Lys Ile Ile Pro Lys Thr Leu Thr Trp His Ser Ala Lys
1115 1120 1125

Arg Glu Cys Leu Lys Ser Asn Met Gln Leu Val Ser Ile Thr Asp
1130 1135 1140

Pro Tyr Gln Gln Ala Phe Leu Ser Val Gln Ala Leu Leu His Asn
1145 1150 1155

Ser Ser Leu Trp Ile Gly Leu Phe Ser Gln Asp Asp Glu Leu Asn
1160 1165 1170

Phe Gly Trp Ser Asp Gly Lys Arg Leu His Phe Ser Arg Trp Ala
1175 1180 1185

Glu Thr Asn Gly Gln Leu Glu Asp Cys Val Val Leu Asp Thr Asp
1190 1195 1200

Gly Phe Trp Lys Thr Val Asp Cys Asn Asp Asn Gln Pro Gly Ala
1205 1210 1215

Ile Cys Tyr Tyr Ser Gly Asn Glu Thr Glu Lys Glu Val Lys Pro
1220 1225 1230

Val Asp Ser Val Lys Cys Pro Ser Pro Val Leu Asn Thr Pro Trp
1235 1240 1245

Ile Pro Phe Gln Asn Cys Cys Tyr Asn Phe Ile Ile Thr Lys Asn
1250 1255 1260

Arg His Met Ala Thr Thr Gln Asp Glu Val His Thr Lys Cys Gln
1265 1270 1275

Lys Leu Asn Pro Lys Ser His Ile Leu Ser Ile Arg Asp Glu Lys
1280 1285 1290

Glu Asn Asn Phe Val Leu Glu Gln Leu Leu Tyr Phe Asn Tyr Met
1295 1300 1305

Ala Ser Trp Val Met Leu Gly Ile Thr Tyr Arg Asn Lys Ser Leu
1310 1315 1320

Met Trp Phe Asp Lys Thr Pro Leu Ser Tyr Thr His Trp Arg Ala
1325 1330 1335

Gly Arg Pro Thr Ile Lys Asn Glu Lys Phe Leu Ala Gly Leu Ser
1340 1345 1350

Thr Asp Gly Phe Trp Asp Ile Gln Thr Phe Lys Val Ile Glu Glu
1355 1360 1365

Ala Val Tyr Phe His Gln His Ser Ile Leu Ala Cys Lys Ile Glu
1370 1375 1380

Met Val Asp Tyr Lys Glu Glu Tyr Asn Thr Thr Leu Pro Gln Phe
1385 1390 1395

Met Pro Tyr Glu Asp Gly Ile Tyr Ser Val Ile Gln Lys Lys Val
1400 1405 1410

Thr Trp Tyr Glu Ala Leu Asn Met Cys Ser Gln Ser Gly Gly His
1415 1420 1425

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

Ile Val Lys Arg Asp Gly Phe Pro Leu Trp Val Gly Leu Ser Ser
1445 1450 1455

His Asp Gly Ser Glu Ser Ser Phe Glu Trp Ser Asp Gly Ser Thr
1460 1465 1470

Phe Asp Tyr Ile Pro Trp Lys Gly Gln Thr Ser Pro Gly Asn Cys
1475 1480 1485

Val Leu Leu Asp Pro Lys Gly Thr Trp Lys His Glu Lys Cys Asn
1490 1495 1500

Ser Val Lys Asp Gly Ala Ile Cys Tyr Lys Pro Thr Lys Ser Lys
1505 1510 1515

Lys Leu Ser Arg Leu Thr Tyr Ser Ser Arg Cys Pro Ala Ala Lys
1520 1525 1530

Glu Asn Gly Ser Arg Trp Ile Gln Tyr Lys Gly His Cys Tyr Lys
1535 1540 1545

Ser Asp Gln Ala Leu His Ser Phe Ser Glu Ala Lys Lys Leu Cys
1550 1555 1560

Ser Lys His Asp His Ser Ala Thr Ile Val Ser Ile Lys Asp Glu
1565 1570 1575

Asp Glu Asn Lys Phe Val Ser Arg Leu Met Arg Glu Asn Asn Asn
1580 1585 1590

Ile Thr Met Arg Val Trp Leu Gly Leu Ser Gln His Ser Val Asp
1595 1600 1605

Gln Ser Trp Ser Trp Leu Asp Gly Ser Glu Val Thr Phe Val Lys
1610 1615 1620

Trp Glu Asn Lys Ser Lys Ser Gly Val Gly Arg Cys Ser Met Leu
1625 1630 1635

Ile Ala Ser Asn Glu Thr Trp Lys Lys Val Glu Cys Glu His Gly
1640 1645 1650

Phe Gly Arg Val Val Cys Lys Val Pro Leu Gly Pro Asp Tyr Thr
1655 1660 1665

Ala Ile Ala Ile Ile Val Ala Thr Leu Ser Ile Leu Val Leu Met
1670 1675 1680

Gly Gly Leu Ile Trp Phe Leu Phe Gln Arg His Arg Leu His Leu
1685 1690 1695

Ala Gly Phe Ser Ser Val Arg Tyr Ala Gln Gly Val Asn Glu Asp
1700 1705 1710

Glu Ile Met Leu Pro Ser Phe His Asp
1715 1720

<210> 165
<211> 23
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 165

Cys Ile Lys Pro Val Tyr Gly Trp Ile Val Ala Asp Asp Cys Asp Glu
1 5 10 15

Thr Glu Asp Lys Leu Trp Lys
20

<210> 166
<211> 11

<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 166

Cys Glu His His Ser Leu Tyr Gly Ala Ala Arg
1 5 10

<210> 167
<211> 34
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 167

Asp Gly His Gly Thr Ala Ile Ser Asn Ala Ser Asp Val Trp Lys Lys
1 5 10 15

Gly Gly Ser Glu Glu Ser Leu Cys Asp Gln Pro Tyr His Glu Ile Tyr
20 25 30

Thr Arg

<210> 168
<211> 19
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 168

Lys Gly Gly Ser Glu Glu Ser Leu Cys Asp Gln Pro Tyr His Glu Ile
1 5 10 15

Tyr Thr Arg

<210> 169
<211> 18
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 169

Gly Gly Ser Glu Glu Ser Leu Cys Asp Gln Pro Tyr His Glu Ile Tyr
1 5 10 15

Thr Arg

<210> 170
<211> 15
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 170

Asp Gly His Gly Thr Ala Ile Ser Asn Ala Ser Asp Val Trp Lys
1 5 10 15

<210> 171
<211> 17
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 171

Trp Gly Ile Cys Leu Lys Pro Glu Asn Gly Cys Glu Asp Asn Trp Glu
1 5 10 15

Lys

<210> 172
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 172

Ile Phe Trp Ile Gly Leu Asn Gln Leu Tyr Ser Ala Arg
1 5 10

<210> 173
<211> 15
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 173

Leu His Asn Glu Asp Ile Lys Glu Glu Val Trp Ile Gly Leu Lys
1 5 10 15

<210> 174
<211> 15
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 174

Thr Pro Asn Cys Val Ser Tyr Leu Gly Glu Leu Gly Gln Trp Lys
1 5 10 15

<210> 175
<211> 23
<212> PRT
<213> Artificial Sequence

<220>

<223> Peptide

<400> 175

Tyr Phe Trp Thr Gly Leu Arg Asp Val Asp Ser Cys Gly Glu Tyr Asn
1 5 10 15

Trp Ala Thr Val Gly Gly Arg
20

<210> 176

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> Peptide

<400> 176

Ser Val Gly Lys Trp Glu Val Lys Asp Cys Arg
1 5 10

<210> 177

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> Peptide

<400> 177

Trp Glu Val Lys Asp Cys Arg Ser Phe Lys
1 5 10

<210> 178

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Peptide

<400> 178

Ala Leu Ser Ile Cys Lys Lys
1 5

<210> 179
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 179

Pro Ala Ser Leu Ser Cys Tyr Lys Val Phe His Ala
1 5 10

<210> 180
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 180

Arg Asn Trp Glu Glu Ala Glu Arg
1 5

<210> 181
<211> 33
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 181

Arg Ser Pro Asp Leu Gln Gly Ser Trp Gln Trp Ser Asp Arg Thr Pro
1 5 10 15

Val Ser Thr Ile Ile Met Pro Asn Glu Phe Gln Gln Asp Tyr Asp Ile
20 25 30

Arg

<210> 182
<211> 19
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 182

Thr	Pro	Val	Ser	Thr	Ile	Ile	Met	Pro	Asn	Glu	Phe	Gln	Gln	Asp	Tyr
1				5					10					15	

Asp Ile Arg

<210> 183
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 183

Pro	Trp	Arg	Arg	Gly	Trp	His	Phe	Tyr	Asp	Asp	Arg	Glu	Phe	Ile	Tyr
1				5					10					15	

Leu Arg Pro Phe
20

<210> 184
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 184

Gly Trp His Phe Tyr Asp Asp Arg
1 5

<210> 185
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 185

Ile Ser Glu Trp Pro Ile Asp Asp His Phe Thr Tyr Ser Arg
1 5 10

<210> 186
<211> 19
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 186

Asp Asp His Phe Thr Tyr Ser Arg Tyr Pro Trp His Arg Phe Pro Val
1 5 10 15

Thr Phe Gly

<210> 187
<211> 30
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 187

Phe Pro Val Thr Phe Gly Glu Glu Cys Leu Tyr Met Ser Ala Lys Thr
1 5 10 15

Trp Leu Ile Asp Leu Gly Lys Pro Thr Asp Cys Ser Thr Lys
20 25 30

<210> 188
<211> 15
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 188

Thr Trp Leu Ile Asp Leu Gly Lys Pro Thr Asp Cys Ser Thr Lys
1 5 10 15

<210> 189
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 189

Tyr Asn Val Ser Ser Leu Glu Lys
1 5

<210> 190
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 190

Val Gln Cys Ser Glu Gln Trp Ile Pro Phe Gln Asn
1 5 10

<210> 191
<211> 12
<212> PRT
<213> Artificial Sequence

<220>

<223> Peptide

<400> 191

Arg Glu Leu Thr Tyr Ser Asn Phe His Pro Leu Leu
1 5 10

<210> 192

<211> 39

<212> PRT

<213> Artificial Sequence

<220>

<223> Peptide

<400> 192

Glu Leu Thr Tyr Ser Asn Phe His Pro Leu Leu Val Ser Gly Arg Leu
1 5 10 15

Arg Ile Pro Glu Asn Phe Phe Glu Glu Glu Ser Arg Tyr His Cys Ala
20 25 30

Leu Ile Leu Asn Leu Gln Lys
35

<210> 193

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> Peptide

<400> 193

Tyr His Cys Ala Leu Ile Leu Asn Leu Gln Lys
1 5 10

<210> 194

<211> 17

<212> PRT

<213> Artificial Sequence

<220>

<223> Peptide

<400> 194

Phe	Thr	Ser	Cys	Ser	Glu	Arg	His	Phe	Val	Ser	Leu	Cys	Gln	Lys	Tyr
1				5					10					15	

Ser

<210> 195

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Peptide

<400> 195

His	Phe	Val	Ser	Leu	Cys	Gln	Lys
1				5			

<210> 196

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> Peptide

<400> 196

Gln	Thr	Leu	Gln	Asn	Ala	Ser	Glu	Thr	Val	Lys
1				5					10	

<210> 197

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

<223> Peptide

<400> 197

Thr Val Lys Tyr Leu Asn Asn Leu Tyr Lys Ile Ile
1 5 10

<210> 198

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Peptide

<400> 198

Thr Leu Thr Trp His Ser Ala Lys
1 5

<210> 199

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> Peptide

<400> 199

Asn Arg His Met Ala Thr Thr Gln Asp Glu Val His Thr Lys
1 5 10

<210> 200

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Peptide

<400> 200

Ser His Ile Leu Ser Ile Arg
1 5

<210> 201

<211> 16

<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 201

Ser	Leu	Met	Trp	Phe	Asp	Lys	Thr	Pro	Leu	Ser	Tyr	Thr	His	Trp	Ang
1				5					10					15	

<210> 202
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 202

Ser	Leu	Met	Trp	Phe	Asp	Lys
1				5		

<210> 203
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 203

Glu	Ala	Val	Tyr	Phe	His	Gln	His	Ser	Ile	Leu
1				5					10	

<210> 204
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 204

Lys Lys Leu Ser Arg Leu Thr Tyr Ser Ser
1 5 10

<210> 205
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 205

Asn Gly Ser Arg Trp Ile Gln Tyr Lys Gly His Cys Tyr Lys Ser Asp
1 5 10 15

Gln Ala Leu His
20

<210> 206
<211> 21
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 206

His Asp His Ser Ala Thr Ile Val Ser Ile Lys Asp Glu Asp Glu Asn
1 5 10 15

Lys Phe Val Ser Arg
20

<210> 207
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide

<400> 207

Val	Glu	Cys	Glu	His	Gly	Phe	Gly	Arg
1				5				