SEQUENCE LISTING

<110> Dyne Therapeutics, Inc.

<120> ANTI-TRANSFERRIN RECEPTOR (TFR) ANTIBODY AND USES THEREOF

<130> D0824.70023WO00

<140> Not Yet Assigned
<141> Concurrently Herewith

<150> US 63/055,405
<151> 2020-07-23

<150> US 62/968,252
<151> 2020-01-31

<160> 290

<170> PatentIn version 3.5

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

Tyr Met Tyr Trp Val Lys Gln Arg Pro Glu Gln Gly Leu Glu Trp Ile
35 40 45

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

Gln Asp Lys Ala Thr Val Thr Ala Asp Thr Ser Ser Asn Thr Ala Tyr
65 70 75 80

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

Thr Leu Trp Leu Arg Arg Gly Leu Asp Tyr Trp Gly Gln Gly Thr Ser
100 105 110

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


Asn Gly Tyr Thr Tyr Leu Phe Trp Phe Leu Gln Arg Pro Gly Gln Ser
        35                  40                  45


Pro Gln Leu Leu Ile Tyr Arg Met Ser Asn Leu Ala Ser Gly Val Pro
    50                  55                  60


Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Ala Phe Thr Leu Arg Ile
65                  70                  75                  80


Ser Arg Val Glu Ala Glu Asp Val Gly Val Tyr Tyr Cys Met Gln His
                85                  90                  95


Leu Glu Tyr Pro Phe Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys
            100                 105                 110
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Gln Asp Ile Ser Asn Phe
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Gln Gln Gly His Thr Leu Pro Tyr Thr
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Asp Val Gln Leu Gln Glu Ser Gly Pro Gly Leu Val Lys Pro Ser Gln
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Ser Leu Ser Leu Thr Cys Ser Val Thr Gly Tyr Ser Ile Thr Ser Gly
            20   25   30

Tyr Tyr Trp Asn Trp Ile Arg Gln Phe Pro Gly Asn Lys Leu Glu Trp
        35   40   45

Met Gly Tyr Ile Thr Phe Asp Gly Ala Asn Asn Tyr Asn Pro Ser Leu
    50   55   60

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


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


Thr Arg Ser Ser Tyr Asp Tyr Asp Val Leu Asp Tyr Trp Gly Gln Gly
            100                 105                 110


Thr Thr Leu Thr Val Ser Ser
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Asp Arg Val Thr Ile Ser Cys Arg Ala Ser Gln Asp Ile Ser Asn Phe
            20                  25                  30


Leu Asn Trp Tyr Gln Gln Arg Pro Asp Gly Thr Val Lys Leu Leu Ile
        35                  40                  45


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


Ser Gly Ser Gly Thr Asp Phe Ser Leu Thr Val Ser Asn Leu Glu Gln
65                  70                  75                  80


Glu Asp Ile Ala Thr Tyr Phe Cys Gln Gln Gly His Thr Leu Pro Tyr
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Gly Tyr Ser Phe Thr Asp Tyr Cys
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Ile Tyr Pro Gly Ser Gly Asn Thr
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Ala Arg Glu Asp Tyr Tyr Pro Tyr His Gly Met Asp Tyr
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Glu Ser Val Asp Gly Tyr Asp Asn Ser Phe
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<400> 21

Arg Ala Ser
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<210> 22
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<400> 22

Gln Gln Ser Ser Glu Asp Pro Trp Thr
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Gln Ile Gln Leu Gln Gln Ser Gly Pro Glu Leu Val Arg Pro Gly Ala
1 5 10 15

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

Cys Ile Asn Trp Val Asn Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile
35 40 45

Gly Trp Ile Tyr Pro Gly Ser Gly Asn Thr Arg Tyr Ser Glu Arg Phe
50 55 60

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

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

Ala Arg Glu Asp Tyr Tyr Pro Tyr His Gly Met Asp Tyr Trp Gly Gln
100 105 110

Gly Thr Ser Val Thr Val Ser Ser
115 120

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

Gln Arg Ala Thr Ile Ser Cys Arg Ala Ser Glu Ser Val Asp Gly Tyr

20 25 30

Asp Asn Ser Phe Met His Trp Tyr Gln Gln Lys Pro Gly Gln Pro Pro
35 40 45

Lys Leu Leu Ile Phe Arg Ala Ser Asn Leu Glu Ser Gly Ile Pro Ala
50 55 60

Arg Phe Ser Gly Ser Gly Ser Arg Thr Asp Phe Thr Leu Thr Ile Asn
65 70 75 80

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

Glu Asp Pro Trp Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys
100 105 110

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Gly Tyr Thr Phe Thr Ser Tyr Trp
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Ile Phe Pro Asn Ser Gly Arg Thr
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Ala Arg Glu Gly Asn Phe Gly Ser Leu Asp Tyr
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Ser Asn Leu Asn Tyr
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Asp Thr Ser
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Gln Gln Trp Ser Arg Asn Pro Leu Thr
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Gln Val His 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

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

Gly Asp Ile Phe Pro Asn Ser Gly Arg Thr Asn Tyr Asp Glu Lys Phe
    50   55   60

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

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

Ala Arg Glu Gly Asn Phe Gly Ser Leu Asp Tyr Trp Gly Gln Gly Thr
            100   105   110

Thr Leu Thr Val Ser Ser
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<400> 32

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

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

Asn Trp Tyr His Gln Lys Ser Gly Thr Ser Pro Lys Arg Trp Ile Tyr
35 40 45

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

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

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

Phe Gly Ala Gly Thr Arg Leu Glu Leu Lys
100 105

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Gly Phe Ser Leu Asn Thr Tyr Asp Val Gly
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Ile Trp Trp Asn Asp Asp Lys
1 5

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Thr Leu Tyr Ser Tyr Asp Gly Gly Phe Ala Tyr
1 5 10

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<400> 36

Ser Ser Val Ser Ser Ser Tyr
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Ser Thr Ser
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<400> 38

His Gln Tyr His Arg Ser Pro Tyr Thr
1 5

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

Thr Leu Ser Leu Thr Cys Ser Phe Ser Gly Phe Ser Leu Asn Thr Tyr
20 25 30

Asp Val Gly Val Gly Trp Ile Arg Gln Pro Ser Gly Lys Gly Leu Glu
35 40 45

Trp Leu Ala Asn Ile Trp Trp Asn Asp Asp Lys Tyr Tyr Asn Ser Ala

50 55 60

Leu Lys Ser Arg Leu Thr Ile Ser Lys Asp Thr Ser Asn Asn Gln Val
65 70 75 80

Phe Leu Lys Ile Ser Ser Val Asp Thr Ala Asp Thr Ala Thr Tyr Tyr
85 90 95

Cys Thr Leu Tyr Ser Tyr Asp Gly Gly Phe Ala Tyr Trp Gly Gln Gly
100 105 110

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

Glu Arg Val Thr Met Thr Cys Thr Ala Ser Ser Ser Val Ser Ser Ser
20 25 30

Tyr Leu His Trp Tyr Gln Gln Lys Pro Gly Ser Ser Pro Lys Leu Trp
35 40 45

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

Gly Ser Gly Ser Gly Thr Ser Tyr Ser Leu Thr Ile Ser Ser Met Glu
65 70 75 80

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

Tyr Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys
100 105

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Gly Tyr Ser Phe Thr Gly Tyr Asn
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Ile Asn Pro Tyr Tyr Gly Ser Thr
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Ala Arg Gly Asp Tyr Gly Tyr Asp Glu Gly Thr Trp Phe Ala Tyr
1 5 10 15

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Gln Ser Leu Leu Asn Ser Arg Thr Arg Lys Asn Tyr
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Trp Ala Ser
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Lys Gln Ser Tyr Asn Leu Pro Phe Thr
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<400> 47

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


Asn Met Asn Trp Val Lys Gln Ser His Gly Lys Ser Leu Glu Trp Ile
        35                  40                  45


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


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


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


Ala Arg Gly Asp Tyr Gly Tyr Asp Glu Gly Thr Trp Phe Ala Tyr Trp
            100                 105                 110


Gly Gln Gly Thr Leu Val Thr Val Ser Ala
        115                 120
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Asp Ile Val Met Ser Gln Ser Pro Ser Ser Leu Ala Val Ser Ala Gly
1               5                   10                  15
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Glu Lys Val Thr Met Ser Cys Lys Ser Ser Gln Ser Leu Leu Asn Ser
20 25 30

Arg Thr Arg Lys Asn Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Glu Gln
35 40 45

Ser Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val
50 55 60

Pro Asp Arg Phe Thr Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr
65 70 75 80

Ile Ser Ser Val Gln Ala Glu Asp Leu Ala Val Tyr Tyr Cys Lys Gln
85 90 95

Ser Tyr Asn Leu Pro Phe Thr Phe Gly Ser Gly Thr Lys Leu Glu Ile
100 105 110

Lys

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<400> 49

Gly Tyr Thr Phe Thr Arg His Trp
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<400> 50

Ile Tyr Pro Gly Ser Gly Arg Thr
1 5

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<400> 51

Ala Arg Asp Gly Tyr Leu Tyr Ile Asn Tyr Phe Asp Tyr
1 5 10

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<400> 52

Ser Ser Val Ser Phe
1 5

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<400> 53

Phe Gln Gly Ser Gly Tyr Pro Tyr Thr
1 5

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<400> 54

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 Arg His
20 25 30

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

Gly Asp Ile Tyr Pro Gly Ser Gly Arg Thr Asn Tyr Asn Glu Lys Phe
50 55 60

Lys Ser Thr Ala Thr Leu Thr Val Asp Thr 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 Asp Gly Tyr Leu Tyr Ile Asn Tyr Phe Asp Tyr Trp Gly Gln
100 105 110

Gly Thr Thr Leu Thr Val Ser Ser
115 120

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

Glu Lys Val Thr Met Thr Cys Ser Ala Ser Ser Ser Val Ser Phe Met
20 25 30

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

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

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

Asp Val Ala Thr Tyr Tyr Cys Phe Gln Gly Ser Gly Tyr Pro Tyr Thr
85 90 95

Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys
100 105

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<400> 56

Gly Phe Asn Ile Val Asp Asp Tyr
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Ile Tyr Pro Glu Asn Ala Asp Thr
1 5

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Thr Thr Ala Thr Gly Thr Gly Trp Phe Ala Tyr
1 5 10

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Gln Ser Leu Leu Asp Ser Asp Gly Lys Thr Tyr
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<400> 60

Leu Val Ser
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Trp Gln Gly Thr His Phe Pro Trp Thr
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<400> 62

Glu Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Val Arg Pro Gly Ala
1 5 10 15

Ser Val Lys Leu Ser Cys Thr Ala Ser Gly Phe Asn Ile Val Asp Asp
20 25 30

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

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

Gln Gly Lys Ala Thr Ile Thr Ala Asp Thr Ser Ser Asn Thr Ala Tyr
65 70 75 80

Leu Gln Leu Ser Ser Leu Thr Ser Glu Asp Thr Ala Val Tyr Tyr Cys

85 90 95

Thr Thr Ala Thr Gly Thr Gly Trp Phe Ala Tyr Trp Gly Gln Gly Thr
100 105 110

Leu Val Thr Val Ser Ala
115

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<400> 63

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

Gln Pro Ala Ser Ile Ser Cys Lys Ser Ser Gln Ser Leu Leu Asp Ser
20 25 30

Asp Gly Lys Thr Tyr Leu Asn Trp Leu Phe Gln Arg Pro Gly Gln Ser
35 40 45

Pro Lys Arg Leu Ile Tyr Leu Val Ser Lys Leu Asp Ser Gly Val Pro
50 55 60

Asp Arg Phe Thr Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Lys Ile
65 70 75 80

Ser Arg Val Glu Thr Glu Asp Leu Gly Val Tyr Tyr Cys Trp Gln Gly
85 90 95

Thr His Phe Pro Trp Thr Phe Gly Gly Gly Ala Lys Leu Glu Ile Lys
100 105 110

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Gly Tyr Thr Phe Ser Asn Tyr Trp
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Ile Leu Pro Gly Ser Gly Ser Thr
1 5

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<400> 66

Ala Arg Arg Gly Ala Tyr Gly Asn Phe His Tyr
1 5 10

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<400> 67

Ser Ser Ile Ser Ser Ser Asn
1 5

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Gly Thr Ser
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Gln Gln Trp Arg Ser Tyr Pro Tyr Thr
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Gln Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Met Lys Pro Gly Ala
1 5 10 15

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


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


Gly Glu Ile Leu Pro Gly Ser Gly Ser Thr Asn Tyr Asn Glu Asn Phe
    50                  55                  60


Lys Gly Lys Ala Thr Phe Thr Ala Asp Thr Ser Ser Asn 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 Arg Gly Ala Tyr Gly Asn Phe His Tyr Trp Gly Gln Gly Thr
            100                 105                 110


Thr Leu Thr Val Ser Ser
        115
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Glu Ile Val Leu Thr Gln Ser Pro Ala Leu Met Ala Ala Ser Pro Gly
1               5                   10                  15


Glu Lys Val Thr Ile Thr Cys Ser Val Ser Ser Ser Ile Ser Ser Ser
            20                  25                  30


Asn Leu His Trp Tyr Gln Gln Lys Ser Glu Thr Ser Pro Lys Pro Trp
        35                  40                  45
```

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

Gly Ser Gly Ser Gly Thr Ser Tyr Ser Leu Thr Ile Ser Ser Met Glu
65 70 75 80

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

Tyr Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys
100 105

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<400> 72

Gly Tyr Thr Phe Thr Asp Tyr Asn
1 5

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<400> 73

Ile Asn Pro Asn Tyr Asp Thr Thr
1 5

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<400> 74

Ala Arg Ser Gly Tyr Tyr Gly Ser Ser Tyr Tyr Trp His Phe Asp Val
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<210> 75
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<212> PRT
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<220>
<223> Synthetic

<400> 75

Gln Ser Leu Leu Tyr Ser Ser Asn Gln Lys Asn Tyr
1 5 10

<210> 76
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 76

Gln Gln Tyr Tyr Asn Tyr Pro Phe Thr
1 5

<210> 77
<211> 123
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 77

Glu Val Gln Leu Gln Gln Phe Gly Ala Glu Leu Val Lys Pro Gly Ala

1 5 10 15

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

Asn Met Ala Trp Val Lys Glu Ser His Gly Lys Ser Leu Glu Trp Ile
35 40 45

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

Lys Gly Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Ser Thr Ala His
65 70 75 80

Met Glu Leu Arg Ser Leu Thr Ser Glu Gly Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Arg Ser Gly Tyr Tyr Gly Ser Ser Tyr Tyr Trp His Phe Asp Val
100 105 110

Trp Gly Thr Gly Thr Thr Val Thr Val Ser Ser
115 120

<210> 78
<211> 113
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 78

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

Glu Lys Val Thr Met Ser Cys Lys Ser Ser Gln Ser Leu Leu Tyr Ser
20 25 30

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

Ser Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val
50 55 60

Pro Asp Arg Phe Thr Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr
65 70 75 80

Ile Ser Ser Val Lys Ala Glu Asp Leu Ala Val Tyr Tyr Cys Gln Gln
85 90 95

Tyr Tyr Asn Tyr Pro Phe Thr Phe Gly Ser Gly Thr Lys Leu Glu Ile
100 105 110

Lys

<210> 79
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 79

Gly Phe Asn Ile Lys Asp Tyr Tyr
1 5

<210> 80
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 80

Ile Asp Pro Glu Ser Gly Asp Thr

1 5

<210> 81
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 81

Tyr Gly His Asp Tyr Arg Val Asp Cys
1 5

<210> 82
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 82

Gln Ser Leu Val His Ser Asn Gly Asn Thr Tyr
1 5 10

<210> 83
<211> 3
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 83

Lys Val Ser
1

<210> 84
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 84

Ser Gln Ser Thr His Ile Pro Trp Thr
1   5

<210> 85
<211> 116
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 85

Glu Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Val Arg Ser Gly Ala
1   5   10   15

Ser Val Lys Leu Ser Cys Thr Ala Ser Gly Phe Asn Ile Lys Asp Tyr
            20   25   30

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

Gly Trp Ile Asp Pro Glu Ser Gly Asp Thr Glu Tyr Ala Pro Lys Phe
    50   55   60

Gln Gly Arg Ala Thr Met Thr Ala Asp Thr Ser Ser Asn Thr Ala Tyr
65   70   75   80

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

Tyr Gly His Asp Tyr Arg Val Asp Cys Trp Gly Gln Gly Thr Ser Val
            100   105   110

Thr Val Ser Ser

115

<210> 86
<211> 112
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 86

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

Asp Gln Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser Leu Val His Ser
20 25 30

Asn Gly Asn Thr Tyr Leu His Trp Tyr Leu Gln Lys Pro Gly Gln Ser
35 40 45

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

Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Lys Ile
65 70 75 80

Ser Arg Val Glu Ala Glu Asp Leu Gly Val Tyr Phe Cys Ser Gln Ser
85 90 95

Thr His Ile Pro Trp Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys
100 105 110

<210> 87
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 87

Ile Asn Pro Asn Tyr Asp Ser Thr
1 5

<210> 88
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 88

Gln Gln Tyr Tyr His Tyr Pro Phe Thr
1 5

<210> 89
<211> 123
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 89

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

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

Asn Met Gly Trp Val Lys Gln Ser His Gly Lys Ser Leu Glu Trp Ile
35 40 45

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

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

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

Ala Arg Ser Gly Tyr Tyr Gly Ser Ser Tyr Tyr Trp His Phe Asp Val
100 105 110

Trp Gly Thr Gly Thr Thr Val Thr Val Ser Ser
115 120

<210> 90
<211> 113
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 90

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

Glu Lys Val Thr Met Ser Cys Lys Ser Ser Gln Ser Leu Leu Tyr Ser
20 25 30

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

Ser Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val
50 55 60

Pro Asp Arg Phe Thr Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr
65 70 75 80

Ile Ser Ser Val Lys Ala Glu Asp Leu Ala Val Tyr Tyr Cys Gln Gln
85 90 95

Tyr Tyr His Tyr Pro Phe Thr Phe Gly Ser Gly Thr Lys Leu Glu Ile
100 105 110

Lys

<210> 91
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 91

Gly Phe Ser Leu Thr Asn Tyr Gly
1 5

<210> 92
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 92

Ile Trp Asn Asp Gly Ser Ala
1 5

<210> 93
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 93

Ala Arg His Glu Ser Ser Asn Pro Phe Ala Tyr
1 5 10

<210> 94

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

<220>
<223> Synthetic

<400> 94

Gln Ser Ile Gly Thr Ser
1 5

<210> 95
<211> 3
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 95

Ser Ala Ser
1

<210> 96
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 96

Gln Gln Asn Asn Arg Trp Pro Tyr Thr
1 5

<210> 97
<211> 117
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 97

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

Ser Leu Ser Ile Thr Cys Thr Val Ser Gly Phe Ser Leu Thr Asn Tyr
20 25 30

Gly Val His Trp Val Arg Gln Pro Pro Gly Lys Gly Leu Glu Trp Leu
35 40 45

Val Val Ile Trp Asn Asp Gly Ser Ala Thr Tyr Asn Ser Ala Leu Glu
50 55 60

Ser Arg Leu Ser Ile Ser Lys Asp Asn Ser Lys Ser Gln Val Phe Leu
65 70 75 80

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

Arg His Glu Ser Ser Asn Pro Phe Ala Tyr Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ala
115

<210> 98
<211> 107
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 98

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

Glu Arg Val Ser Phe Ser Cys Arg Ala Ser Gln Ser Ile Gly Thr Ser

20 25 30

Ile His Trp Tyr Gln Gln Arg Thr Asn Gly Ser Pro Arg Leu Leu Ile
35 40 45

Lys Ser Ala Ser Glu Ser Ile Ala Gly Ile Pro Ser Arg Phe Ser Gly
50 55 60

Ser Gly Ser Gly Thr Asp Phe Thr Leu Ser Ile Asn Ser Val Glu Ser
65 70 75 80

Glu Asp Ile Ala Asp Tyr Tyr Cys Gln Gln Asn Asn Arg Trp Pro Tyr
85 90 95

Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys
100 105

<210> 99
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 99

Asp Phe Asn Ile Lys Asp Asp Tyr
1 5

<210> 100
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 100

Ile Asp Pro Ala Asn Gly Asn Thr
1 5

<210> 101
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 101

Ala Leu Gly Tyr Thr Tyr
1 5

<210> 102
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 102

Gln Ser Leu Leu His Ser Tyr Gly Lys Thr Tyr
1 5 10

<210> 103
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 103

Leu Gln Thr Thr His Phe Pro Gln Thr
1 5

<210> 104
<211> 113
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 104

```
Glu Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Val Arg Pro Gly Ala
1               5                  10                  15


Ser Val Lys Leu Ser Cys Thr Ala Ser Asp Phe Asn Ile Lys Asp Asp
            20                  25                  30


Tyr Ile His Trp Val Lys Gln Arg Pro Glu Gln Gly Leu Glu Trp Ile
        35                  40                  45


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


Gln Asp Lys Ala Thr Ile Thr Ala Asp Thr Ser Ser Asn Thr Ala Tyr
65                  70                  75                  80


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


Ala Leu Gly Tyr Thr Tyr Trp Gly Gln Gly Thr Thr Leu Thr Val Ser
            100                 105                 110


Ser
```

<210> 105
<211> 112
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 105

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

```
Gln Pro Ala Ser Ile Ser Cys Lys Ser Ser Gln Ser Leu Leu His Ser
            20                  25                  30

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

Pro Lys Leu Leu Ile Tyr Leu Val Ser Lys Leu Glu Ser Gly Val Pro
    50                  55                  60

Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Lys Ile
65                  70                  75                  80

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

Thr His Phe Pro Gln Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys
            100                 105                 110
```

<210> 106
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 106

```
Gly Phe Thr Phe Ser Asp Tyr Gly
1               5
```

<210> 107
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 107

Ile Asn Ser Gly Ser Ser Thr Ile
1 5

<210> 108
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 108

Ala Arg Pro Gly Asp Tyr Asp Asn Tyr Ala Met Asp Tyr
1 5 10

<210> 109
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 109

Gln Asp Val Ser Val Ala
1 5

<210> 110
<211> 3
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 110

Trp Ala Tyr
1

<210> 111
<211> 10

<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 111

Gln Gln His Tyr Asn Thr Pro Pro Trp Thr
1 5 10

<210> 112
<211> 120
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 112

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

Ser Leu Lys Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Asp Tyr
20 25 30

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

Ala Tyr Ile Asn Ser Gly Ser Ser Thr Ile Tyr Tyr Ala Asp Thr Val
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Thr Leu Phe
65 70 75 80

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

Ala Arg Pro Gly Asp Tyr Asp Asn Tyr Ala Met Asp Tyr Trp Gly Gln
100 105 110

Gly Thr Ser Val Thr Val Ser Ser
        115   120

<210> 113
<211> 108
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 113

Asp Ile Val Met Thr Gln Ser His Lys Phe Leu Ser Thr Ser Val Gly
1   5   10   15

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

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

Tyr Trp Ala Tyr Thr Arg His Thr Gly Val Pro Asp Arg Phe Thr Gly
    50   55   60

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

Glu Asp Leu Ala Leu Tyr Tyr Cys Gln Gln His Tyr Asn Thr Pro Pro
                85   90   95

Trp Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys
            100   105

<210> 114
<211> 8
<212> PRT
<213> Artificial Sequence

<220>

<223> Synthetic

<400> 114

Ile Asp Pro Glu Asn Ala Asp Thr
1 5

<210> 115
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 115

Tyr Ala Trp Asp Tyr Ser Met Asp Tyr
1 5

<210> 116
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 116

Ser Gln Asn Thr His Val Pro Tyr Thr
1 5

<210> 117
<211> 116
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 117

Glu Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Val Arg Ser Gly Ala
1 5 10 15

Ser Val Lys Leu Ser Cys Thr Ala Ser Gly Phe Asn Ile Lys Asp Tyr
20 25 30

Tyr Ile His Trp Val Lys Gln Arg Pro Glu Gln Gly Leu Glu Trp Ile
35 40 45

Gly Trp Ile Asp Pro Glu Asn Ala Asp Thr Glu Tyr Ala Pro Lys Phe
50 55 60

Gln Gly Lys Ala Thr Met Thr Pro Asp Thr Ser Ser Asn Thr Ala Tyr
65 70 75 80

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

Tyr Ala Trp Asp Tyr Ser Met Asp Tyr Trp Gly Gln Gly Thr Ser Val
100 105 110

Thr Val Ser Ser
115

<210> 118
<211> 112
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 118

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

Asp Gln Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser Leu Val His Ser
20 25 30

Asn Gly Asn Thr Tyr Leu His Trp Tyr Leu Gln Lys Pro Gly Gln Ser
35 40 45

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

Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Ile Leu Lys Ile
65 70 75 80

Ser Arg Val Glu Ala Glu Asp Leu Gly Val Tyr Phe Cys Ser Gln Asn
85 90 95

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

<210> 119
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 119

Gly Phe Thr Phe Thr Asp Tyr Gly
1 5

<210> 120
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 120

Ile Tyr Pro Ser Ser Gly Asn Ser
1 5

<210> 121
<211> 12
<212> PRT

<213> Artificial Sequence

<220>
<223> Synthetic

<400> 121

Ala Arg Ser Thr Tyr Tyr Gly Ser Pro Ile Asp Tyr
1 5 10

<210> 122
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 122

Gln Asp Val Asp Thr Thr
1 5

<210> 123
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 123

Gln Gln Tyr Ser Thr Tyr Pro Leu Thr
1 5

<210> 124
<211> 119
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 124

Gln Val Gln Leu Gln Gln Ser Gly Thr Glu Leu Ala Arg Pro Gly Ala
1 5 10 15

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

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

Gly Glu Ile Tyr Pro Ser Ser Gly Asn Ser Tyr Tyr Asn Glu Lys Phe
50 55 60

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

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

Ala Arg Ser Thr Tyr Tyr Gly Ser Pro Ile Asp Tyr Trp Gly Gln Gly
100 105 110

Thr Thr Leu Thr Val Ser Ser
115

<210> 125
<211> 107
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 125

Asp Ile Val Met Thr Gln Ser His Lys Phe Met Ser Thr Pro Val Gly
1 5 10 15

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

```
Val Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Lys Leu Leu Ile
        35                  40                  45


Tyr Trp Ala Ser Thr Arg Gln Ile Gly Val Pro Asp Arg Phe Thr Gly
    50                  55                  60


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


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


Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys
            100                 105
```

<210> 126
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 126

```
Gly Phe Ser Leu Thr Ser Tyr Ala
1               5
```

<210> 127
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 127

```
Ile Trp Thr Gly Gly Gly Thr
1               5
```

<210> 128
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 128

Ala Arg Ile Tyr Asp Gly Tyr Tyr Arg Tyr Phe Asp Val
1 5 10

<210> 129
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 129

Gln Ser Val Ser Asn Asp
1 5

<210> 130
<211> 3
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 130

Tyr Ala Ser
1

<210> 131
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 131

Gln Gln Asp Tyr Ser Ser Pro Trp Thr
1 5

<210> 132
<211> 119
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 132

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

Ser Leu Ser Ile Thr Cys Thr Val Ser Gly Phe Ser Leu Thr Ser Tyr
20 25 30

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

Gly Leu Ile Trp Thr Gly Gly Gly Thr Asn Tyr Asn Ser Ala Leu Lys
50 55 60

Ser Arg Leu Ser Ile Ser Lys Asp Asn Ser Lys Ser Gln Val Phe Leu
65 70 75 80

Lys Met Asn Ser Leu Gln Thr Asp Asp Thr Ala Arg Tyr Tyr Cys Ala
85 90 95

Arg Ile Tyr Asp Gly Tyr Tyr Arg Tyr Phe Asp Val Trp Gly Thr Gly
100 105 110

Thr Thr Val Thr Val Ser Ser
115

<210> 133
<211> 107
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 133

Arg Ile Val Leu Thr Gln Thr Pro Lys Phe Leu Leu Val Ser Ala Gly
1 5 10 15

Asp Arg Val Thr Met Thr Cys Lys Ala Ser Gln Ser Val Ser Asn Asp
20 25 30

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

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

Ser Gly Tyr Gly Thr Asp Phe Thr Phe Thr Ile Ser Thr Val Gln Ala
65 70 75 80

Glu Asp Leu Ala Val Tyr Phe Cys Gln Gln Asp Tyr Ser Ser Pro Trp
85 90 95

Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys
100 105

<210> 134
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 134

Asn Val Leu Thr Met Pro Thr Ala Tyr

1 5

<210> 135
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 135

Gln Gln Tyr Tyr Ser Tyr Pro Tyr Thr
1 5

<210> 136
<211> 116
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 136

Glu Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Val Arg Ser Gly Ala
1 5 10 15

Ser Val Lys Leu Ser Cys Thr Ala Ser Gly Phe Asn Ile Lys Asp Tyr
20 25 30

Tyr Met His Trp Val Lys Gln Arg Pro Glu Gln Gly Leu Asp Trp Ile
35 40 45

Gly Trp Ile Asp Pro Glu Asn Gly Asp Thr Glu Tyr Ala Pro Lys Phe
50 55 60

Gln Gly Lys Ala Thr Met Thr Ala Asp Thr Ser Ser Asn Thr Ala Tyr
65 70 75 80

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

Asn Val Leu Thr Met Pro Thr Ala Tyr Trp Gly Gln Gly Thr Leu Val
100 105 110

Thr Val Ser Ala
115

<210> 137
<211> 113
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 137

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

Glu Lys Val Ile Met Ser Cys Lys Ser Ser Gln Ser Leu Leu Tyr Ser
20 25 30

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

Ser Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val
50 55 60

Pro Asp Arg Phe Thr Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr
65 70 75 80

Ile Ser Ser Val Lys Ala Glu Asp Leu Ala Val Tyr Tyr Cys Gln Gln
85 90 95

Tyr Tyr Ser Tyr Pro Tyr Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile
100 105 110

Lys

<210> 138
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 138

Gly Phe Thr Phe Ser Ser Tyr Gly
1 5

<210> 139
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 139

Ile Thr Ser Gly Gly Ser Tyr Thr
1 5

<210> 140
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 140

Ala Leu Trp Ser Leu Asp Tyr
1 5

<210> 141
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 141

Ser Ser Leu Ser Tyr
1   5

<210> 142
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 142

His Gln Arg Arg Ser Tyr Pro Trp Thr
1   5

<210> 143
<211> 114
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 143

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

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

Gly Leu Ser Trp Val Arg Gln Thr Pro Asp Lys Arg Leu Glu Trp Val
        35   40   45

Ala Thr Ile Thr Ser Gly Gly Ser Tyr Thr Tyr Tyr Pro Asp Ser Val
    50   55   60

```
Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Arg Asn Thr Leu Tyr
65                  70                  75                  80


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


Ala Leu Trp Ser Leu Asp Tyr Trp Gly Gln Gly Thr Thr Leu Thr Val
            100                 105                 110


Ser Ser
```

<210> 144
<211> 106
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 144

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


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


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


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


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


Asp Ala Ala Thr Tyr Tyr Cys His Gln Arg Arg Ser Tyr Pro Trp Thr
                85                  90                  95
```

Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys
100 105

<210> 145
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 145

Asp Asp Tyr Met Tyr
1 5

<210> 146
<211> 17
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 146

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

Asp

<210> 147
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 147

Trp Leu Arg Arg Gly Leu Asp Tyr

1 5

<210> 148
<211> 16
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 148

Arg Ser Ser Lys Ser Leu Leu His Ser Asn Gly Tyr Thr Tyr Leu Phe
1 5 10 15

<210> 149
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 149

Arg Met Ser Asn Leu Ala Ser
1 5

<210> 150
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 150

Gly Phe Asn Ile Lys Asp Asp
1 5

<210> 151
<211> 3
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 151

Glu Asn Gly
1

<210> 152
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 152

Leu Arg Arg Gly Leu Asp
1   5

<210> 153
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 153

Ser Lys Ser Leu Leu His Ser Asn Gly Tyr Thr Tyr
1   5   10

<210> 154
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 154

His Leu Glu Tyr Pro Phe

1 5

<210> 155
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 155

Ser Gly Tyr Tyr Trp Asn
1 5

<210> 156
<211> 16
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 156

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

<210> 157
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 157

Ser Ser Tyr Asp Tyr Asp Val Leu Asp Tyr
1 5 10

<210> 158
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 158

Arg Ala Ser Gln Asp Ile Ser Asn Phe Leu Asn
1   5   10

<210> 159
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 159

Tyr Thr Ser Arg Leu His Ser
1   5

<210> 160
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 160

Gly Tyr Ser Ile Thr Ser Gly Tyr
1   5

<210> 161
<211> 3
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 161

Phe Asp Gly

1

<210> 162
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 162

Ser Tyr Asp Tyr Asp Val Leu Asp
1 5

<210> 163
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 163

Ser Gln Asp Ile Ser Asn Phe
1 5

<210> 164
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 164

Gly His Thr Leu Pro Tyr
1 5

<210> 165
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 165

Asp Tyr Cys Ile Asn
1 5

<210> 166
<211> 17
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 166

Trp Ile Tyr Pro Gly Ser Gly Asn Thr Arg Tyr Ser Glu Arg Phe Lys
1 5 10 15

Gly

<210> 167
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 167

Glu Asp Tyr Tyr Pro Tyr His Gly Met Asp Tyr
1 5 10

<210> 168
<211> 15
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 168

Arg Ala Ser Glu Ser Val Asp Gly Tyr Asp Asn Ser Phe Met His
1 5 10 15

<210> 169
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 169

Arg Ala Ser Asn Leu Glu Ser
1 5

<210> 170
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 170

Gly Tyr Ser Phe Thr Asp Tyr
1 5

<210> 171
<211> 3
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 171

Gly Ser Gly
1

<210> 172
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 172

Asp Tyr Tyr Pro Tyr His Gly Met Asp
1 5

<210> 173
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 173

Ser Glu Ser Val Asp Gly Tyr Asp Asn Ser Phe
1 5 10

<210> 174
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 174

Ser Ser Glu Asp Pro Trp
1 5

<210> 175
<211> 330
<212> PRT
<213> Homo sapiens

<400> 175

```
Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys
1               5                   10                  15

Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr
            20                  25                  30

Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser
        35                  40                  45

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

Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr
65                  70                  75                  80

Tyr Ile Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys
                85                  90                  95

Lys Val Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys
            100                 105                 110

Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro
        115                 120                 125

Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys
    130                 135                 140

Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp
145                 150                 155                 160

Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu
                165                 170                 175

Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu
            180                 185                 190

His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn
```

195 200 205

Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly
210 215 220

Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu
225 230 235 240

Leu Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr
245 250 255

Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn
260 265 270

Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe
275 280 285

Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn
290 295 300

Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr
305 310 315 320

Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys
325 330

<210> 176
<211> 330
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 176

Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys
1 5 10 15

```
Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr
            20                  25                  30

Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser
        35                  40                  45

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

Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr
65                  70                  75                  80

Tyr Ile Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys
                85                  90                  95

Lys Val Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys
            100                 105                 110

Pro Ala Pro Glu Ala Ala Gly Gly Pro Ser Val Phe Leu Phe Pro Pro
        115                 120                 125

Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys
    130                 135                 140

Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp
145                 150                 155                 160

Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu
                165                 170                 175

Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu
            180                 185                 190

His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn
        195                 200                 205

Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly
```

210 215 220

Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu
225 230 235 240

Leu Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr
245 250 255

Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn
260 265 270

Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe
275 280 285

Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn
290 295 300

Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr
305 310 315 320

Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys
325 330

<210> 177
<211> 107
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 177

Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu
1 5 10 15

Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe
20 25 30

Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln
35 40 45

Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser
50 55 60

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

Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser
85 90 95

Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys
100 105

<210> 178
<211> 447
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 178

Glu Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Val Arg Pro Gly Ala
1 5 10 15

Ser Val Lys Leu Ser Cys Thr Ala Ser Gly Phe Asn Ile Lys Asp Asp
20 25 30

Tyr Met Tyr Trp Val Lys Gln Arg Pro Glu Gln Gly Leu Glu Trp Ile
35 40 45

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

Gln Asp Lys Ala Thr Val Thr Ala Asp Thr Ser Ser Asn Thr Ala Tyr
65 70 75 80

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

Thr Leu Trp Leu Arg Arg Gly Leu Asp Tyr Trp Gly Gln Gly Thr Ser
            100                 105                 110

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

Ala Pro Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys
    130                 135                 140

Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser
145                 150                 155                 160

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

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

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

Thr Lys Val Asp Lys Lys Val Glu Pro Lys Ser Cys Asp Lys Thr His
    210                 215                 220

Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val
225                 230                 235                 240

Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr
                245                 250                 255

Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro Glu
            260                 265                 270
```

```
Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys
        275                 280                 285

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

Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys
305                 310                 315                 320

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

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

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

Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn
    370                 375                 380

Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser
385                 390                 395                 400

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

Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala Leu
            420                 425                 430

His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys
        435                 440                 445
```

<210> 179
<211> 219
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 179

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


Glu Ser Val Ser Ile Ser Cys Arg Ser Ser Lys Ser Leu Leu His Ser
            20                  25                  30


Asn Gly Tyr Thr Tyr Leu Phe Trp Phe Leu Gln Arg Pro Gly Gln Ser
        35                  40                  45


Pro Gln Leu Leu Ile Tyr Arg Met Ser Asn Leu Ala Ser Gly Val Pro
    50                  55                  60


Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Ala Phe Thr Leu Arg Ile
65                  70                  75                  80


Ser Arg Val Glu Ala Glu Asp Val Gly Val Tyr Tyr Cys Met Gln His
                85                  90                  95


Leu Glu Tyr Pro Phe Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys
            100                 105                 110


Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu
        115                 120                 125


Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe
    130                 135                 140


Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln
145                 150                 155                 160


Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser
                165                 170                 175
```

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

Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser
195 200 205

Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys
210 215

<210> 180
<211> 449
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 180

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

Ser Leu Ser Leu Thr Cys Ser Val Thr Gly Tyr Ser Ile Thr Ser Gly
20 25 30

Tyr Tyr Trp Asn Trp Ile Arg Gln Phe Pro Gly Asn Lys Leu Glu Trp
35 40 45

Met Gly Tyr Ile Thr Phe Asp Gly Ala Asn Asn Tyr Asn Pro Ser Leu
50 55 60

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

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

Thr Arg Ser Ser Tyr Asp Tyr Asp Val Leu Asp Tyr Trp Gly Gln Gly
100 105 110

```
Thr Thr Leu Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe
        115                 120                 125

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

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

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

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

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

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

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

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

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

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

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

```
Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu
305                 310                 315                 320

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

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

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

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

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

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

Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu
            420                 425                 430

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

Lys
```

<210> 181
<211> 214
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 181

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

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

Leu Asn Trp Tyr Gln Gln Arg Pro Asp Gly Thr Val Lys Leu Leu Ile
        35                  40                  45

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

Ser Gly Ser Gly Thr Asp Phe Ser Leu Thr Val Ser Asn Leu Glu Gln
65                  70                  75                  80

Glu Asp Ile Ala Thr Tyr Phe Cys Gln Gln Gly His Thr Leu Pro Tyr
                85                  90                  95

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

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

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

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

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

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

Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro Val Thr Lys Ser
195 200 205

Phe Asn Arg Gly Glu Cys
210

<210> 182
<211> 450
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 182

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

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

Cys Ile Asn Trp Val Asn Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile
35 40 45

Gly Trp Ile Tyr Pro Gly Ser Gly Asn Thr Arg Tyr Ser Glu Arg Phe
50 55 60

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

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

Ala Arg Glu Asp Tyr Tyr Pro Tyr His Gly Met Asp Tyr Trp Gly Gln
100 105 110

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

Gly Lys
450

<210> 183
<211> 218
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 183

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

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

Asp Asn Ser Phe Met His Trp Tyr Gln Gln Lys Pro Gly Gln Pro Pro
        35                  40                  45

Lys Leu Leu Ile Phe Arg Ala Ser Asn Leu Glu Ser Gly Ile Pro Ala
    50                  55                  60

Arg Phe Ser Gly Ser Gly Ser Arg Thr Asp Phe Thr Leu Thr Ile Asn
65                  70                  75                  80

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

Glu Asp Pro Trp Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Arg
            100                 105                 110

Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln
        115                 120                 125

Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr
    130                 135                 140

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

Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr
                165                 170                 175

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

His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro
        195                 200                 205
```

Val Thr Lys Ser Phe Asn Arg Gly Glu Cys
210 215

<210> 184
<211> 108
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 184

Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys
1 5 10 15

Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr
20 25 30

Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser
35 40 45

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

Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr
65 70 75 80

Tyr Ile Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys
85 90 95

Lys Val Glu Pro Lys Ser Cys Asp Lys Thr His Thr
100 105

<210> 185
<211> 225
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 185

```
Glu Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Val Arg Pro Gly Ala
1               5                  10                  15

Ser Val Lys Leu Ser Cys Thr Ala Ser Gly Phe Asn Ile Lys Asp Asp
            20                  25                  30

Tyr Met Tyr Trp Val Lys Gln Arg Pro Glu Gln Gly Leu Glu Trp Ile
        35                  40                  45

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

Gln Asp Lys Ala Thr Val Thr Ala Asp Thr Ser Ser Asn Thr Ala Tyr
65                  70                  75                  80

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

Thr Leu Trp Leu Arg Arg Gly Leu Asp Tyr Trp Gly Gln Gly Thr Ser
            100                 105                 110

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

Ala Pro Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys
    130                 135                 140

Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser
145                 150                 155                 160

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

Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser
            180                 185                 190
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Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val Asn His Lys Pro Ser Asn
195 200 205

Thr Lys Val Asp Lys Lys Val Glu Pro Lys Ser Cys Asp Lys Thr His
210 215 220

Thr
225

<210> 186
<211> 227
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 186

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

Ser Leu Ser Leu Thr Cys Ser Val Thr Gly Tyr Ser Ile Thr Ser Gly
20 25 30

Tyr Tyr Trp Asn Trp Ile Arg Gln Phe Pro Gly Asn Lys Leu Glu Trp
35 40 45

Met Gly Tyr Ile Thr Phe Asp Gly Ala Asn Asn Tyr Asn Pro Ser Leu
50 55 60

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

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

Thr Arg Ser Ser Tyr Asp Tyr Asp Val Leu Asp Tyr Trp Gly Gln Gly

100 105 110

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

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

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

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

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

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

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

Thr His Thr
225

<210> 187
<211> 228
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 187

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

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

Cys Ile Asn Trp Val Asn Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile
        35                  40                  45

Gly Trp Ile Tyr Pro Gly Ser Gly Asn Thr Arg Tyr Ser Glu Arg Phe
    50                  55                  60

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

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

Ala Arg Glu Asp Tyr Tyr Pro Tyr His Gly Met Asp Tyr Trp Gly Gln
            100                 105                 110

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

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

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

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

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

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

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

Lys Thr His Thr
225

<210> 188
<211> 351
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 188
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tcctgcacag cttctggctt taacattaaa gacgactata tgtactgggt gaagcagagg 120

cctgagcagg gcctggagtg gattggatgg attgatcctg agaatggtga tactgaatat 180

gcctcgaagt tccaggacaa ggccactgta acagcagaca catcctccaa cacagcctac 240

ctgcagctca gcagcctgac atctgaggac actgccgtct attactgtac tttatggtta 300

cgccgagggt tggactactg gggtcaagga acctcagtca ccgtctcctc a 351

<210> 189
<211> 336
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 189
gatattgtga tgactcaggc tgcaccctct gtacctgtca ctcctggaga gtcagtatcc 60

atctcctgca ggtctagtaa gagtctcctg catagtaatg gctacactta cttgttttgg 120

ttcctgcaga ggccaggcca gtctcctcag ctcctgatat atcggatgtc caaccttgcc 180

tcaggagtcc cagacaggtt cagtggcagt gggtcaggaa ctgctttcac actgagaatc 240

agtagagtgg aggctgagga tgtgggtgtt tattactgta tgcaacatct agaatatccg 300

ttcacgttcg gaggggggac caagctggaa ataaaa 336

<210> 190
<211> 357
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 190
gatgtacagc ttcaggagtc aggacctggc ctcgtgaaac cttctcagtc tctgtctctc 60

acctgctctg tcactggcta ctccatcacc agtggttatt actggaactg gatccggcag 120

tttccaggaa acaaactgga atggatgggc tacataacct tcgatggtgc caataactac 180

aacccatctc tcaaaaatcg catctccatc actcgtgaca catctaagaa ccagtttttc 240

ctgaagttga cttctgtgac tactgaggac acagccacat attactgtac aagatcctcc 300

tatgattacg acgtccttga ctactggggc caaggcacca ctctcacagt ctcctca 357

<210> 191
<211> 321
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 191
gatatccaga tgacacagac tacatcctcc ctgtctgcct ctctgggaga cagagtcacc 60

atcagttgca gggcaagtca ggacattagc aatttttaa actggtatca gcagagacca 120

gatggaactg ttaaactcct gatctactac acatcaagat tacactcagg agtcccatca 180

aggttcagtg gcagtgggtc tgggacagat ttttctctca ccgttagcaa cctggagcaa 240

gaagatattg ccacttactt ttgccaacag ggtcatacgc ttccgtacac gttcggaggg 300

gggaccaagc tggaaataaa a 321

<210> 192
<211> 360
<212> DNA

<213> Artificial Sequence

<220>
<223> Synthetic

<400> 192
cagatccagc tgcagcagtc tggacctgag ttggtgaggc ctggggcttc agtgaagata 60

tcctgcaagg cttctggcta ctccttcact gactactgta taaattgggt gaatcagagg 120

cctggacagg gacttgagtg gattggatgg atttatcctg gaagcggtaa tactaggtac 180

agtgagaggt tcaagggcaa ggccacattg actgtagaca catcttccaa cacagcctac 240

atgcagctca gcagcctgac ctctgaggac tctgcggtct atttctgtgc aagagaggat 300

tactatccgt accatggtat ggactactgg ggtcaaggca cctcagtcac cgtctcctca 360

<210> 193
<211> 333
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 193
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atatcctgca gagccagtga aagtgttgat ggttatgaca atagttttat gcactggtac 120

cagcagaaac caggacagcc acccaaactc ctcatctttc gtgcatccaa cctagaatct 180

gggatccctg ccaggttcag tggcagtggg tctaggacag acttcaccct caccattaat 240

cctgtggagg ctgctgatgt tgcaacctat tactgtcagc aaagtagtga ggatccgtgg 300

acgttcggtg gaggcaccaa gctggaaatc aaa 333

<210> 194
<211> 354
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 194
caggtccacc tgcagcagcc tggggctgag cttgtgaagc ctggggcttc agtgaagatg 60

tcctgcaagg cttctggcta caccttcacc agctactgga taacctgggt gaagcagagg 120

cctggacaag gccttgagtg gattggagat atttttccta atagtggtag aactaactac 180

gatgagaagt tcaagagcaa ggccacactg actgtagaca catcctccag caccgcctac 240

atgcagctca gcagcctgac atctgaggac tctgcggtct atttctgtgc aagggaggga 300

aacttcggta gtcttgacta ctggggccaa ggcaccactc tcacagtctc ctca 354

<210> 195
<211> 318
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 195
caaattgttc tcacccagtc tccagcaatc atgtctgcat ctccagggga gaaggtcacc 60

atgacctgca gtgccaactc aaatttaaat tacatgaact ggtaccacca gaagtcaggc 120

acctccccca aaagatggat ttatgacaca tccaaactgg cttctggagt ccctgctcgc 180

ttcagtgcca gtgggtctgg gacctcttac tctctcacaa tcagcagcat ggaggctgaa 240

gatgctgcca cttattactg ccagcagtgg agtagaaacc ctctcacgtt cggtgctggg 300

accaggctgg agctgaaa 318

<210> 196
<211> 357
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 196
caggtcactc tgaaagagtc tggccctggg atgttgcagc cctcccagac cctcagtctg 60

acttgttctt tctctgggtt ttcactgaac acttatgatg tgggtgtagg ttggattcgt 120

cagccttcag ggaagggtct ggagtggctg gccaacattt ggtggaatga tgataagtac 180

tataactcag ccctgaagag ccggctcaca atctccaagg atacctccaa caaccaggtc 240

ttcctcaaga tctccagtgt ggacactgca gatactgcca catactactg taccctatat 300

agttacgacg gagggtttgc ttactggggc caagggactc tggtcactgt ctctgca 357

<210> 197
<211> 324
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 197
caaattgttc tcacccagtc tccagcaatc atgtctgcat ctctaggggа acgggtcacc 60

atgacctgca ctgccagctc aagtgtaagt tccagttact tgcactggta ccagcagaag 120

ccaggatcct cccccaaact ctggatttat agcacatcca acctggcttc tggagtccca 180

gctcgcttca gtggcagtgg gtctgggacc tcttactctc tcacaatcag cagcatggag 240

gctgaagatg ctgccactta ttactgccac cagtatcatc gttccccgta cacgttcgga 300

gggggggacca agctggaaat aaaa 324

<210> 198
<211> 366
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 198
gagatccaga tgaagcagtc tggagctgag ctggtgaagc ctggggcttc agtgaagata 60

tcctgcaagg cttctggtta ctcattcact ggctacaata tgaactgggt gaagcagagc 120

catggaaaga gccttgagtg gattggaaat attaatcctt actatggtag tactggctac 180

aatcagaaat tcaagggcaa ggccacattg actgtagaca aatcttccag cacagcctac 240

atgcagctca acagcctgac atctgaggac tctgcagtct attactgtgc aagaggggac 300

tatggttacg acgaggggac ctggtttgct tactggggcc aagggactct ggtcactgtc 360

tctgca 366

<210> 199
<211> 339
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 199
gacattgtga tgtcacagtc tccatcctcc ctggctgtgt cagcaggaga gaaggtcact 60

atgagctgca aatccagtca gagtctgctc aacagtagaa cccgaaagaa ctacttggct 120

tggtaccagc agaaaccaga gcagtctcct aaactgctga tctactgggc atccactagg 180

gaatctgggg tccctgatcg cttcacaggc agtggatctg ggacagattt cactctcacc 240

atcagcagtg tgcaggctga agacctggca gtttattact gcaagcaatc ttataatctt 300

ccattcacgt tcggctcggg gacaaaattg gaaataaaa 339

<210> 200
<211> 360
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 200
caggtccaac tgcagcagcc tggggctgag cttgtgaagc ctggggcttc agtgaagatg 60

tcctgcaagg cttctggcta caccttcacc aggcactgga taacctgggt gaagcagagg 120

cctggacaag gccttgagtg gattggagat atttatcctg gtagtggtag aactaactac 180

aatgagaagt tcaagagcac ggccacactg actgtagaca catcctccag cacagcctac 240

atgcagctca gcagcctgac atctgaggac tctgcggtct attactgtgc acgagacggt 300

tacttatata taaactactt tgactactgg ggccaaggaa ccactctcac agtctcctca 360

<210> 201
<211> 318
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 201
gaaaatgttc tcacccagtc tccagcaatc atgtctgcat ctccagggga aaaggtcacc 60

atgacctgca gtgccagctc aagtgtaagt ttcatgcact ggttccagca gaagtcaagc 120

acctccccca aactctggat ttatgacaca tccaaactgg cttctggagt cccaggtcgc 180

ttcagtggca gtgggtctgg aagctcttac tctctcacga tcagcagcat ggcggctgaa 240

gatgttgcca cttattactg ttttcagggg agtgggtacc cgtacacgtt cggaggggg 300

accaagctgg aaataaaa 318

<210> 202
<211> 354
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 202
gaggttcagc tgcagcagtc tggggctgag cttgtgaggc caggggcctc agtcaagttg 60

tcctgcacag cttctggctt taacattgta gacgactata tgcactgggt gaagcagagg 120

cctgaacagg gcctggagtg gattggatgg atttatcctg agaatgctga tactgaatat 180

gcctcgaagt tccagggcaa ggccactata acagcagaca catcctccaa cacagcctac 240

ctgcagctca gcagcctaac atctgaggac actgccgtct attactgtac taccgccact 300

gggacgggct ggtttgctta ctggggccaa gggactctgg tcactgtctc tgca 354

<210> 203
<211> 336
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 203
gatgttgtga tgacccagac tccactcact ttgtcggtta ccattggaca accagcctcc 60

atctcttgca agtcaagtca gagcctctta gatagtgatg gaaagacata tttgaattgg 120

ttgtttcaga ggccaggcca gtctccaaag cgcctaatct atctggtgtc taaactggac 180

tctggagtcc ctgacaggtt cactggcagt ggatcaggga cagatttcac actgaaaatc 240

agcagagtgg agactgagga tttgggagtt tattattgct ggcaaggtac acattttccg 300

tggacgttcg gtggaggcgc caagctggaa atcaaa 336

<210> 204
<211> 354
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 204
caggttcagc tgcagcagtc tggagctgag ctgatgaagc ctggggcctc agtgaagata 60

tcctgcaagg ctactggcta cacattcagt aactactgga tagagtgggt aaagcagagg 120

cctggacatg gccttgagtg gattggagag attttacctg gaagtggtag tactaactat 180

aatgagaact tcaagggcaa ggccacattc actgcagata catcctccaa cacagcctac 240

atgcaactca gcagcctgac atctgaggac tctgccgtct attactgtgc acgaaggggg 300

gcctatggta actttcacta ctggggccaa ggcaccactc tcacagtctc ctca 354

<210> 205
<211> 324
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 205

gaaattgtgc tcacccagtc tccagcactc atggctgcat ctccagggga gaaggtcacc 60

atcacctgca gtgtcagctc aagtataagt tccagcaact tgcactggta ccagcagaag 120

tcagaaacct cccccaaacc ctggatttat ggcacatcca acctggcttc tggagtccct 180

gttcgcttca gtggcagtgg atctgggacc tcttattctc tcacaatcag cagcatggag 240

gctgaagatg ctgccactta ttactgtcaa cagtggcgta gttacccgta cacgttcgga 300

gggggggacca agctggaaat aaaa 324

<210> 206
<211> 369
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 206
gaggtccagc tgcaacagtt tggagctgag ctggtgaagc ctggggcttc agtgaagata 60

tcctgcaagg cttctggcta cacattcact gactacaaca tggcctgggt gaaggagagc 120

catggaaaga gccttgagtg gattggagat attaatccta actatgatac tactagctac 180

aaccagaagt tcaagggaaa ggccacattg actgtagaca agtcctccag cacagcccac 240

atggagctcc gcagcctgac atctgagggc actgcagtct attactgtgc aagatcgggt 300

tactacggta gtagctacta ctggcacttc gatgtctggg gcacagggac cacggtcacc 360

gtctcctca 369

<210> 207
<211> 339
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 207
gacattgtga tgtcacagtc tccatcctcc ctagctgtgt cagttggaga gaaggttact 60

atgagctgca agtccagtca gagcctttta tatagtagca atcaaaagaa ctacttggcc 120

tggtaccagc agaaaccagg gcagtctcct aaactgctga tttactgggc atccactagg 180

gaatctgggg tccctgatcg cttcacaggc agtggatctg ggacagattt cactctcacc 240

atcagcagtg tgaaggctga agacctggca gtttattact gtcagcaata ttataactat 300

ccattcacgt tcggctcggg gacaaagttg gaaataaaa 339

<210> 208
<211> 348
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 208
gaggttcagc tgcagcagtc tggggcagag ctagtgaggt caggggcctc agtcaagttg 60

tcctgcacag cttctggctt caacattaaa gactactata tgcactgggt gaaacagagg 120

cctgaacagg gcctggagtg gattggatgg attgatcctg agagtggtga tactgaatat 180

gccccgaagt tccagggcag ggccactatg actgcagaca catcctccaa cacagcctac 240

atgcagctca gcagcctgac atctgaggac actgccgtct attactgtta tggacatgat 300

tatagggtgg actgctgggg tcaaggaacc tcagtcactg tctcctca 348

<210> 209
<211> 336
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 209
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atctcttgca gatctagtca gagccttgta cacagtaatg gaaacaccta tttacattgg 120

tacctgcaga agccaggcca gtctccaaag ctcctgatct acaaagtttc caaccgattt 180

tctggggtcc cagacaggtt cagtggcagt ggatcaggga cagatttcac actcaagatc 240

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tggacgttcg gtggaggcac caagctggaa atcaaa 336

<210> 210
<211> 369
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 210
gaggtccagt tgcaacagtt tggagctgag ctggtgaagc ctggggcttc agtgaagata 60

tcctgcaagg cttctggcta cacattcact gactacaaca tgggctgggt gaagcagagc 120

catggaaaga gccttgagtg gattggagat attaatccta actatgatag tactagctac 180

acccagaagt tcaagggaaa ggccacattg actgtagaca agtcctccag cacagcctac 240

atggaactcc gcagcctgac atctgaggac actgcagtct attactgtgc aagatcgggt 300

tactacggta gtagctacta ctggcacttc gatgtctggg gcacagggac cacggtcacc 360

gtctcctca 369

<210> 211
<211> 339
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 211
gacattgtga tgtcacagtc tccatcctcc ctagctgtgt cagttggaga gaaggttact 60

atgagctgca agtccagtca gagcctttta tatagtagca atcaaaagaa ctacttggcc 120

tggtaccagc agaaaccagg gcagtctcct aaactgctga tttactgggc atccactagg 180

gaatctgggg tccctgatcg cttcacaggc agtggatctg ggacagattt cactctcacc 240

atcagcagtg tgaaggctga agacctggca gtttattact gtcaacaata ttatcactat 300

ccattcacgt tcggctcggg gacaaagttg gaaataaaa 339

<210> 212
<211> 351
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 212
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acatgcaccg tctcagggtt ctcattaacc aactatggtg ttcactgggt tcgccagcct 120

ccagggaagg gtctggaatg gctggtcgtg atatggaatg atggaagcgc aacctataat 180

tcagctctcg aatccagact gagcatcagc aaggacaact ccaagagcca agttttctta 240

aaaatgaaca gtctccaaac tgatgacaca gccatgtact actgtgccag acatgaatct 300

agtaacccgt ttgcttactg gggccaaggg actctggtca ctgtctctgc a 351

<210> 213
<211> 321
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 213
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ttctcctgca gggccagtca gagcattggc acaagcatcc actggtatca gcaaagaaca 120

aatggctctc caaggcttct cataaagtct gcttctgagt ctatcgctgg gattccttcc 180

aggtttagtg gcagtggatc agggacagat tttactctta gcatcaacag tgtggagtct 240

gaagatattg cagattatta ctgtcaacaa aataataggt ggccgtacac gttcggaggg 300

gggaccaagc tggaaataaa a 321

<210> 214
<211> 339
<212> DNA

<213> Artificial Sequence

<220>
<223> Synthetic

<400> 214
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tcctgcacag cttctgactt taacattaaa gacgactata tacattgggt gaaacagagg 120

cctgaacagg gcctggagtg gattggaagg attgatcctg cgaatggtaa tactaaatat 180

gccccgaagt tccaggacaa ggccactata actgcagaca catcctccaa cacagcctac 240

ctgcagctca gcagcctgac ttctgaggac actgccgtct attactgtgc tctgggctac 300

acctactggg gccaaggcac cactctcaca gtctcctca 339

<210> 215
<211> 336
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 215
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atctcttgca agtcaagtca gagcctctta catagttatg gaaagacata tttgaattgg 120

ttattacaga ggccaggcca gtctccaaag ctcctaatct atctggtgtc taaactggaa 180

tctggagtcc ctgacaggtt cagtggcagt ggatcaggga cagatttcac actgaaaatc 240

agcagagtgg aggctgagga tttgggagtt tattactgct tgcaaactac acattttcct 300

cagacgttcg gtggaggcac caagctggaa atcaag 336

<210> 216
<211> 360
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 216
gaggtgcagc tggtggagtc tgggggagac ttagtgaagc ctggagggtc cctgaaactc 60

tcctgtgcag cctctggatt cactttcagt gactatggaa tgcactgggt tcgtcagggt 120

ccagagaagg ggctggagtg ggttgcatac attaatagtg gcagtagtac catctactat 180

gcagacacag tgaagggccg attcaccatc tccagagaca atgccaagaa caccctgttc 240

ctgcaaatga ccagtctgag gtctgaggac acggccatgt attactgtgc aaggccgggg 300

gattacgaca actatgctat ggactactgg ggtcaaggaa cctcagtcac cgtctcctca 360

<210> 217
<211> 324
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 217
gacattgtga tgacccagtc tcacaaattc ctgtccacat cagtaggaga cagggtcagc 60

atcacctgca aggccagtca ggatgtgagt gttgctgtag cctggtatca acaaaaacca 120

gggcaatctc ctaaactact gatttactgg gcatacaccc ggcacactgg agtccctgat 180

cgcttcacag gcagtggatc tgggacagaa tatactctca ccatcagcag tgtgcaggct 240

gaagacctgg cactttatta ctgtcagcaa cattataaca ctcctccgtg gacgttcggt 300

ggaggcacca agctggaaat caaa 324

<210> 218
<211> 348
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 218
gaggttcagc tgcaacagtc tggggcagag cttgtgaggt caggggcctc agtcaagttg 60

tcctgcacag cttctggctt caacattaaa gactactata tacactgggt gaagcagagg 120

cctgaacagg gcctggagtg gattggatgg attgatcctg agaatgctga tactgaatat 180

gccccgaagt tccagggcaa ggccactatg actccagaca catcctccaa cacagcctac 240

ctgcagctca gcagcctgac atctgaggac actgccgtct attactgtta tgcctgggac 300

tattctatgg actactgggg tcaaggaacc tcagtcaccg tctcctca 348

<210> 219
<211> 336
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 219
gatgttgtga tgacccaaac tccactctcc ctgtctgtca gtcttggaga tcaagcctcc 60

atctcttgca gatctagtca gagccttgta cacagtaatg gaaacaccta tttacattgg 120

tacctgcaga agccgggcca gtctccaaag ctcctgatct acaaagtttc caaccgattt 180

tctggggtcc cagacaggtt cagtggcagt ggatcaggga cagatttcat actcaagatc 240

agtagagtgg aggctgagga tctgggagtt tatttctgct ctcaaaatac acatgttccg 300

tacacgttcg gaggggggac caggctggaa ataaaa 336

<210> 220
<211> 357
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 220
caggttcagc tgcagcagtc tggaactgag ctggcgaggc ctggggcttc agtgaagctg 60

tcctgcaagg cttctggctt caccttcaca gactatggta taaactgggt gaaacagaga 120

actggacagg gccttgagtg gattggagag atttatccta gcagtggtaa ttcttactac 180

aatgagaaat tcaaggccaa ggccacactg actgcagaca aatcctccag cacagcgtac 240

atggagctcc gcagtctgac atctgaggac tctgcggtct atttctgtgc aagatcgact 300

tactacggta gcccaattga ctactggggc caaggcacca ctctcacagt ctcctca 357

<210> 221
<211> 321
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 221
gacattgtga tgacccagtc tcacaaattc atgtccacac cagtaggaga cagggtcagc 60

atcacctgca aggccagtca ggatgtggat actactgtag cctggtatca acagaaacca 120

gggcaatctc cgaaactact gatttactgg gcatccaccc ggcaaattgg agtccctgat 180

cgcttcacag gcagtggatc tgggacagat ttcactctca ccattagtaa tgtgcagtct 240

gaagacttgg cagattattt ctgtcagcaa tatagcacct atcctctgac gttcggtgga 300

ggcaccaagc tggaaatcaa a 321

<210> 222
<211> 357
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 222
caggtgcagc tgaaggagtc aggacctggc ctggtggcgc cctcacagag cctgtccatt 60

acatgcactg tctctgggtt ctcattgacc agctatgcta taacctgggt tcgccagtca 120

ccaggaaagg gtctggagtg gcttggatta tatggactg gtggaggcac aaattataat 180

tcagcgctca aatccagact gagcatcagc aaagacaact ccaagagtca agttttctta 240

aaaatgaaca gtctgcaaac tgatgacaca gccaggtact actgtgccag aatctatgat 300

ggttactacc ggtacttcga tgtctggggc acagggacca cggtcaccgt ctcctca 357

<210> 223

<211> 321
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 223
cgtattgtgt tgacccagac tcccaaattc ctgcttgtat cagcaggaga cagggttacc 60

atgacctgca aggccagtca gagtgtgagt aatgatgtag cttggtacca acagaagcca 120

gggcagtctc ctaaactgct gatatactat gcatccaatc gttacactgg agtccctgat 180

cgcttcactg gcagtggata tgggacggat ttcactttca ccatcagcac tgtgcaggct 240

gaagacctgg cagtttattt ctgtcagcag gattatagct ctccgtggac gttcggtgga 300

ggcaccaagc tggaaatcaa a 321

<210> 224
<211> 348
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 224
gaggttcagc tgcagcagtc tggggcagag cttgtgaggt caggggcctc agtcaagttg 60

tcctgcacag cttctggctt caacattaag gactactata tgcactgggt gaagcagagg 120

cctgaacagg gcctggactg gattggatgg attgatcctg agaatggtga tactgaatat 180

gccccgaaat tccagggcaa ggccactatg actgcagaca catcctccaa cacagcctac 240

ctgcagctca gcagcctgac atctgaggac actgccgtct attactgtaa tgtccttact 300

atgcctacgg cttactgggg ccaagggact ctggtcactg tctctgca 348

<210> 225
<211> 339
<212> DNA
<213> Artificial Sequence

<220>

<223> Synthetic

<400> 225
gacattgtga tgtcacagtc tccatcctcc ctagctgtgt cagttggaga gaaggttatt 60

atgagctgca agtccagtca gagcctttta tatagtagca atcaaaagaa ctatttggcc 120

tggtaccagc agaaaccagg gcagtctcct aaactgctga tttactgggc atccactagg 180

gaatctgggg tccctgatcg cttcacaggc agtggatctg ggacagattt cactctcacc 240

atcagcagtg tgaaggctga agacctggca gtttattact gtcagcaata ttatagctat 300

ccgtacacgt tcggaggggg gaccaagctg gaaataaaa 339

<210> 226
<211> 342
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 226
gaggtgcaac tgatggaatc tgggggagac ttagtgaagc ctggagggtc cctgaaactc 60

tcctgtgcag cctctggatt cactttcagt agctatggct tgtcttgggt tcgccagact 120

ccagacaaga ggctggagtg ggtcgcaacc attactagtg gtggtagtta tacctactat 180

ccagacagtg tgaaggggcg attcaccatc tccagagaca atgccaggaa caccctgtac 240

ctgcaaatgt tcagtctgaa gtctgaggac acagccatgt attactgtgc actatggtcc 300

cttgactact ggggccaagg caccactctc acagtctcct ca 342

<210> 227
<211> 318
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 227
caaattgttc tcacccagtc tccagcaatc atgtctgcat ctccagggga gaaggtcacc 60

atgacctgca gtgccaattc aagtttaagt tacatgcact ggtaccagca gaagccaggc 120

acctccccca aaagatggat ttatgacaca tccgaactgg cttctggagt ccctgctcgc 180

ttcagtggca gtgggtctgg gacctcttat tctctcacaa tcagcagcat ggaggctgaa 240

gatgctgcca cttattactg ccatcagcgg aggagttacc catggacgtt cggtggaggc 300

accaagctgg aaatcaaa 318

<210> 228
<211> 760
<212> PRT
<213> Homo sapiens

<400> 228

Met Met Asp Gln Ala Arg Ser Ala Phe Ser Asn Leu Phe Gly Gly Glu
1 5 10 15

Pro Leu Ser Tyr Thr Arg Phe Ser Leu Ala Arg Gln Val Asp Gly Asp
20 25 30

Asn Ser His Val Glu Met Lys Leu Ala Val Asp Glu Glu Glu Asn Ala
35 40 45

Asp Asn Asn Thr Lys Ala Asn Val Thr Lys Pro Lys Arg Cys Ser Gly
50 55 60

Ser Ile Cys Tyr Gly Thr Ile Ala Val Ile Val Phe Phe Leu Ile Gly
65 70 75 80

Phe Met Ile Gly Tyr Leu Gly Tyr Cys Lys Gly Val Glu Pro Lys Thr
85 90 95

Glu Cys Glu Arg Leu Ala Gly Thr Glu Ser Pro Val Arg Glu Glu Pro
100 105 110

Gly Glu Asp Phe Pro Ala Ala Arg Arg Leu Tyr Trp Asp Asp Leu Lys
115 120 125

```
Arg Lys Leu Ser Glu Lys Leu Asp Ser Thr Asp Phe Thr Gly Thr Ile
    130                 135                 140

Lys Leu Leu Asn Glu Asn Ser Tyr Val Pro Arg Glu Ala Gly Ser Gln
145                 150                 155                 160

Lys Asp Glu Asn Leu Ala Leu Tyr Val Glu Asn Gln Phe Arg Glu Phe
                165                 170                 175

Lys Leu Ser Lys Val Trp Arg Asp Gln His Phe Val Lys Ile Gln Val
            180                 185                 190

Lys Asp Ser Ala Gln Asn Ser Val Ile Ile Val Asp Lys Asn Gly Arg
        195                 200                 205

Leu Val Tyr Leu Val Glu Asn Pro Gly Gly Tyr Val Ala Tyr Ser Lys
    210                 215                 220

Ala Ala Thr Val Thr Gly Lys Leu Val His Ala Asn Phe Gly Thr Lys
225                 230                 235                 240

Lys Asp Phe Glu Asp Leu Tyr Thr Pro Val Asn Gly Ser Ile Val Ile
                245                 250                 255

Val Arg Ala Gly Lys Ile Thr Phe Ala Glu Lys Val Ala Asn Ala Glu
            260                 265                 270

Ser Leu Asn Ala Ile Gly Val Leu Ile Tyr Met Asp Gln Thr Lys Phe
        275                 280                 285

Pro Ile Val Asn Ala Glu Leu Ser Phe Phe Gly His Ala His Leu Gly
    290                 295                 300

Thr Gly Asp Pro Tyr Thr Pro Gly Phe Pro Ser Phe Asn His Thr Gln
305                 310                 315                 320
```

```
Phe Pro Pro Ser Arg Ser Ser Gly Leu Pro Asn Ile Pro Val Gln Thr
                325                 330                 335

Ile Ser Arg Ala Ala Ala Glu Lys Leu Phe Gly Asn Met Glu Gly Asp
            340                 345                 350

Cys Pro Ser Asp Trp Lys Thr Asp Ser Thr Cys Arg Met Val Thr Ser
        355                 360                 365

Glu Ser Lys Asn Val Lys Leu Thr Val Ser Asn Val Leu Lys Glu Ile
    370                 375                 380

Lys Ile Leu Asn Ile Phe Gly Val Ile Lys Gly Phe Val Glu Pro Asp
385                 390                 395                 400

His Tyr Val Val Val Gly Ala Gln Arg Asp Ala Trp Gly Pro Gly Ala
                405                 410                 415

Ala Lys Ser Gly Val Gly Thr Ala Leu Leu Leu Lys Leu Ala Gln Met
            420                 425                 430

Phe Ser Asp Met Val Leu Lys Asp Gly Phe Gln Pro Ser Arg Ser Ile
        435                 440                 445

Ile Phe Ala Ser Trp Ser Ala Gly Asp Phe Gly Ser Val Gly Ala Thr
    450                 455                 460

Glu Trp Leu Glu Gly Tyr Leu Ser Ser Leu His Leu Lys Ala Phe Thr
465                 470                 475                 480

Tyr Ile Asn Leu Asp Lys Ala Val Leu Gly Thr Ser Asn Phe Lys Val
                485                 490                 495

Ser Ala Ser Pro Leu Leu Tyr Thr Leu Ile Glu Lys Thr Met Gln Asn
            500                 505                 510

Val Lys His Pro Val Thr Gly Gln Phe Leu Tyr Gln Asp Ser Asn Trp
```

```
        515                 520                 525

Ala Ser Lys Val Glu Lys Leu Thr Leu Asp Asn Ala Ala Phe Pro Phe
    530                 535                 540

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

Thr Asp Tyr Pro Tyr Leu Gly Thr Thr Met Asp Thr Tyr Lys Glu Leu
                565                 570                 575

Ile Glu Arg Ile Pro Glu Leu Asn Lys Val Ala Arg Ala Ala Ala Glu
            580                 585                 590

Val Ala Gly Gln Phe Val Ile Lys Leu Thr His Asp Val Glu Leu Asn
        595                 600                 605

Leu Asp Tyr Glu Arg Tyr Asn Ser Gln Leu Leu Ser Phe Val Arg Asp
    610                 615                 620

Leu Asn Gln Tyr Arg Ala Asp Ile Lys Glu Met Gly Leu Ser Leu Gln
625                 630                 635                 640

Trp Leu Tyr Ser Ala Arg Gly Asp Phe Phe Arg Ala Thr Ser Arg Leu
                645                 650                 655

Thr Thr Asp Phe Gly Asn Ala Glu Lys Thr Asp Arg Phe Val Met Lys
            660                 665                 670

Lys Leu Asn Asp Arg Val Met Arg Val Glu Tyr His Phe Leu Ser Pro
        675                 680                 685

Tyr Val Ser Pro Lys Glu Ser Pro Phe Arg His Val Phe Trp Gly Ser
    690                 695                 700

Gly Ser His Thr Leu Pro Ala Leu Leu Glu Asn Leu Lys Leu Arg Lys
705                 710                 715                 720
```

```
Gln Asn Asn Gly Ala Phe Asn Glu Thr Leu Phe Arg Asn Gln Leu Ala
                725                 730                 735


Leu Ala Thr Trp Thr Ile Gln Gly Ala Ala Asn Ala Leu Ser Gly Asp
            740                 745                 750


Val Trp Asp Ile Asp Asn Glu Phe
        755                 760


<210>  229
<211>  760
<212>  PRT
<213>  Macaca mulatta

<400>  229

Met Met Asp Gln Ala Arg Ser Ala Phe Ser Asn Leu Phe Gly Gly Glu
1               5                   10                  15


Pro Leu Ser Tyr Thr Arg Phe Ser Leu Ala Arg Gln Val Asp Gly Asp
            20                  25                  30


Asn Ser His Val Glu Met Lys Leu Gly Val Asp Glu Glu Glu Asn Thr
        35                  40                  45


Asp Asn Asn Thr Lys Pro Asn Gly Thr Lys Pro Lys Arg Cys Gly Gly
    50                  55                  60


Asn Ile Cys Tyr Gly Thr Ile Ala Val Ile Ile Phe Phe Leu Ile Gly
65                  70                  75                  80


Phe Met Ile Gly Tyr Leu Gly Tyr Cys Lys Gly Val Glu Pro Lys Thr
                85                  90                  95


Glu Cys Glu Arg Leu Ala Gly Thr Glu Ser Pro Ala Arg Glu Glu Pro
            100                 105                 110
```

```
Glu Glu Asp Phe Pro Ala Ala Pro Arg Leu Tyr Trp Asp Asp Leu Lys
        115                 120                 125

Arg Lys Leu Ser Glu Lys Leu Asp Thr Thr Asp Phe Thr Ser Thr Ile
    130                 135                 140

Lys Leu Leu Asn Glu Asn Leu Tyr Val Pro Arg Glu Ala Gly Ser Gln
145                 150                 155                 160

Lys Asp Glu Asn Leu Ala Leu Tyr Ile Glu Asn Gln Phe Arg Glu Phe
                165                 170                 175

Lys Leu Ser Lys Val Trp Arg Asp Gln His Phe Val Lys Ile Gln Val
            180                 185                 190

Lys Asp Ser Ala Gln Asn Ser Val Ile Ile Val Asp Lys Asn Gly Gly
        195                 200                 205

Leu Val Tyr Leu Val Glu Asn Pro Gly Gly Tyr Val Ala Tyr Ser Lys
    210                 215                 220

Ala Ala Thr Val Thr Gly Lys Leu Val His Ala Asn Phe Gly Thr Lys
225                 230                 235                 240

Lys Asp Phe Glu Asp Leu Asp Ser Pro Val Asn Gly Ser Ile Val Ile
                245                 250                 255

Val Arg Ala Gly Lys Ile Thr Phe Ala Glu Lys Val Ala Asn Ala Glu
            260                 265                 270

Ser Leu Asn Ala Ile Gly Val Leu Ile Tyr Met Asp Gln Thr Lys Phe
        275                 280                 285

Pro Ile Val Lys Ala Asp Leu Ser Phe Phe Gly His Ala His Leu Gly
    290                 295                 300

Thr Gly Asp Pro Tyr Thr Pro Gly Phe Pro Ser Phe Asn His Thr Gln
```

305 310 315 320

Phe Pro Pro Ser Gln Ser Ser Gly Leu Pro Asn Ile Pro Val Gln Thr
325 330 335

Ile Ser Arg Ala Ala Ala Glu Lys Leu Phe Gly Asn Met Glu Gly Asp
340 345 350

Cys Pro Ser Asp Trp Lys Thr Asp Ser Thr Cys Lys Met Val Thr Ser
355 360 365

Glu Asn Lys Ser Val Lys Leu Thr Val Ser Asn Val Leu Lys Glu Thr
370 375 380

Lys Ile Leu Asn Ile Phe Gly Val Ile Lys Gly Phe Val Glu Pro Asp
385 390 395 400

His Tyr Val Val Val Gly Ala Gln Arg Asp Ala Trp Gly Pro Gly Ala
405 410 415

Ala Lys Ser Ser Val Gly Thr Ala Leu Leu Leu Lys Leu Ala Gln Met
420 425 430

Phe Ser Asp Met Val Leu Lys Asp Gly Phe Gln Pro Ser Arg Ser Ile
435 440 445

Ile Phe Ala Ser Trp Ser Ala Gly Asp Phe Gly Ser Val Gly Ala Thr
450 455 460

Glu Trp Leu Glu Gly Tyr Leu Ser Ser Leu His Leu Lys Ala Phe Thr
465 470 475 480

Tyr Ile Asn Leu Asp Lys Ala Val Leu Gly Thr Ser Asn Phe Lys Val
485 490 495

Ser Ala Ser Pro Leu Leu Tyr Thr Leu Ile Glu Lys Thr Met Gln Asp
500 505 510

```
Val Lys His Pro Val Thr Gly Arg Ser Leu Tyr Gln Asp Ser Asn Trp
        515                 520                 525

Ala Ser Lys Val Glu Lys Leu Thr Leu Asp Asn Ala Ala Phe Pro Phe
    530                 535                 540

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

Thr Asp Tyr Pro Tyr Leu Gly Thr Thr Met Asp Thr Tyr Lys Glu Leu
                565                 570                 575

Val Glu Arg Ile Pro Glu Leu Asn Lys Val Ala Arg Ala Ala Ala Glu
            580                 585                 590

Val Ala Gly Gln Phe Val Ile Lys Leu Thr His Asp Thr Glu Leu Asn
        595                 600                 605

Leu Asp Tyr Glu Arg Tyr Asn Ser Gln Leu Leu Leu Phe Leu Arg Asp
    610                 615                 620

Leu Asn Gln Tyr Arg Ala Asp Val Lys Glu Met Gly Leu Ser Leu Gln
625                 630                 635                 640

Trp Leu Tyr Ser Ala Arg Gly Asp Phe Phe Arg Ala Thr Ser Arg Leu
                645                 650                 655

Thr Thr Asp Phe Arg Asn Ala Glu Lys Arg Asp Lys Phe Val Met Lys
            660                 665                 670

Lys Leu Asn Asp Arg Val Met Arg Val Glu Tyr Tyr Phe Leu Ser Pro
        675                 680                 685

Tyr Val Ser Pro Lys Glu Ser Pro Phe Arg His Val Phe Trp Gly Ser
    690                 695                 700
```

Gly Ser His Thr Leu Ser Ala Leu Leu Glu Ser Leu Lys Leu Arg Arg
705 710 715 720

Gln Asn Asn Ser Ala Phe Asn Glu Thr Leu Phe Arg Asn Gln Leu Ala
725 730 735

Leu Ala Thr Trp Thr Ile Gln Gly Ala Ala Asn Ala Leu Ser Gly Asp
740 745 750

Val Trp Asp Ile Asp Asn Glu Phe
755 760

<210> 230
<211> 760
<212> PRT
<213> Macaca fascicularis

<400> 230

Met Met Asp Gln Ala Arg Ser Ala Phe Ser Asn Leu Phe Gly Gly Glu
1 5 10 15

Pro Leu Ser Tyr Thr Arg Phe Ser Leu Ala Arg Gln Val Asp Gly Asp
20 25 30

Asn Ser His Val Glu Met Lys Leu Gly Val Asp Glu Glu Glu Asn Thr
35 40 45

Asp Asn Asn Thr Lys Ala Asn Gly Thr Lys Pro Lys Arg Cys Gly Gly
50 55 60

Asn Ile Cys Tyr Gly Thr Ile Ala Val Ile Ile Phe Phe Leu Ile Gly
65 70 75 80

Phe Met Ile Gly Tyr Leu Gly Tyr Cys Lys Gly Val Glu Pro Lys Thr
85 90 95

Glu Cys Glu Arg Leu Ala Gly Thr Glu Ser Pro Ala Arg Glu Glu Pro

```
            100                 105                 110

Glu Glu Asp Phe Pro Ala Ala Pro Arg Leu Tyr Trp Asp Asp Leu Lys
        115                 120                 125

Arg Lys Leu Ser Glu Lys Leu Asp Thr Thr Asp Phe Thr Ser Thr Ile
    130                 135                 140

Lys Leu Leu Asn Glu Asn Leu Tyr Val Pro Arg Glu Ala Gly Ser Gln
145                 150                 155                 160

Lys Asp Glu Asn Leu Ala Leu Tyr Ile Glu Asn Gln Phe Arg Glu Phe
                165                 170                 175

Lys Leu Ser Lys Val Trp Arg Asp Gln His Phe Val Lys Ile Gln Val
            180                 185                 190

Lys Asp Ser Ala Gln Asn Ser Val Ile Ile Val Asp Lys Asn Gly Gly
        195                 200                 205

Leu Val Tyr Leu Val Glu Asn Pro Gly Gly Tyr Val Ala Tyr Ser Lys
    210                 215                 220

Ala Ala Thr Val Thr Gly Lys Leu Val His Ala Asn Phe Gly Thr Lys
225                 230                 235                 240

Lys Asp Phe Glu Asp Leu Asp Ser Pro Val Asn Gly Ser Ile Val Ile
                245                 250                 255

Val Arg Ala Gly Lys Ile Thr Phe Ala Glu Lys Val Ala Asn Ala Glu
            260                 265                 270

Ser Leu Asn Ala Ile Gly Val Leu Ile Tyr Met Asp Gln Thr Lys Phe
        275                 280                 285

Pro Ile Val Lys Ala Asp Leu Ser Phe Phe Gly His Ala His Leu Gly
    290                 295                 300
```

```
Thr Gly Asp Pro Tyr Thr Pro Gly Phe Pro Ser Phe Asn His Thr Gln
305                 310                 315                 320

Phe Pro Pro Ser Gln Ser Ser Gly Leu Pro Asn Ile Pro Val Gln Thr
                325                 330                 335

Ile Ser Arg Ala Ala Ala Glu Lys Leu Phe Gly Asn Met Glu Gly Asp
            340                 345                 350

Cys Pro Ser Asp Trp Lys Thr Asp Ser Thr Cys Lys Met Val Thr Ser
        355                 360                 365

Glu Asn Lys Ser Val Lys Leu Thr Val Ser Asn Val Leu Lys Glu Thr
    370                 375                 380

Lys Ile Leu Asn Ile Phe Gly Val Ile Lys Gly Phe Val Glu Pro Asp
385                 390                 395                 400

His Tyr Val Val Val Gly Ala Gln Arg Asp Ala Trp Gly Pro Gly Ala
                405                 410                 415

Ala Lys Ser Ser Val Gly Thr Ala Leu Leu Leu Lys Leu Ala Gln Met
            420                 425                 430

Phe Ser Asp Met Val Leu Lys Asp Gly Phe Gln Pro Ser Arg Ser Ile
        435                 440                 445

Ile Phe Ala Ser Trp Ser Ala Gly Asp Phe Gly Ser Val Gly Ala Thr
    450                 455                 460

Glu Trp Leu Glu Gly Tyr Leu Ser Ser Leu His Leu Lys Ala Phe Thr
465                 470                 475                 480

Tyr Ile Asn Leu Asp Lys Ala Val Leu Gly Thr Ser Asn Phe Lys Val
                485                 490                 495
```

```
Ser Ala Ser Pro Leu Leu Tyr Thr Leu Ile Glu Lys Thr Met Gln Asp
            500                 505                 510

Val Lys His Pro Val Thr Gly Arg Ser Leu Tyr Gln Asp Ser Asn Trp
        515                 520                 525

Ala Ser Lys Val Glu Lys Leu Thr Leu Asp Asn Ala Ala Phe Pro Phe
    530                 535                 540

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

Thr Asp Tyr Pro Tyr Leu Gly Thr Thr Met Asp Thr Tyr Lys Glu Leu
                565                 570                 575

Val Glu Arg Ile Pro Glu Leu Asn Lys Val Ala Arg Ala Ala Ala Glu
            580                 585                 590

Val Ala Gly Gln Phe Val Ile Lys Leu Thr His Asp Thr Glu Leu Asn
        595                 600                 605

Leu Asp Tyr Glu Arg Tyr Asn Ser Gln Leu Leu Leu Phe Leu Arg Asp
    610                 615                 620

Leu Asn Gln Tyr Arg Ala Asp Val Lys Glu Met Gly Leu Ser Leu Gln
625                 630                 635                 640

Trp Leu Tyr Ser Ala Arg Gly Asp Phe Phe Arg Ala Thr Ser Arg Leu
                645                 650                 655

Thr Thr Asp Phe Arg Asn Ala Glu Lys Arg Asp Lys Phe Val Met Lys
            660                 665                 670

Lys Leu Asn Asp Arg Val Met Arg Val Glu Tyr Tyr Phe Leu Ser Pro
        675                 680                 685
```

Tyr Val Ser Pro Lys Glu Ser Pro Phe Arg His Val Phe Trp Gly Ser
690 695 700

Gly Ser His Thr Leu Ser Ala Leu Leu Glu Ser Leu Lys Leu Arg Arg
705 710 715 720

Gln Asn Asn Ser Ala Phe Asn Glu Thr Leu Phe Arg Asn Gln Leu Ala
725 730 735

Leu Ala Thr Trp Thr Ile Gln Gly Ala Ala Asn Ala Leu Ser Gly Asp
740 745 750

Val Trp Asp Ile Asp Asn Glu Phe
755 760

<210> 231
<211> 763
<212> PRT
<213> Mus musculus

<400> 231

Met Met Asp Gln Ala Arg Ser Ala Phe Ser Asn Leu Phe Gly Gly Glu
1 5 10 15

Pro Leu Ser Tyr Thr Arg Phe Ser Leu Ala Arg Gln Val Asp Gly Asp
20 25 30

Asn Ser His Val Glu Met Lys Leu Ala Ala Asp Glu Glu Glu Asn Ala
35 40 45

Asp Asn Asn Met Lys Ala Ser Val Arg Lys Pro Lys Arg Phe Asn Gly
50 55 60

Arg Leu Cys Phe Ala Ala Ile Ala Leu Val Ile Phe Phe Leu Ile Gly
65 70 75 80

Phe Met Ser Gly Tyr Leu Gly Tyr Cys Lys Arg Val Glu Gln Lys Glu
85 90 95

```
Glu Cys Val Lys Leu Ala Glu Thr Glu Glu Thr Asp Lys Ser Glu Thr
            100                 105                 110

Met Glu Thr Glu Asp Val Pro Thr Ser Ser Arg Leu Tyr Trp Ala Asp
        115                 120                 125

Leu Lys Thr Leu Leu Ser Glu Lys Leu Asn Ser Ile Glu Phe Ala Asp
    130                 135                 140

Thr Ile Lys Gln Leu Ser Gln Asn Thr Tyr Thr Pro Arg Glu Ala Gly
145                 150                 155                 160

Ser Gln Lys Asp Glu Ser Leu Ala Tyr Tyr Ile Glu Asn Gln Phe His
                165                 170                 175

Glu Phe Lys Phe Ser Lys Val Trp Arg Asp Glu His Tyr Val Lys Ile
            180                 185                 190

Gln Val Lys Ser Ser Ile Gly Gln Asn Met Val Thr Ile Val Gln Ser
        195                 200                 205

Asn Gly Asn Leu Asp Pro Val Glu Ser Pro Glu Gly Tyr Val Ala Phe
    210                 215                 220

Ser Lys Pro Thr Glu Val Ser Gly Lys Leu Val His Ala Asn Phe Gly
225                 230                 235                 240

Thr Lys Lys Asp Phe Glu Glu Leu Ser Tyr Ser Val Asn Gly Ser Leu
                245                 250                 255

Val Ile Val Arg Ala Gly Glu Ile Thr Phe Ala Glu Lys Val Ala Asn
            260                 265                 270

Ala Gln Ser Phe Asn Ala Ile Gly Val Leu Ile Tyr Met Asp Lys Asn
        275                 280                 285
```

```
Lys Phe Pro Val Val Glu Ala Asp Leu Ala Leu Phe Gly His Ala His
    290                 295                 300

Leu Gly Thr Gly Asp Pro Tyr Thr Pro Gly Phe Pro Ser Phe Asn His
305                 310                 315                 320

Thr Gln Phe Pro Pro Ser Gln Ser Ser Gly Leu Pro Asn Ile Pro Val
                325                 330                 335

Gln Thr Ile Ser Arg Ala Ala Ala Glu Lys Leu Phe Gly Lys Met Glu
            340                 345                 350

Gly Ser Cys Pro Ala Arg Trp Asn Ile Asp Ser Ser Cys Lys Leu Glu
        355                 360                 365

Leu Ser Gln Asn Gln Asn Val Lys Leu Ile Val Lys Asn Val Leu Lys
    370                 375                 380

Glu Arg Arg Ile Leu Asn Ile Phe Gly Val Ile Lys Gly Tyr Glu Glu
385                 390                 395                 400

Pro Asp Arg Tyr Val Val Val Gly Ala Gln Arg Asp Ala Leu Gly Ala
                405                 410                 415

Gly Val Ala Ala Lys Ser Ser Val Gly Thr Gly Leu Leu Leu Lys Leu
            420                 425                 430

Ala Gln Val Phe Ser Asp Met Ile Ser Lys Asp Gly Phe Arg Pro Ser
        435                 440                 445

Arg Ser Ile Ile Phe Ala Ser Trp Thr Ala Gly Asp Phe Gly Ala Val
    450                 455                 460

Gly Ala Thr Glu Trp Leu Glu Gly Tyr Leu Ser Ser Leu His Leu Lys
465                 470                 475                 480
```

```
Ala Phe Thr Tyr Ile Asn Leu Asp Lys Val Val Leu Gly Thr Ser Asn
                485                 490                 495

Phe Lys Val Ser Ala Ser Pro Leu Leu Tyr Thr Leu Met Gly Lys Ile
            500                 505                 510

Met Gln Asp Val Lys His Pro Val Asp Gly Lys Ser Leu Tyr Arg Asp
        515                 520                 525

Ser Asn Trp Ile Ser Lys Val Glu Lys Leu Ser Phe Asp Asn Ala Ala
    530                 535                 540

Tyr Pro Phe Leu Ala Tyr Ser Gly Ile Pro Ala Val Ser Phe Cys Phe
545                 550                 555                 560

Cys Glu Asp Ala Asp Tyr Pro Tyr Leu Gly Thr Arg Leu Asp Thr Tyr
                565                 570                 575

Glu Ala Leu Thr Gln Lys Val Pro Gln Leu Asn Gln Met Val Arg Thr
            580                 585                 590

Ala Ala Glu Val Ala Gly Gln Leu Ile Ile Lys Leu Thr His Asp Val
        595                 600                 605

Glu Leu Asn Leu Asp Tyr Glu Met Tyr Asn Ser Lys Leu Leu Ser Phe
    610                 615                 620

Met Lys Asp Leu Asn Gln Phe Lys Thr Asp Ile Arg Asp Met Gly Leu
625                 630                 635                 640

Ser Leu Gln Trp Leu Tyr Ser Ala Arg Gly Asp Tyr Phe Arg Ala Thr
                645                 650                 655

Ser Arg Leu Thr Thr Asp Phe His Asn Ala Glu Lys Thr Asn Arg Phe
            660                 665                 670

Val Met Arg Glu Ile Asn Asp Arg Ile Met Lys Val Glu Tyr His Phe
```

675 680 685

Leu Ser Pro Tyr Val Ser Pro Arg Glu Ser Pro Phe Arg His Ile Phe
690 695 700

Trp Gly Ser Gly Ser His Thr Leu Ser Ala Leu Val Glu Asn Leu Lys
705 710 715 720

Leu Arg Gln Lys Asn Ile Thr Ala Phe Asn Glu Thr Leu Phe Arg Asn
725 730 735

Gln Leu Ala Leu Ala Thr Trp Thr Ile Gln Gly Val Ala Asn Ala Leu
740 745 750

Ser Gly Asp Ile Trp Asn Ile Asp Asn Glu Phe
755 760

<210> 232
<211> 19
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 232

Met Gly Trp Ser Cys Ile Ile Leu Phe Leu Val Ala Thr Ala Thr Gly
1 5 10 15

Val His Ser

<210> 233
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 233

Ile Asp Pro Glu Thr Gly Asp Thr
1 5

<210> 234
<211> 17
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 234

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

Asp

<210> 235
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<220>
<221> misc_feature
<222> (3)..(3)
<223> Xaa can be any naturally occurring amino acid

<400> 235

Leu Pro Xaa Thr Gly
1 5

<210> 236
<211> 17
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 236

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

Asp

<210> 237
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 237

Gly Tyr Ser Phe Thr Asp Tyr Tyr
1   5

<210> 238
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 238

Asp Tyr Tyr Ile Asn
1   5

<210> 239
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 239

Gly Tyr Ser Phe Thr Asp Tyr Asp
1 5

<210> 240
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 240

Asp Tyr Asp Ile Asn
1 5

<210> 241
<211> 25
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 241

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

Ser Leu Arg Leu Ser Cys Ala Ala Ser
20 25

<210> 242
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 242

Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val

1 5 10

<210> 243
<211> 30
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 243

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

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

<210> 244
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 244

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

<210> 245
<211> 25
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 245

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

Ser Val Lys Val Ser Cys Lys Ala Ser
20 25

<210> 246
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 246

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

<210> 247
<211> 30
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 247

Arg Val Thr Ile Thr Ala Asp Thr Ser Thr Ser Thr Ala Tyr Met Glu
1 5 10 15

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

<210> 248
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 248

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

<210> 249
<211> 25
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 249

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

Thr Leu Ser Leu Thr Cys Thr Val Ser
20 25

<210> 250
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 250

Trp Ile Arg Gln Pro Pro Gly Lys Gly Leu Glu Trp Ile
1 5 10

<210> 251
<211> 30
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 251

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

Leu Ser Ser Val Thr Ala Ala Asp Thr Ala Val Tyr Tyr Cys
20 25 30

<210> 252
<211> 23
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 252

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

Glu Arg Ala Thr Ile Asn Cys
20

<210> 253
<211> 23
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 253

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
20

<210> 254
<211> 15
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 254

Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile Tyr

1 5 10 15

<210> 255
<211> 15
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 255

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

<210> 256
<211> 32
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 256

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

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

<210> 257
<211> 23
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 257

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

Glu Pro Ala Ser Ile Ser Cys
20

<210> 258
<211> 15
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 258

Trp Tyr Leu Gln Lys Pro Gly Gln Ser Pro Gln Leu Leu Ile Tyr
1 5 10 15

<210> 259
<211> 32
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 259

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

Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Val Gly Val Tyr Tyr Cys
20 25 30

<210> 260
<211> 32
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 260

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

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

<210> 261
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 261

Gly Tyr Thr Phe Thr Asp Tyr Gly
1 5

<210> 262
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 262

Gly Phe Ser Ile Thr Ser Gly Tyr Tyr
1 5

<210> 263
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 263

Ile Tyr Pro Ser Ser Gly Asn Thr
1 5

<210> 264
<211> 7

<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 264

Ile Asn Phe Asp Gly Asn Asn
1 5

<210> 265
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 265

Ile Asn Phe Asp Gly Asn Ser
1 5

<210> 266
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 266

Ala Arg Ser Asn Tyr Tyr Gly Ser Pro Ile Asp Phe
1 5 10

<210> 267
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 267

Ala Arg Phe Tyr Tyr Asp Tyr Asp Tyr Phe Asp Tyr
1 5 10

<210> 268
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 268

Ala Arg Ile Tyr Tyr Asp Tyr Asp Tyr Phe Asp Tyr
1 5 10

<210> 269
<211> 447
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 269

Glu Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Val Arg Pro Gly Ala
1 5 10 15

Ser Val Lys Leu Ser Cys Thr Ala Ser Gly Phe Asn Ile Lys Asp Asp
20 25 30

Tyr Met Tyr Trp Val Lys Gln Arg Pro Glu Gln Gly Leu Glu Trp Ile
35 40 45

Gly Trp Ile Asp Pro Glu Thr Gly Asp Thr Glu Tyr Ala Ser Lys Phe
50 55 60

Gln Asp Lys Ala Thr Val Thr Ala Asp Thr Ser Ser Asn Thr Ala Tyr
65 70 75 80

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

Thr Leu Trp Leu Arg Arg Gly Leu Asp Tyr Trp Gly Gln Gly Thr Ser
            100                 105                 110

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

Ala Pro Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys
    130                 135                 140

Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser
145                 150                 155                 160

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

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

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

Thr Lys Val Asp Lys Lys Val Glu Pro Lys Ser Cys Asp Lys Thr His
    210                 215                 220

Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val
225                 230                 235                 240

Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr
                245                 250                 255

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

Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys
```

275 280 285

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

Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys
305 310 315 320

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

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

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

Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn
370 375 380

Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser
385 390 395 400

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

Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala Leu
420 425 430

His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys
435 440 445

<210> 270
<211> 447
<212> PRT
<213> Artificial Sequence

<220>

<223> Synthetic

<400> 270

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Glu Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Val Arg Pro Gly Ala
1               5                  10                  15

Ser Val Lys Leu Ser Cys Thr Ala Ser Gly Phe Asn Ile Lys Asp Asp
            20                  25                  30

Tyr Met Tyr Trp Val Lys Gln Arg Pro Glu Gln Gly Leu Glu Trp Ile
        35                  40                  45

Gly Trp Ile Asp Pro Glu Ser Gly Asp Thr Glu Tyr Ala Ser Lys Phe
    50                  55                  60

Gln Asp Lys Ala Thr Val Thr Ala Asp Thr Ser Ser Asn Thr Ala Tyr
65                  70                  75                  80

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

Thr Leu Trp Leu Arg Arg Gly Leu Asp Tyr Trp Gly Gln Gly Thr Ser
            100                 105                 110

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

Ala Pro Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys
    130                 135                 140

Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser
145                 150                 155                 160

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

Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser
```

```
            180                 185                 190

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

Thr Lys Val Asp Lys Lys Val Glu Pro Lys Ser Cys Asp Lys Thr His
    210                 215                 220

Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val
225                 230                 235                 240

Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr
                245                 250                 255

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

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

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

Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys
305                 310                 315                 320

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

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

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

Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn
    370                 375                 380
```

Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser
385 390 395 400

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

Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala Leu
420 425 430

His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys
435 440 445

<210> 271
<211> 450
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 271

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

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

Tyr Ile Asn Trp Val Asn Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile
35 40 45

Gly Trp Ile Tyr Pro Gly Ser Gly Asn Thr Arg Tyr Ser Glu Arg Phe
50 55 60

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

Met Gln Leu Ser Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Phe Cys

```
                85                  90                  95

Ala Arg Glu Asp Tyr Tyr Pro Tyr His Gly Met Asp Tyr Trp Gly Gln
            100                 105                 110

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

Gly Lys
    450
```

<210> 272
<211> 450
<212> PRT

<213> Artificial Sequence

<220>
<223> Synthetic

<400> 272

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


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


Asp Ile Asn Trp Val Asn Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile
        35                  40                  45


Gly Trp Ile Tyr Pro Gly Ser Gly Asn Thr Arg Tyr Ser Glu Arg Phe
    50                  55                  60


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


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


Ala Arg Glu Asp Tyr Tyr Pro Tyr His Gly Met Asp Tyr Trp Gly Gln
            100                 105                 110


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


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


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


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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

Gly Lys
    450
```

<210> 273
<211> 225
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic

<400> 273

```
Glu Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Val Arg Pro Gly Ala
1               5                   10                  15

Ser Val Lys Leu Ser Cys Thr Ala Ser Gly Phe Asn Ile Lys Asp Asp
            20                  25                  30

Tyr Met Tyr Trp Val Lys Gln Arg Pro Glu Gln Gly Leu Glu Trp Ile
        35                  40                  45

Gly Trp Ile Asp Pro Glu Thr Gly Asp Thr Glu Tyr Ala Ser Lys Phe
    50                  55                  60
```

```
Gln Asp Lys Ala Thr Val Thr Ala Asp Thr Ser Ser Asn Thr Ala Tyr
65                  70                  75                  80

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

Thr Leu Trp Leu Arg Arg Gly Leu Asp Tyr Trp Gly Gln Gly Thr Ser
            100                 105                 110

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

Ala Pro Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys
    130                 135                 140

Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser
145                 150                 155                 160

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

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

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

Thr Lys Val Asp Lys Lys Val Glu Pro Lys Ser Cys Asp Lys Thr His
    210                 215                 220

Thr
225
```

<210> 274
<211> 225
<212> PRT

<213> Artificial Sequence

<220>
<223> Synthetic

<400> 274

```
Glu Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Val Arg Pro Gly Ala
1               5                   10                  15


Ser Val Lys Leu Ser Cys Thr Ala Ser Gly Phe Asn Ile Lys Asp Asp
            20                  25                  30


Tyr Met Tyr Trp Val Lys Gln Arg Pro Glu Gln Gly Leu Glu Trp Ile
        35                  40                  45


Gly Trp Ile Asp Pro Glu Ser Gly Asp Thr Glu Tyr Ala Ser Lys Phe
    50                  55                  60


Gln Asp Lys Ala Thr Val Thr Ala Asp Thr Ser Ser Asn Thr Ala Tyr
65                  70                  75                  80


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


Thr Leu Trp Leu Arg Arg Gly Leu Asp Tyr Trp Gly Gln Gly Thr Ser
            100                 105                 110


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


Ala Pro Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys
    130                 135                 140


Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser
145                 150                 155                 160


Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu Gln Ser
                165                 170                 175
```

```
Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser
            180                 185                 190


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


Thr Lys Val Asp Lys Lys Val Glu Pro Lys Ser Cys Asp Lys Thr His
    210                 215                 220


Thr
225


<210>  275
<211>  228
<212>  PRT
<213>  Artificial Sequence

<220>
<223>  Synthetic

<400>  275

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


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


Tyr Ile Asn Trp Val Asn Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile
        35                  40                  45


Gly Trp Ile Tyr Pro Gly Ser Gly Asn Thr Arg Tyr Ser Glu Arg Phe
    50                  55                  60


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


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

Ala Arg Glu Asp Tyr Tyr Pro Tyr His Gly Met Asp Tyr Trp Gly Gln
100 105 110

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

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

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

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

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

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

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

Lys Thr His Thr
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Ser Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr Ser Phe Thr Asp Tyr
            20                  25                  30


Asp Ile Asn Trp Val Asn Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile
        35                  40                  45


Gly Trp Ile Tyr Pro Gly Ser Gly Asn Thr Arg Tyr Ser Glu Arg Phe
    50                  55                  60


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


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


Ala Arg Glu Asp Tyr Tyr Pro Tyr His Gly Met Asp Tyr Trp Gly Gln
            100                 105                 110


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


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


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


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


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


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

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

Lys Thr His Thr
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Gln Asp Ile Ser Asn Tyr
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Gln Gln Gly Asn Thr Leu Pro Trp Thr
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Gln Gln Gly Asn Thr Leu Pro Tyr Thr
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Gln Gln Gly His Ala Tyr Pro Tyr Thr
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Gly Tyr Ser Ile Thr Ser Gly Tyr Xaa
1 5

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Ile Xaa Phe Asp Gly Xaa Xaa
1 5

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Xaa Arg Xaa Xaa Tyr Asp Tyr Asp Xaa Xaa Asp Xaa
1 5 10

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1   5

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Gln Gln Gly Xaa Xaa Xaa Pro Xaa Thr
1   5