

<110> MEDIMMUNE LIMITED

<120> ANTICUERPOS ANTI-PAR2 Y SUS USOS

<130> 1848081-0002-093-W01

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<150> 62/637,766

<151> 2018-03-02

<150> 62/472,462

<151> 2017-03-16

<160> 841

<170> PatentIn versión 3.5

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

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

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

Leu Gln Met Asn	Ser Leu Arg Ala	Glu Asp Thr Ala	Val Tyr Tyr Cys	
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Ala Arg Ile Asn	Asn Asp Pro Met	Asp Val Trp Gly	Gln Gly Thr Leu	
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Val Thr Val Ser	Ser			
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Ser Tyr Ala Met	Asn			
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Gln Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Val Leu Val Ile Tyr
 35 40 45

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys Gln Thr Trp Asp Gly Asn Pro Thr Thr
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Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
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Ala Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ser Thr Ile Ser Tyr His Gly Ser His Ile Ser Tyr His Asp Ser Val
50 55 60

His His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
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Gln Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Val Leu Val Ile Tyr
 35 40 45

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

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

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
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 20 25 30

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

Ser Thr Ile Ser Tyr Ser Gly Ser Leu Ile His Tyr Ala Asp Ser Val
 50 55 60

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
 100 105 110

Val Thr Val Ser Ser
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 Ser Tyr Ala Met Asn
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Gly

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 20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

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

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

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

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

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

Ala Arg Ile Asn His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
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Ser Tyr Ala Met Asn
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Gly

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

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

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

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20 25 30

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

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

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
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Val Thr Val Ser Ser
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20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
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20 25 30

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

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

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

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
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Ala Arg Ile His His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
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Gly

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20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
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Asp Glu Ala Asp Tyr Tyr Cys Gln Thr Trp Asp Gly Asn Pro Thr Thr
85 90 95

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

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

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

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

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

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
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Ala Arg Ile His His Asp Pro Met Asp His Trp Gly Gln Gly Thr Leu
 100 105 110

Val Thr Val Ser Ser
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 20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys Gln Thr Trp Asp Gly Asn Pro Thr Thr
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Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

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 Asp Asp Ser Asn Arg Pro Ser
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 20 25 30

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

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

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

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys 95
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Val Thr Val Ser Ser 115
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<210> 74
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Gly

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 1 5

<210> 76
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 20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

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

Gly Glu Thr Asn His Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

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 ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac 300
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 <213> Homo sapiens

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1515648ENSP_Sequences.txt

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

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

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

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Ser Tyr Ala Met Asn
1 5

<210> 84
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<212> PRT
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Gly

<210> 85
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<212> PRT
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<400> 85
Ile Asn Asn Asp Pro Met Asp Val
1 5

<210> 86
<211> 333
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ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
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cacttcggcg gagggaccaa gttaaccgtc cta 333

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<213> Homo sapiens

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20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys His His Trp His Gly Asn His His Thr
85 90 95

Gly Glu Thr Asn His Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

<210> 88
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<400> 88
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<210> 89
<211> 7
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<400> 89
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1 5

<210> 90
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<400> 90
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<212> ADN
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ccaggcaagg gcctggagtg ggtgtccacc atcagctaca gcgggtcgct gatcagctac 180
gccgacagcg tgaagggcag gttcaccatc agcagggaca acagcaagaa caccctgtac 240
ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac 300
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<210> 92
<211> 117
<212> PRT
<213> Homo sapiens

<400> 92
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1 5 10 15

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

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

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

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

<210> 93
<211> 5
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<213> Homo sapiens

<400> 93
Ser Tyr Ala Met Asn
1 5

<210> 94
<211> 17
<212> PRT
<213> Homo sapiens

<400> 94
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1 5 10 15

Gly

<210> 95
<211> 8
<212> PRT
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<400> 95
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<210> 96
<211> 333
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ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
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gtgttcggcg gagggaccaa gttaaccgtc cta 333

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<211> 111
<212> PRT
<213> Homo sapiens

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys His Thr Trp His His His Pro Thr His
85 90 95

Gly Glu His Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

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<210> 99
<211> 7
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<400> 99
Asp Asp Ser Asn Arg Pro Ser
1 5

<210> 100
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 ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac 300
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 <212> PRT
 <213> Homo sapiens

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

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

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

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
 100 105 110

Val Thr Val Ser Ser
 115

<210> 103
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 <213> Homo sapiens

<400> 103
 Ser Tyr Ala Met Asn
 1 5

<210> 104
 <211> 17
 <212> PRT
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<400> 104
 Thr Ile Ser Tyr Ser Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val Lys
 1 5 10 15

Gly

<210> 105
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 <212> PRT
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cagtcccccg tgctggtgat ctacgacgac agcaacaggc ccagcggcat ccccgagagg      180
ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg      240
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<400> 107

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          20          25          30

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

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

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Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
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Asp Glu Ala Asp Tyr Tyr Cys Gln Thr Trp Asp Gly His His Thr His
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Gly Glu His His Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
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<210> 108

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<212> PRT

<213> Homo sapiens

<400> 108

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Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
1           5           10

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<210> 109

<211> 7

<212> PRT

<213> Homo sapiens

<400> 109

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Asp Asp Ser Asn Arg Pro Ser
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<210> 110

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<212> PRT

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Gln Thr Trp Asp Gly His His Thr His Gly Glu His His Val
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<210> 111

<211> 351

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ccaggcaagg gcctggagtg ggtgtccacc atcagctaca gcgggtcgct gatcagctac      180
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ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac      300
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<211> 117

<212> PRT

<213> Homo sapiens

<400> 112

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

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

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

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
 100 105 110

Val Thr Val Ser Ser
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<210> 113

<211> 5

<212> PRT

<213> Homo sapiens

<400> 113

Ser Tyr Ala Met Asn
 1 5

<210> 114

<211> 17

<212> PRT

<213> Homo sapiens

<400> 114

Thr Ile Ser Tyr Ser Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val Lys
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Gly

<210> 115

<211> 8

<212> PRT

<213> Homo sapiens

<400> 115

Ile Asn Asn Asp Pro Met Asp Val
 1 5

<210> 116

<211> 333

<212> ADN

<213> Homo sapiens

<400> 116

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 ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
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 gtgttcggcg gagggaccaa gtaaccgtc cta 333

<210> 117

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<213> Homo sapiens

<400> 117

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 1 5 10 15

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

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

35

40

45

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

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

Gly His Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

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<400> 118
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<210> 119
<211> 7
<212> PRT
<213> Homo sapiens

<400> 119
Asp Asp Ser Asn Arg Pro Ser
1 5

<210> 120
<211> 14
<212> PRT
<213> Homo sapiens

<400> 120
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1 5 10

<210> 121
<211> 351
<212> ADN
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<400> 121
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<210> 122
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<213> Homo sapiens

<400> 122
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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
20 25 30

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

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

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser

115

<210> 123
 <211> 5
 <212> PRT
 <213> Homo sapiens

<400> 123
 Ser Tyr Ala Met Asn
 1 5

<210> 124
 <211> 17
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<400> 124
 Thr Ile Ser Tyr Ser Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val Lys
 1 5 10 15

Gly

<210> 125
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 <212> PRT
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<400> 125
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<210> 126
 <211> 333
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 ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
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<400> 127
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Thr Ala Ser Ile Thr Cys Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val
 20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys His Thr Trp His His His His His Thr
 85 90 95

Gly Glu His Asn His Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

<210> 128
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 <212> PRT
 <213> Homo sapiens

<400> 128
 Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
 1 5 10

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<213> Homo sapiens

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Asp Asp Ser Asn Arg Pro Ser
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<400> 130

His Thr Trp His His His His Thr Gly Glu His Asn His
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<213> Homo sapiens

<400> 132

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

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

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

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

<210> 133

<211> 5

<212> PRT

<213> Homo sapiens

<400> 133

Ser Tyr Ala Met Asn
1 5

<210> 134

<211> 17

<212> PRT

<213> Homo sapiens

<400> 134

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Gly

<210> 135

<211> 8

<212> PRT

<213> Homo sapiens

<400> 135
Ile Asn Asn Asp Pro Met Asp Val
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<210> 136
<211> 333
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gtgttcggcg gagggaccaa gttaaccgtc cta 333

<210> 137
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<400> 137
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Thr Ala Ser Ile Thr Cys Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val
20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys Gln His Trp His Gly His Pro His His
85 90 95

Gly Glu His Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

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Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
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Asp Asp Ser Asn Arg Pro Ser
1 5

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<212> PRT
<213> Homo sapiens

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

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

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

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

<210> 143
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<213> Homo sapiens

<400> 143
Ser Tyr Ala Met Asn
1 5

<210> 144
<211> 17
<212> PRT
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<400> 144
Thr Ile Ser Tyr Ser Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val Lys
1 5 10 15

Gly

<210> 145
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<212> PRT
<213> Homo sapiens

<400> 145
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1 5

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cacttcggcg gagggaccaa gttaaccgtc cta 333

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys Gln His Trp Asp Gly Asn His His His
85 90 95

Gly Glu Thr Asn His Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

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<400> 148
Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
1 5 10

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<211> 7
<212> PRT
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<400> 149
Asp Asp Ser Asn Arg Pro Ser
1 5

<210> 150
<211> 14
<212> PRT
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1 5 10

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ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac 300
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<212> PRT
<213> Homo sapiens

<400> 152
Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
20 25 30

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

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

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
 100 105 110

Val Thr Val Ser Ser
 115

<210> 153
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<400> 153
 Ser Tyr Ala Met Asn
 1 5

<210> 154
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 Thr Ile Ser Tyr Ser Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val Lys
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Gly

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 gtgttcggcg gagggaccaa gtaaccgtc cta 333

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 20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys His Thr Trp Asp His His His Thr Thr
 85 90 95

Gly Glu His Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

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Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
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Asp Asp Ser Asn Arg Pro Ser
1 5

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<213> Homo sapiens

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20 25 30

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

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

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
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Ser Tyr Ala Met Asn
1 5

<210> 164
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Thr Ile Ser Tyr Ser Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val Lys
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Gly

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 cacttcggcg gagggaccaa gttaaccgtc cta 333

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 20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys His His Trp His Gly His His Thr Thr
 85 90 95

Gly Glu Thr Asn His Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

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 Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
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<400> 169
 Asp Asp Ser Asn Arg Pro Ser
 1 5

<210> 170
 <211> 14
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 1 5 10

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 20 25 30

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

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

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
 100 105 110

Val Thr Val Ser Ser
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 Ser Tyr Ala Met Asn
 1 5

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 Thr Ile Ser Tyr Ser Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val Lys
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Gly

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 20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys His His Trp His Gly His Pro His Thr
 85 90 95

Gly Glu His Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

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 Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
 1 5 10

<210> 179
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 <212> PRT
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<400> 179
 Asp Asp Ser Asn Arg Pro Ser
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<210> 180
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 20 25 30

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

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

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

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Ser Tyr Ala Met Asn
1 5

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Gly

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1 5

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cacttcggcg gagggaccaa gtaaccgtc cta 333

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys Gln His Trp Asp Gly His His Thr Thr
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Gly Glu Thr Asn His Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

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 Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
 1 5 10

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 Asp Asp Ser Asn Arg Pro Ser
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 gccgacagcg tgaagggcag gttcaccatc agcagggaca acagcaagaa caccctgtac 240
 ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac 300
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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
 20 25 30

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

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

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
 100 105 110

Val Thr Val Ser Ser
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 20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

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 85 90 95

Gly Glu Thr Asn His Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

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 cacgacagcc atcatggcag gttcaccatc agcagggaca acagcaagaa caccctgtac 240
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 20 25 30

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

Ser Thr Ile Ser His Ser Gly His His Ile Ser His His Asp Ser His
 50 55 60

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

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

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 100 105 110

Val Thr Val Ser Ser
 115

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<210> 204
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Gly

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 cagtcccccg tgctggtgat ctacgacgac agcaacaggc ccagcggcat ccccgagagg 180
 ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
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1515648ENSP_Sequences.txt

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20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

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85 90 95

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

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1 5 10

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<211> 7
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Asp Asp Ser Asn Arg Pro Ser
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ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac 300
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20 25 30

1515648ENSP_Sequences.txt

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

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

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<400> 213
Ser Tyr Ala Met Asn
1 5

<210> 214
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Gly

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<400> 215
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1 5

<210> 216
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ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
gacgaggccg actactactg ccagacctgg gacgggaacc cgacgacggg cgagaccaac 300
gtgttcggcg gagggaccaa gttaaccgtc cta 333

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20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

1515648ENSP_Sequences.txt
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Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
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<210> 219
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<213> Homo sapiens

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<210> 220
<211> 14
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ccaggcaagg gcctggagtg ggtgtccacc atccaccaca gcgggcacct gcacagccac 180
gccgaccacg tgaagggcag gttcaccatc agcagggaca acagcaagaa caccctgtac 240
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20 25 30

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

Ser Thr Ile His His Ser Gly His Leu His Ser His Ala Asp His Val
50 55 60

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

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Ser Tyr Ala Met Asn
1 5

<210> 224
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 Thr Ile His His Ser Gly His Leu His Ser His Ala Asp His Val Lys
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Gly

<210> 225
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 ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
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 gtgttcggcg gagggaccaa gtaaccgtc cta 333

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 20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

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<210> 229
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<210> 230
 <211> 14
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 <213> Homo sapiens

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 20 25 30

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

Ser Thr Ile Ser His Ser Gly His Leu Ile His His Ala Asp Ser His
 50 55 60

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
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Val Thr Val Ser Ser
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Gly

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ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg    240
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Gln Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Val Leu Val Ile Tyr
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Asp Glu Ala Asp Tyr Tyr Cys Gln Thr Trp Asp Gly Asn Pro Thr Thr
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<212> PRT
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<210> 240
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Ala Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45
Ser Thr Ile Ser Tyr His Gly Ser His Ile Ser Tyr His His Ser His
50 55 60
Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95
Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110
Val Thr Val Ser Ser
115

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<212> PRT
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<400> 243
Ser Tyr Ala Met Asn
1 5

<210> 244
<211> 17
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<400> 244
Thr Ile Ser Tyr His Gly Ser His Ile Ser Tyr His His Ser His Lys
1 5 10 15

Gly

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<400> 245
Ile Asn Asn Asp Pro Met Asp Val
1 5

<210> 246
<211> 333
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<213> Homo sapiens

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20 25 30

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

Asp Asp Ser Asn Arg Pro Ser Gly Ile Pro Glu Arg Phe Ser Gly Ser
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1          5          10

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Asp Asp Ser Asn Arg Pro Ser
1          5

<210> 250
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<212> ADN
<213> Homo sapiens

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ccaggcaagg gcctggagtg ggtgtccacc atcagctacc acgggtcgca catcagctac      180
gccgaccacc ataagcacag gttcaccatc agcagggaca acagcaagaa caccctgtac      240
ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac      300
aacgacccta tggatgtgtg gggccagggc accctggtga cagtgaagctc a          351

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<212> PRT
<213> Homo sapiens

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

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

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

Ser Thr Ile Ser Tyr His Gly Ser His Ile Ser Tyr Ala Asp His His
50          55          60

Lys His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65          70          75          80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
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Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100         105         110

Val Thr Val Ser Ser
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His

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 1 5

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

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

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

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 Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
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<210> 259
 <211> 7
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<400> 259
 Asp Asp Ser Asn Arg Pro Ser

1 5

<210> 260
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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
 20 25 30

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

Ser Thr Ile Ser Tyr Ser Gly His Leu Ile His Tyr Ala Asp His His
 50 55 60

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
 100 105 110

Val Thr Val Ser Ser
 115

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<400> 263
 Ser Tyr Ala Met Asn
 1 5

<210> 264
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 <212> PRT
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<400> 264
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Gly

<210> 265
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<210> 266
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 20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

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

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 100 105 110

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 Asp Asp Ser Asn Arg Pro Ser
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 <213> Homo sapiens

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

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

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

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

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

 Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
 100 105 110

 Val Thr Val Ser Ser
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 <400> 273
 Ser Tyr Ala Met Asn
 1 5

<210> 274
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 <212> PRT
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 <400> 274
 Thr Ile Ser Tyr His Gly Ser His Ile Ser His Ala Asp His Val His
 1 5 10 15

Gly

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 <212> PRT
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<400> 275
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 1 5

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<400> 277
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 20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

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

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

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<400> 279
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1 5

<210> 280
<211> 14
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1 5 10

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

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

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

Lys His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

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<400> 283
Ser Tyr Ala Met Asn
1 5

<210> 284
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Thr Ile Ser Tyr His Gly His His Ile Ser His Ala Asp His Val Lys
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His

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20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

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85 90 95

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

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

Ser Thr Ile Ser His Ser Gly His Leu Ile His His His His Ser Val
 50 55 60

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

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

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

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100 105 110

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 20 25 30

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

Ser Thr Ile Ser Tyr His Gly Ser His Ile Ser Tyr His Asp Ser His
 50 55 60

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

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

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Val Thr Val Ser Ser
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<400> 303
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Gly

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20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
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<400> 308

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Asp Asp Ser Asn Arg Pro Ser
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Gln Thr Trp Asp Gly Asn Pro Thr Thr Gly Glu Thr Asn Val
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20 25 30

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35 40 45

Ser Thr Ile Ser Tyr His His His Leu Ile Ser His Ala Asp Ser Val
50 55 60

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

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
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Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
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<400> 313
Ser Tyr Ala Met Asn
1 5

<210> 314
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1 5 10 15

Gly

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<212> PRT
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<212> PRT
<213> Homo sapiens

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

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

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

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<213> Homo sapiens

<400> 318

Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
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<212> PRT

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Asp Asp Ser Asn Arg Pro Ser
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<400> 320

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<211> 351

<212> ADN

<213> Homo sapiens

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ccaggcaagg gcctggagtg ggtgtccacc atcagctacc acgggcacca catcagccac	180
gccgaccacc ataagggcag gttcaccatc agcagggaca acagcaagaa caccctgtac	240
ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac	300
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<211> 117

<212> PRT

<213> Homo sapiens

<400> 322

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20 25 30

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

Ser Thr Ile Ser Tyr His Gly His His Ile Ser His Ala Asp His His
50 55 60

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

<210> 323

<211> 5

<212> PRT

<213> Homo sapiens

<400> 323

Ser Tyr Ala Met Asn
1 5

<210> 324

<211> 17

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<213> Homo sapiens

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Thr Ile Ser Tyr His Gly His His Ile Ser His Ala Asp His His Lys
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 <212> PRT
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 Ile Asn Asn Asp Pro Met Asp Val
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<210> 326
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<400> 327
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Thr Ala Ser Ile Thr Cys Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val
 20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

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

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

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 1 5 10

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<400> 329
 Asp Asp Ser Asn Arg Pro Ser
 1 5

<210> 330
 <211> 14
 <212> PRT
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<400> 330
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<210> 331
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ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac 300
aacgacccta tggatgtgtg gggccagggc accctggtga cagttagctc a 351

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<212> PRT
<213> Homo sapiens

<400> 332
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20 25 30

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

Ser Thr Ile Ser Tyr His Gly His His Ile Ser Tyr Ala Asp Ser His
50 55 60

His His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

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<213> Homo sapiens

<400> 333
Ser Tyr Ala Met Asn
1 5

<210> 334
<211> 17
<212> PRT
<213> Homo sapiens

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Thr Ile Ser Tyr His Gly His His Ile Ser Tyr Ala Asp Ser His His
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His

<210> 335
<211> 8
<212> PRT
<213> Homo sapiens

<400> 335
Ile Asn Asn Asp Pro Met Asp Val
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<210> 336
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ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
gacgaggccg actactactg ccagacctgg gacgggaacc cgacgacggg cgagaccaac 300
gtgttcggcg gagggaccaa gttaaccgtc cta 333

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65      70      75      80

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

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
      100      105      110

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<400> 338
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1      5      10

<210> 339
<211> 7
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<400> 339
Asp Asp Ser Asn Arg Pro Ser
1      5

<210> 340
<211> 14
<212> PRT
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<400> 340
Gln Thr Trp Asp Gly Asn Pro Thr Thr Gly Glu Thr Asn Val
1      5      10

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ccaggcaagg gcctggagtg ggtgtccacc atccaccaca gcgggcacct gcacagctac      180
caccaccacg tgaagggcag gttcaccatc agcagggaaca acagcaagaa caccctgtac      240
ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac      300
aacgacccta tggatgtgtg gggccagggc accctggtga cagtgaagctc a      351

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<213> Homo sapiens

<400> 342
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      20      25      30

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

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1515648ENSP_Sequences.txt
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Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

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<211> 5
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<213> Homo sapiens

<400> 343
Ser Tyr Ala Met Asn
1 5

<210> 344
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Thr Ile His His Ser Gly His Leu His Ser Tyr His His His Val Lys
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Gly

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<400> 345
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1 5

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ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
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gtgttcggcg gagggaccaa gttaaccgtc cta 333

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<212> PRT
<213> Homo sapiens

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20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

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

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

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<210> 349
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<400> 349
 Asp Asp Ser Asn Arg Pro Ser
 1 5

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 20 25 30

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

Ser Thr Ile Ser His Ser Gly His Leu His His Tyr Ala Asp Ser His
 50 55 60

Lys His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
 65 70 75 80

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
 100 105 110

Val Thr Val Ser Ser
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 <213> Homo sapiens

<400> 353
 Ser Tyr Ala Met Asn
 1 5

<210> 354
 <211> 17

<212> PRT

<213> Homo sapiens

<400> 354

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His

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 ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
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 gtgttcggcg gagggaccaa gttaaccgtc cta 333

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<213> Homo sapiens

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 20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

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

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
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<213> Homo sapiens

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Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
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<210> 359

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 ccaggcaagg gcctggagtg ggtgtccacc atcagctacc acgggtcgca catcagctac 180
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 ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac 300
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 <212> PRT
 <213> Homo sapiens

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 20 25 30
 Ala Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
 35 40 45
 Ser Thr Ile Ser Tyr His Gly Ser His Ile Ser Tyr His Asp Ser Val
 50 55 60
 His His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
 65 70 75 80
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
 85 90 95
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 Val Thr Val Ser Ser
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 Ser Tyr Ala Met Asn
 1 5

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His

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 20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
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Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
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aacgacccta tggatgtgtg gggccagggc accctggtga cagtgcgctc a	351

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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
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Ala Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45
Ser Thr Ile Ser Tyr His Gly His His Ile Ser His Ala His His Val
50 55 60
His Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95
Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110
Val Thr Val Ser Ser
115

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Ser Tyr Ala Met Asn
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<210> 374
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Gly

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ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
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gtgttcggcg gagggaccaa gttaaccgtc cta 333

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20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
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65              70              75              80
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<210> 380
<211> 14
<212> PRT
<213> Homo sapiens

<400> 380
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1      5              10

<210> 381
<211> 351
<212> ADN
<213> Homo sapiens

<400> 381
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ccaggcaagg gcctggagtg ggtgtccacc atcagctaca gcgggtcgct gatcagctac      180
gccgacagcg tgaagggcag gttcaccatc agcagggaca acagcaagaa caccctgtac      240
ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatccac      300
caccaccctc acgatcactg gggccagggc accctggtga cagtgaagctc a      351

<210> 382
<211> 117
<212> PRT
<213> Homo sapiens

<400> 382
Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1      5              10             15

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

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

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

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

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

Ala Arg Ile His His His Pro His Asp His Trp Gly Gln Gly Thr Leu
      100             105             110

Val Thr Val Ser Ser
      115

<210> 383
<211> 5
<212> PRT
<213> Homo sapiens

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<400> 383
Ser Tyr Ala Met Asn
1 5

<210> 384
<211> 17
<212> PRT
<213> Homo sapiens

<400> 384
Thr Ile Ser Tyr Ser Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val Lys
1 5 10 15

Gly

<210> 385
<211> 8
<212> PRT
<213> Homo sapiens

<400> 385
Ile His His His Pro His Asp His
1 5

<210> 386
<211> 333
<212> ADN
<213> Homo sapiens

<400> 386
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ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
gacgaggccg actactactg ccagacctgg gacgggaacc cgacgacggg cgagaccaac 300
gtgttcggcg gagggaccaa gttaaccgtc cta 333

<210> 387
<211> 111
<212> PRT
<213> Homo sapiens

<400> 387
Ser Ile Glu Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly Gln
1 5 10 15

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

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

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

<210> 388
<211> 11
<212> PRT
<213> Homo sapiens

<400> 388
Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
1 5 10

<210> 389
<211> 7
<212> PRT
<213> Homo sapiens

<400> 389
Asp Asp Ser Asn Arg Pro Ser
1 5

<210> 390

<211> 14
 <212> PRT
 <213> Homo sapiens

<400> 390
 Gln Thr Trp Asp Gly Asn Pro Thr Thr Gly Glu Thr Asn Val
 1 5 10

<210> 391
 <211> 351
 <212> ADN
 <213> Homo sapiens

<400> 391
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 gccgacagcg tgaagggcag gttcaccatc agcagggaca acagcaagaa caccctgtac 240
 ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac 300
 cacgacccta tggatcactg gggccagggc accctggtga cagtgaagctc a 351

<210> 392
 <211> 117
 <212> PRT
 <213> Homo sapiens

<400> 392
 Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1 5 10 15

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

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

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

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

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

Ala Arg Ile Asn His Asp Pro Met Asp His Trp Gly Gln Gly Thr Leu
 100 105 110

Val Thr Val Ser Ser
 115

<210> 393
 <211> 5
 <212> PRT
 <213> Homo sapiens

<400> 393
 Ser Tyr Ala Met Asn
 1 5

<210> 394
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 394
 Thr Ile Ser Tyr Ser Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val Lys
 1 5 10 15

Gly

<210> 395
 <211> 8
 <212> PRT
 <213> Homo sapiens

<400> 395
 Ile Asn His Asp Pro Met Asp His
 1 5

<210> 396
 <211> 333
 <212> ADN

<213> Homo sapiens

<400> 396

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cagtcccccg tgctggtgat ctacgacgac agcaacaggc ccagcggcat ccccgagagg      180
ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg      240
gacgaggccg actactactg ccagacctgg gacgggaacc cgacgacggg cgagaccaac      300
gtgttcggcg gagggaccaa gttaaccgtc cta                                  333

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<210> 397

<211> 111

<212> PRT

<213> Homo sapiens

<400> 397

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

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

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

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

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Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65          70          75          80

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

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Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
          100          105          110

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<210> 398

<211> 11

<212> PRT

<213> Homo sapiens

<400> 398

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Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
1           5           10

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<210> 399

<211> 7

<212> PRT

<213> Homo sapiens

<400> 399

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Asp Asp Ser Asn Arg Pro Ser
1           5

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<210> 400

<211> 14

<212> PRT

<213> Homo sapiens

<400> 400

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Gln Thr Trp Asp Gly Asn Pro Thr Thr Gly Glu Thr Asn Val
1           5           10

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<210> 401

<211> 351

<212> ADN

<213> Homo sapiens

<400> 401

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ccaggcaagg gcctggagtg ggtgtccacc atcagctaca gcgggtcgct gatcagctac      180
gccgacagcg tgaagggcag gttcaccatc agcagggaca acagcaagaa caccctgtac      240
ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac      300
aacgacccta tggatgtgtg gggccagggc accctggtga cagttagctc a              351

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<210> 402

<211> 117

<212> PRT

<213> Homo sapiens

<400> 402
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 1 5 10 15

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

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

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

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
 100 105 110

Val Thr Val Ser Ser
 115

<210> 403
 <211> 5
 <212> PRT
 <213> Homo sapiens

<400> 403
 Ser Tyr Ala Met Asn
 1 5

<210> 404
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 404
 Thr Ile Ser Tyr Ser Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val Lys
 1 5 10 15

Gly

<210> 405
 <211> 8
 <212> PRT
 <213> Homo sapiens

<400> 405
 Ile Asn Asn Asp Pro Met Asp Val
 1 5

<210> 406
 <211> 333
 <212> ADN
 <213> Homo sapiens

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 cagtcccccg tgctggtgat ctacgacgac agcaacaggc ccagcggcat ccccgagagg 180
 ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
 gacgaggccg actactactg ccacacctgg gacgggcacc acacgacggg ggagaccaac 300
 cacttcggcg gagggaccaa gttaaccgtc cta 333

<210> 407
 <211> 111
 <212> PRT
 <213> Homo sapiens

<400> 407
 Ser Ile Glu Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly Gln
 1 5 10 15

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys His Thr Trp Asp Gly His His Thr Thr
85 90 95

Gly Glu Thr Asn His Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

<210> 408
<211> 11
<212> PRT
<213> Homo sapiens

<400> 408
Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
1 5 10

<210> 409
<211> 7
<212> PRT
<213> Homo sapiens

<400> 409
Asp Asp Ser Asn Arg Pro Ser
1 5

<210> 410
<211> 14
<212> PRT
<213> Homo sapiens

<400> 410
His Thr Trp Asp Gly His His Thr Thr Gly Glu Thr Asn His
1 5 10

<210> 411
<211> 351
<212> ADN
<213> Homo sapiens

<400> 411
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ccaggcaagg gcctggagtg ggtgtccacc atcagctaca gcgggtcgct gatcagctac 180
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ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcdc ccgcatcaac 300
aacgacccta tggatgtgtg gggccagggc accctggtga cagtgaagctc a 351

<210> 412
<211> 117
<212> PRT
<213> Homo sapiens

<400> 412
Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

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

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

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

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

<210> 413
 <211> 5
 <212> PRT
 <213> Homo sapiens

<400> 413
 Ser Tyr Ala Met Asn
 1 5

<210> 414
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 414
 Thr Ile Ser Tyr Ser Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val Lys
 1 5 10 15

Gly

<210> 415
 <211> 8
 <212> PRT
 <213> Homo sapiens

<400> 415
 Ile Asn Asn Asp Pro Met Asp Val
 1 5

<210> 416
 <211> 333
 <212> ADN
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<400> 416
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 acctgcagcg gcgacaacct gggcaagaaa tacgtgcagt ggtatcagca gaagcccggc 120
 cagtcccccg tgctggtgat ctacgacgac agcaacaggc ccagcggcat ccccgagagg 180
 ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
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 gtgttcggcg gagggaccaa gttaaccgtc cta 333

<210> 417
 <211> 111
 <212> PRT
 <213> Homo sapiens

<400> 417
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 1 5 10 15

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys Gln His Trp Asp Gly His His His Thr
 85 90 95

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

<210> 418
 <211> 11
 <212> PRT
 <213> Homo sapiens

<400> 418
 Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
 1 5 10

<210> 419
 <211> 7
 <212> PRT
 <213> Homo sapiens

<400> 419
 Asp Asp Ser Asn Arg Pro Ser
 1 5

<210> 420
 <211> 14
 <212> PRT
 <213> Homo sapiens

<400> 420
 Gln His Trp Asp Gly His His His Thr Gly Glu Thr Asn Val
 1 5 10

<210> 421
 <211> 351
 <212> ADN
 <213> Homo sapiens

<400> 421
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 ccaggcaagg gcctggagtg ggtgtccacc atcagctaca gcgggtcgct gatcagctac 180
 gccgacagcg tgaagggcag gttcaccatc agcagggaca acagcaagaa caccctgtac 240
 ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac 300
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 <211> 117
 <212> PRT
 <213> Homo sapiens

<400> 422
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 1 5 10 15

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

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

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

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
 100 105 110

Val Thr Val Ser Ser
 115

<210> 423
 <211> 5
 <212> PRT
 <213> Homo sapiens

<400> 423
 Ser Tyr Ala Met Asn
 1 5

<210> 424
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 424
 Thr Ile Ser Tyr Ser Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val Lys
 1 5 10 15

Gly

<210> 425
 <211> 8
 <212> PRT
 <213> Homo sapiens

<400> 425

Ile Asn Asn Asp Pro Met Asp Val
1 5

<210> 426
<211> 333
<212> ADN
<213> Homo sapiens

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ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
gacgagggcg actactactg ccagcactgg cacgggaacc cgcacacggg ggagcacaac 300
cacttcggcg gagggaccaa gttaaccgtc cta 333

<210> 427
<211> 111
<212> PRT
<213> Homo sapiens

<400> 427
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1 5 10 15

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

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

Gly Glu His Asn His Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

<210> 428
<211> 11
<212> PRT
<213> Homo sapiens

<400> 428
Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
1 5 10

<210> 429
<211> 7
<212> PRT
<213> Homo sapiens

<400> 429
Asp Asp Ser Asn Arg Pro Ser
1 5

<210> 430
<211> 14
<212> PRT
<213> Homo sapiens

<400> 430
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<210> 431
<211> 351
<212> ADN
<213> Homo sapiens

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<211> 117
<212> PRT
<213> Homo sapiens

<400> 432
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1 5 10 15

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

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

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

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

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<211> 5
<212> PRT
<213> Homo sapiens

<400> 433
Ser Tyr Ala Met Asn
1 5

<210> 434
<211> 17
<212> PRT
<213> Homo sapiens

<400> 434
Thr Ile Ser Tyr Ser Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val Lys
1 5 10 15

Gly

<210> 435
<211> 8
<212> PRT
<213> Homo sapiens

<400> 435
Ile Asn Asn Asp Pro Met Asp Val
1 5

<210> 436
<211> 333
<212> ADN
<213> Homo sapiens

<400> 436
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ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
gacgaggccg actactactg ccacacctgg gaccacaacc cgcaccatca tcacaccaac 300
cacttcggcg gagggaccaa gtaaccgtc cta 333

<210> 437
<211> 111
<212> PRT
<213> Homo sapiens

<400> 437
Ser Ile Glu Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly Gln
1 5 10 15

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

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

His His Thr Asn His Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

<210> 438
<211> 11
<212> PRT
<213> Homo sapiens

<400> 438
Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
1 5 10

<210> 439
<211> 7
<212> PRT
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<400> 439
Asp Asp Ser Asn Arg Pro Ser
1 5

<210> 440
<211> 14
<212> PRT
<213> Homo sapiens

<400> 440
His Thr Trp Asp His Asn Pro His His His His Thr Asn His
1 5 10

<210> 441
<211> 351
<212> ADN
<213> Homo sapiens

<400> 441
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ccaggcaagg gcctggagtg ggtgtccacc atcagctaca gcgggtcgct gatcagctac 180
gccgacagcg tgaagggcag gttcaccatc agcagggaca acagcaagaa caccctgtac 240
ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac 300
aacgacccta tggatgtgtg gggccagggc accctggtga cagtgaagctc a 351

<210> 442
<211> 117
<212> PRT
<213> Homo sapiens

<400> 442
Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

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

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

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

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
 100 105 110

Val Thr Val Ser Ser
 115

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 Ser Tyr Ala Met Asn
 1 5

<210> 444
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 Thr Ile Ser Tyr Ser Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val Lys
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Gly

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 1 5

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<400> 447
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 1 5 10 15

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 20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

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

His Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

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1              5              10

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1              5

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ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac      300
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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
              20              25              30

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

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

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

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100              105              110

Val Thr Val Ser Ser
115

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<400> 453
Ser Tyr Ala Met Asn
1              5

<210> 454
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Thr Ile Ser Tyr Ser Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val Lys
1              5              10              15

Gly

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 1 5

<210> 456
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 20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys Gln His Trp His Gly His His His Thr
 85 90 95

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

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 <212> PRT
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<400> 458
 Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
 1 5 10

<210> 459
 <211> 7
 <212> PRT
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<400> 459
 Asp Asp Ser Asn Arg Pro Ser
 1 5

<210> 460
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 <212> PRT
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<400> 460
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 1 5 10

<210> 461
 <211> 351
 <212> ADN
 <213> Homo sapiens

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1515648ENSP_Sequences.txt

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ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac 300
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<212> PRT
<213> Homo sapiens

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20 25 30

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

Ser Thr Ile Ser Tyr Ser Gly Ser Leu Ile His Tyr Ala Asp Ser Val
50 55 60

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

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

Ala Arg Ile Asn His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

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Ser Tyr Ala Met Asn
1 5

<210> 464
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Gly

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1 5

<210> 466
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ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggccatg 240
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gtgttcggcg gagggaccaa gttaaccgtc cta 333

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<211> 111

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<212> PRT
<213> Homo sapiens

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      20     25     30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65      70      75      80

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

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
      100    105    110

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<212> PRT
<213> Homo sapiens

<400> 468
Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
1      5      10

<210> 469
<211> 7
<212> PRT
<213> Homo sapiens

<400> 469
Asp Asp Ser Asn Arg Pro Ser
1      5

<210> 470
<211> 14
<212> PRT
<213> Homo sapiens

<400> 470
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<213> Homo sapiens

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ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcdc ccgcatcaac      300
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<212> PRT
<213> Homo sapiens

<400> 472
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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
      20     25     30

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

Ser Thr Ile Ser His Ser Gly His His Ile Ser His His Asp Ser His
      50     55     60

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

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

Ala Arg Ile Asn His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
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Ser Tyr Ala Met Asn
1 5

<210> 474
<211> 17
<212> PRT
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Gly

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<212> PRT
<213> Homo sapiens

<400> 475
Ile Asn His Asp Pro Met Asp Val
1 5

<210> 476
<211> 333
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cagtcccccg tgctggtgat ctacgacgac agcaacaggc ccagcggcat ccccgagagg 180
ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
gacgaggccg actactactg ccagacctgg gacgggaacc cgacgacggg cgagaccaac 300
gtgttcggcg gagggaccaa gttaaccgtc cta 333

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<211> 111
<212> PRT
<213> Homo sapiens

<400> 477
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1 5 10 15

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

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

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

<210> 478
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 1 5 10

<210> 479
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 <212> PRT
 <213> Homo sapiens

<400> 479
 Asp Asp Ser Asn Arg Pro Ser
 1 5

<210> 480
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<400> 480
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 gccgacagcc ataagggcag gttcaccatc agcagggaca acagcaagaa caccctgtac 240
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 <213> Homo sapiens

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

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

Ser Thr Ile Ser His Ser Gly His Leu Ile His His Ala Asp Ser His
 50 55 60

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

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

Ala Arg Ile Asn His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
 100 105 110

Val Thr Val Ser Ser
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 1 5

<210> 484
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Gly

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 1 5

<210> 486
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 gtgttcggcg gagggaccaa gttaaccgtc cta 333

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<400> 487
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 1 5 10 15

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

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

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

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<210> 489
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<400> 489
 Asp Asp Ser Asn Arg Pro Ser
 1 5

<210> 490
 <211> 14
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<210> 491
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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
          20          25          30

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

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

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

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Ala Arg Ile Asn His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
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Val Thr Val Ser Ser
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<210> 493

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<212> PRT

<213> Homo sapiens

<400> 493

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Ser Tyr Ala Met Asn
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<210> 494

<211> 17

<212> PRT

<213> Homo sapiens

<400> 494

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Thr Ile Ser Tyr Ser Gly His Leu Ile His Tyr Ala Asp His His His
1          5          10          15

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Gly

<210> 495

<211> 8

<212> PRT

<213> Homo sapiens

<400> 495

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Ile Asn His Asp Pro Met Asp Val
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<210> 496

<211> 333

<212> ADN

<213> Homo sapiens

<400> 496

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cagtcccccg tgctggtgat ctacgacgac agcaacaggc ccagcggcat ccccagagagg      180
ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg      240

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gtgttcggcg gagggaccaa gttaaccgtc cta 333

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<212> PRT
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20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
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Asp Glu Ala Asp Tyr Tyr Cys Gln Thr Trp Asp Gly Asn Pro Thr Thr
85 90 95

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

<210> 498
<211> 11
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<400> 498
Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
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<210> 499
<211> 7
<212> PRT
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<400> 499
Asp Asp Ser Asn Arg Pro Ser
1 5

<210> 500
<211> 14
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<400> 500
Gln Thr Trp Asp Gly Asn Pro Thr Thr Gly Glu Thr Asn Val
1 5 10

<210> 501
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<212> ADN
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<213> Homo sapiens

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

Ala Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
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35

40

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

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

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

Ala Arg Ile Asn His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

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<213> Homo sapiens

<400> 503
Ser Tyr Ala Met Asn
1 5

<210> 504
<211> 17
<212> PRT
<213> Homo sapiens

<400> 504
Thr Ile Ser Tyr His Gly Ser His Ile Ser His Ala Asp His Val His
1 5 10 15

Gly

<210> 505
<211> 8
<212> PRT
<213> Homo sapiens

<400> 505
Ile Asn His Asp Pro Met Asp Val
1 5

<210> 506
<211> 333
<212> ADN
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35 40 45

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
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Asp Glu Ala Asp Tyr Tyr Cys Gln Thr Trp Asp Gly Asn Pro Thr Thr
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His Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
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<211> 351
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gccgaccacg tgcattggcag gttcaccatc agcagggaaca acagcaagaa caccctgtac 240
ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatccac 300
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<213> Homo sapiens

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1 5 10 15

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

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

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

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

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

Ala Arg Ile His His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

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Ser Tyr Ala Met Asn
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<210> 574
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<212> PRT
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<400> 574
Thr Ile Ser Tyr His Gly Ser His Ile Ser His Ala Asp His Val His
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Gly

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<212> PRT
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ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
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1 5 10 15

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

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

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

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<211> 7
 <212> PRT
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<400> 579
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 1 5

<210> 580
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 ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatccac 300
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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
 20 25 30

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

Ser Thr Ile Ser Tyr His Gly Ser His Ile Ser Tyr His Asp Ser His
 50 55 60

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

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

Ala Arg Ile His His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
 100 105 110

Val Thr Val Ser Ser
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 Ser Tyr Ala Met Asn
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<210> 584
 <211> 17
 <212> PRT
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Gly

<210> 585
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Ile His His Asp Pro Met Asp Val
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cagtcccccg tgctggtgat ctacgacgac agcaacaggc ccagcggcat ccccgagagg	180
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35 40 45Asp Asp Ser Asn Arg Pro Ser Gly Ile Pro Glu Arg Phe Ser Gly Ser
50 55 60Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
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100 105 110

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<211> 11

<212> PRT

<213> Homo sapiens

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<210> 589

<211> 7

<212> PRT

<213> Homo sapiens

<400> 589

Asp Asp Ser Asn Arg Pro Ser
1 5

<210> 590

<211> 14

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<400> 590

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<211> 351

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<213> Homo sapiens

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ccaggcaagg gcctggagtg ggtgtccacc atcagctacc acgggcacca catcagctac	180
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 20 25 30

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

Ser Thr Ile Ser Tyr His Gly His His Ile Ser Tyr Ala Asp Ser His
 50 55 60

His His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
 65 70 75 80

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

Ala Arg Ile His His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
 100 105 110

Val Thr Val Ser Ser
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<210> 594
 <211> 17
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His

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20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

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

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
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<400> 599
Asp Asp Ser Asn Arg Pro Ser
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<210> 600
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<210> 601
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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
20 25 30

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

Ser Thr Ile Ser Tyr His Gly His His Ile Ser Tyr Ala Asp Ser His
50 55 60

His His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

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

Ala Arg Ile His His Asp Pro Met Asp His Trp Gly Gln Gly Thr Leu
 100 105 110

Val Thr Val Ser Ser
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 20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys Gln Thr Trp Asp Gly Asn Pro Thr Thr
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Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
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 Asp Asp Ser Asn Arg Pro Ser
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 ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcmc ccgcatccac 300
 cagcacccta tggatcactg gggccagggc accctggtga cagtgcgctc a 351

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 20 25 30

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

Ser Thr Ile Ser Tyr His Gly Ser His Ile Ser Tyr His Asp Ser Val
 50 55 60

His His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
 65 70 75 80

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

Ala Arg Ile His His Asp Pro Met Asp His Trp Gly Gln Gly Thr Leu
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Val Thr Val Ser Ser
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 1 5

<210> 614
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His

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 20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys Gln Thr Trp Asp Gly Asn Pro Thr Thr
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Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

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 <212> PRT
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 <212> PRT
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<400> 619
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 <211> 14
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<400> 620
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 1 5 10

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<400> 621

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ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac      300
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Ala Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
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Ser Thr Ile Ser Tyr Ser Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val
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Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
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Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
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Val Thr Val Ser Ser
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 20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

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

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
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<210> 629
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 <211> 14
 <212> PRT
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 <212> ADN
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 Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1 5 10 15

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

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

Ser Thr Ile Ser Tyr Ser Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val

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50          55          60
Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
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Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
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Val Thr Val Ser Ser
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Ser Tyr Ala Met Asn
1          5

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Gly

<210> 635
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<400> 635
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<210> 636
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Gln Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Val Leu Val Ile Tyr
35         40         45

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65         70         75         80

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

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu

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100

105

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<400> 638
Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
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1 5

<210> 640
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<213> Homo sapiens

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

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

Ser Thr Ile Ser Tyr His Gly His His Ile Ser Tyr Ala Asp Ser His
50 55 60

His His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

<210> 643
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Ser Tyr Ala Met Asn
1 5

<210> 644
<211> 17
<212> PRT

<213> Homo sapiens

<400> 644

Thr Ile Ser Tyr His Gly His His Ile Ser Tyr Ala Asp Ser His His
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His

<210> 645

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<212> PRT

<213> Homo sapiens

<400> 645

Ile Asn Asn Asp Pro Met Asp Val
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<210> 646

<211> 333

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20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

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

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
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<400> 648

Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
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<210> 649

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<213> Homo sapiens

<400> 649

Asp Asp Ser Asn Arg Pro Ser
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 20 25 30

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

Ser Thr Ile Ser Tyr His Gly His His Ile Ser Tyr Ala Asp Ser His 60
 50 55 60

His His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr 80
 65 70 75 80

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

Ala Arg Ile Asn His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu 110
 100 105 110

Val Thr Val Ser Ser 115

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<400> 653
 Ser Tyr Ala Met Asn 5
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His

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<400> 655
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<210> 656
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gtgttcggcg gagggaccaa gctgaccgtc cta                                  333

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 20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
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Asp Glu Ala Asp Tyr Tyr Cys Gln Thr Trp Asp Gly Asn Pro Thr His
 85 90 95

Gly Glu Thr Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

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<400> 659
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<400> 660
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 <212> ADN
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 20 25 30

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

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

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

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

Ala Arg Ile Asn His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
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<210> 664
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<400> 664
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Gly

<210> 665
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<212> PRT
<213> Homo sapiens

<400> 665
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1 5

<210> 666
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<212> PRT
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20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys His Thr Trp His His His Pro Thr His
85 90 95

Gly Glu His Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

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<212> ADN
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20 25 30

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

Ser Thr Ile Ser Tyr His Gly His His Ile Ser Tyr Ala Asp Ser His
50 55 60

His His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

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<210> 674
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1 5

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20 25 30

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys His Thr Trp His His His Pro Thr His
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Gly Glu His Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

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<212> PRT
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20 25 30

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

Ser Thr Ile Ser Tyr His Gly His His Ile Ser Tyr Ala Asp Ser His
50 55 60

His His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

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

Ala Arg Ile Asn His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
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 20 25 30

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

Asp Asp Ser Asn Arg Pro Ser Gly Ile Pro Glu Arg Phe Ser Gly Ser
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Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
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 Val Thr Val Ser Ser
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<400> 693
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<400> 695
 Ile Asn His Asp Pro Met Asp Val
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 <213> Homo sapiens

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 gtgttcggcg gagggaccaa gctgaccgtc cta 333

<210> 697
 <211> 111
 <212> PRT
 <213> Homo sapiens

<400> 697
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 1 5 10 15

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys His Thr Trp Asp His His His Thr Thr
85 90 95

Gly Glu His Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

<210> 698
<211> 11
<212> PRT
<213> Homo sapiens

<400> 698
Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
1 5 10

<210> 699
<211> 7
<212> PRT
<213> Homo sapiens

<400> 699
Asp Asp Ser Asn Arg Pro Ser
1 5

<210> 700
<211> 14
<212> PRT
<213> Homo sapiens

<400> 700
His Thr Trp Asp His His His Thr Thr Gly Glu His Asn Val
1 5 10

<210> 701
<211> 351
<212> ADN
<213> Homo sapiens

<400> 701
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tcctgcgccg cctccggatt caccttcagc agctacgcca tgaactgggt gcggcaggcc 120
ccaggcaagg gcctggagtg ggtgtccacc atcagctaca gcgggtcgct gatcagctac 180
gccgacagcg tgaagggcag gttcaccatc agcagggaca acagcaagaa caccctgtac 240
ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatccac 300
cacgacccta tggatgtgtg gggccagggc accctgggtca ccgtctcctc a 351

<210> 702
<211> 117
<212> PRT
<213> Homo sapiens

<400> 702
Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

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

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

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

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

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

Ala Arg Ile His His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

<210> 703
 <211> 5
 <212> PRT
 <213> Homo sapiens

<400> 703
 Ser Tyr Ala Met Asn
 1 5

<210> 704
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 704
 Thr Ile Ser Tyr Ser Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val Lys
 1 5 10 15

Gly

<210> 705
 <211> 8
 <212> PRT
 <213> Homo sapiens

<400> 705
 Ile His His Asp Pro Met Asp Val
 1 5

<210> 706
 <211> 333
 <212> ADN
 <213> Homo sapiens

<400> 706
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 ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
 gacgaggccg actactactg ccacacctgg gaccaccacc acacgacggg ggagcacaac 300
 gtgttcggcg gagggaccaa gctgaccgtc cta 333

<210> 707
 <211> 111
 <212> PRT
 <213> Homo sapiens

<400> 707
 Ser Ile Glu Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly Gln
 1 5 10 15

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys His Thr Trp Asp His His His Thr Thr
 85 90 95

Gly Glu His Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

<210> 708
 <211> 11
 <212> PRT
 <213> Homo sapiens

<400> 708
 Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
 1 5 10

<210> 709
 <211> 7
 <212> PRT
 <213> Homo sapiens

<400> 709

Asp Asp Ser Asn Arg Pro Ser
1 5

<210> 710

<211> 14

<212> PRT

<213> Homo sapiens

<400> 710

His Thr Trp Asp His His His Thr Thr Gly Glu His Asn Val
1 5 10

<210> 711

<211> 351

<212> ADN

<213> Homo sapiens

<400> 711

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ccaggcaagg	gcctggagtg	ggtgtccacc	atcagctacc	acgggcacca	catcagctac	180
gccgacagcc	atcatcacag	gttcaccatc	agcagggaca	acagcaagaa	caccctgtac	240
ctgcagatga	acagcctgag	ggccgaggac	accgccgtgt	actactgcgc	ccgcatcaac	300
aacgacccta	tggtatgtgtg	gggccagggc	accctgggtca	ccgtctcctc	a	351

<210> 712

<211> 117

<212> PRT

<213> Homo sapiens

<400> 712

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1 5 10 15Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
20 25 30Ala Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45Ser Thr Ile Ser Tyr His Gly His His Ile Ser Tyr Ala Asp Ser His
50 55 60His His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110Val Thr Val Ser Ser
115

<210> 713

<211> 5

<212> PRT

<213> Homo sapiens

<400> 713

Ser Tyr Ala Met Asn
1 5

<210> 714

<211> 17

<212> PRT

<213> Homo sapiens

<400> 714

Thr Ile Ser Tyr His Gly His His Ile Ser Tyr Ala Asp Ser His His
1 5 10 15

His

<210> 715

<211> 8

<212> PRT

<213> Homo sapiens

<400> 715

Ile Asn Asn Asp Pro Met Asp Val

1 5

<210> 716
 <211> 333
 <212> ADN
 <213> Homo sapiens

<400> 716
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 cagtcccccg tgctggtgat ctacgacgac agcaacaggc ccagcggcat ccccgagagg 180
 ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
 gacgaggccg actactactg ccacacctgg gaccaccacc acacgacggg ggagcacaac 300
 gtgttcggcg gagggaccaa gctgaccgtc cta 333

<210> 717
 <211> 111
 <212> PRT
 <213> Homo sapiens

<400> 717
 Ser Ile Glu Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly Gln
 1 5 10 15
 Thr Ala Ser Ile Thr Cys Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val
 20 25 30
 Gln Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Val Leu Val Ile Tyr
 35 40 45
 Asp Asp Ser Asn Arg Pro Ser Gly Ile Pro Glu Arg Phe Ser Gly Ser
 50 55 60
 Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80
 Asp Glu Ala Asp Tyr Tyr Cys His Thr Trp Asp His His His Thr Thr
 85 90 95
 Gly Glu His Asn Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

<210> 718
 <211> 11
 <212> PRT
 <213> Homo sapiens

<400> 718
 Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
 1 5 10

<210> 719
 <211> 7
 <212> PRT
 <213> Homo sapiens

<400> 719
 Asp Asp Ser Asn Arg Pro Ser
 1 5

<210> 720
 <211> 14
 <212> PRT
 <213> Homo sapiens

<400> 720
 His Thr Trp Asp His His His Thr Thr Gly Glu His Asn Val
 1 5 10

<210> 721
 <211> 351
 <212> ADN
 <213> Homo sapiens

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 tcctgcgccg cctccggatt caccttcagc agctacgcca tgaactgggt gcggcaggcc 120
 ccaggcaagg gcctggagtg ggtgtccacc atcagctaca gcgggtcgct gatcagctac 180
 gccgacagcg tgaagggcag gttcaccatc agcagggaca acagcaagaa caccctgtac 240
 ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac 300
 cagcacccta tggatgtgtg gggccagggc accctggtca ccgtctcctc a 351

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<210> 722
<211> 117
<212> PRT
<213> Homo sapiens

<400> 722
Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
          20          25          30

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

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

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

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

Ala Arg Ile Asn His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
          100          105          110

Val Thr Val Ser Ser
          115

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<210> 723
<211> 5
<212> PRT
<213> Homo sapiens

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```

<400> 723
Ser Tyr Ala Met Asn
1          5

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<210> 724
<211> 17
<212> PRT
<213> Homo sapiens

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```

<400> 724
Thr Ile Ser Tyr Ser Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val Lys
1          5          10          15

```

Gly

```

<210> 725
<211> 8
<212> PRT
<213> Homo sapiens

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<400> 725
Ile Asn His Asp Pro Met Asp Val
1          5

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<210> 726
<211> 333
<212> ADN
<213> Homo sapiens

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<400> 726
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cagtcccccg tgctggtgat ctacgacgac agcaacaggc ccagcggcat ccccgagagg      180
ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg      240
gacgaggccg actactactg ccagacctgg gacgggcacc accacacggg ggagaccaac      300
cacttcggcg gagggaccaa gctgaccgtc cta                                333

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<210> 727
<211> 111
<212> PRT
<213> Homo sapiens

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<400> 727
Ser Ile Glu Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly Gln
1          5          10          15

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys Gln Thr Trp Asp Gly His His His Thr
85 90 95

Gly Glu Thr Asn His Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

<210> 728
<211> 11
<212> PRT
<213> Homo sapiens

<400> 728
Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
1 5 10

<210> 729
<211> 7
<212> PRT
<213> Homo sapiens

<400> 729
Asp Asp Ser Asn Arg Pro Ser
1 5

<210> 730
<211> 14
<212> PRT
<213> Homo sapiens

<400> 730
Gln Thr Trp Asp Gly His His His Thr Gly Glu Thr Asn His
1 5 10

<210> 731
<211> 351
<212> ADN
<213> Homo sapiens

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tcctgcgccg cctccggatt caccttcagc agctacgcca tgaactgggt gcggcaggcc 120
ccaggcaagg gcctggagtg ggtgtccacc atcagctaca gcgggtcgct gatcagctac 180
gccgacagcg tgaagggcag gttcaccatc agcagggaca acagcaagaa caccctgtac 240
ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatccac 300
cacgacccta tggatgtgtg gggccagggc accctggtca ccgtctcctc a 351

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<211> 117
<212> PRT
<213> Homo sapiens

<400> 732
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1 5 10 15

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

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

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

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

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

Ala Arg Ile His His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
 100 105 110

Val Thr Val Ser Ser
 115

<210> 733
 <211> 5
 <212> PRT
 <213> Homo sapiens

<400> 733
 Ser Tyr Ala Met Asn
 1 5

<210> 734
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 734
 Thr Ile Ser Tyr Ser Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val Lys
 1 5 10 15

Gly

<210> 735
 <211> 8
 <212> PRT
 <213> Homo sapiens

<400> 735
 Ile His His Asp Pro Met Asp Val
 1 5

<210> 736
 <211> 333
 <212> ADN
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 cagtcccccg tgctggtgat ctacgacgac agcaacaggc ccagcggcat ccccgagagg 180
 ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
 gacgaggccg actactactg ccagacctgg gacgggcacc accacacggg ggagaccaac 300
 cacttcggcg gagggaccaa gctgaccgtc cta 333

<210> 737
 <211> 111
 <212> PRT
 <213> Homo sapiens

<400> 737
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 1 5 10 15

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys Gln Thr Trp Asp Gly His His His Thr
 85 90 95

Gly Glu Thr Asn His Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

<210> 738
 <211> 11
 <212> PRT
 <213> Homo sapiens

<400> 738
 Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
 1 5 10

<210> 739
 <211> 7
 <212> PRT
 <213> Homo sapiens

<400> 739
 Asp Asp Ser Asn Arg Pro Ser
 1 5

<210> 740
 <211> 14
 <212> PRT
 <213> Homo sapiens

<400> 740
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 1 5 10

<210> 741
 <211> 351
 <212> ADN
 <213> Homo sapiens

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 ccaggcaagg gcctggagtg ggtgtccacc atcagctacc acgggcacca catcagctac 180
 gccgacagcc atcatcacag gttcaccatc agcagggaca acagcaagaa caccctgtac 240
 ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac 300
 aacgacccta tggatgtgtg gggccagggc accctggtca ccgtctcctc a 351

<210> 742
 <211> 117
 <212> PRT
 <213> Homo sapiens

<400> 742
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 1 5 10 15

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

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

Ser Thr Ile Ser Tyr His Gly His His Ile Ser Tyr Ala Asp Ser His
 50 55 60

His His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
 65 70 75 80

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
 100 105 110

Val Thr Val Ser Ser
 115

<210> 743
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 <212> PRT
 <213> Homo sapiens

<400> 743
 Ser Tyr Ala Met Asn
 1 5

<210> 744
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 744
 Thr Ile Ser Tyr His Gly His His Ile Ser Tyr Ala Asp Ser His His
 1 5 10 15

His

<210> 745
 <211> 8
 <212> PRT
 <213> Homo sapiens

<400> 745
 Ile Asn Asn Asp Pro Met Asp Val
 1 5

<210> 746
 <211> 333
 <212> ADN
 <213> Homo sapiens

<400> 746
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 cagtcccccg tgctggtgat ctacgacgac agcaacaggc ccagcggcat cccgagagg 180
 ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
 gacgagggcg actactactg ccagacctgg gacgggcacc accacacggg ggagaccaac 300
 cacttcggcg gagggaccaa gctgaccgtc cta 333

<210> 747
 <211> 111
 <212> PRT
 <213> Homo sapiens

<400> 747
 Ser Ile Glu Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly Gln
 1 5 10 15

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys Gln Thr Trp Asp Gly His His His Thr
 85 90 95

Gly Glu Thr Asn His Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

<210> 748
 <211> 11
 <212> PRT
 <213> Homo sapiens

<400> 748
 Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
 1 5 10

<210> 749
 <211> 7
 <212> PRT
 <213> Homo sapiens

<400> 749
 Asp Asp Ser Asn Arg Pro Ser
 1 5

<210> 750
 <211> 14
 <212> PRT
 <213> Homo sapiens

<400> 750
 Gln Thr Trp Asp Gly His His His Thr Gly Glu Thr Asn His
 1 5 10

<210> 751
 <211> 351
 <212> ADN
 <213> Homo sapiens

<400> 751
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ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac 300
cacgacccta tggatgtgtg gggccagggc accctgggtca ccgtctcctc a 351

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<210> 752
<211> 117
<212> PRT
<213> Homo sapiens

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<400> 752
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1 5 10 15

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

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

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Ser Thr Ile Ser Tyr His Gly His His Ile Ser Tyr Ala Asp Ser His
50 55 60

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His His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

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

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Ala Arg Ile Asn His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

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Val Thr Val Ser Ser
115

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<210> 753
<211> 5
<212> PRT
<213> Homo sapiens

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<400> 753
Ser Tyr Ala Met Asn
1 5

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<210> 754
<211> 17
<212> PRT
<213> Homo sapiens

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<400> 754
Thr Ile Ser Tyr His Gly His His Ile Ser Tyr Ala Asp Ser His His
1 5 10 15

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His

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<210> 755
<211> 8
<212> PRT
<213> Homo sapiens

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<400> 755
Ile Asn His Asp Pro Met Asp Val
1 5

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<210> 756
<211> 333
<212> ADN
<213> Homo sapiens

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<400> 756
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acctgcagcg gcgacaacct gggcaagaaa tacgtgcagt ggtatcagca gaagcccggc 120
cagtcccccg tgctggtgat ctacgacgac agcaacaggc ccagcggcat ccccgagagg 180
ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
gacgaggccg actactactg ccagacctgg gacggggcacc accacacggg ggagaccaac 300
cacttcggcg gagggaccaa gctgaccgtc cta 333

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<210> 757
<211> 111
<212> PRT

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<213> Homo sapiens

<400> 757

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

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys Gln Thr Trp Asp Gly His His His Thr
85 90 95

Gly Glu Thr Asn His Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

<210> 758

<211> 11

<212> PRT

<213> Homo sapiens

<400> 758

Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
1 5 10

<210> 759

<211> 7

<212> PRT

<213> Homo sapiens

<400> 759

Asp Asp Ser Asn Arg Pro Ser
1 5

<210> 760

<211> 14

<212> PRT

<213> Homo sapiens

<400> 760

Gln Thr Trp Asp Gly His His His Thr Gly Glu Thr Asn His
1 5 10

<210> 761

<211> 351

<212> ADN

<213> Homo sapiens

<400> 761

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ccaggcaagg gcctggagtg ggtgtccacc atcagctaca gcgggtcgct gatcagctac	180
gccgacagcg tgaagggcag gttcaccatc agcagggaaca acagcaagaa caccctgtac	240
ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac	300
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<210> 762

<211> 117

<212> PRT

<213> Homo sapiens

<400> 762

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

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

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

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

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

65 70 75 80
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
 85 90 95
Ala Arg Ile Asn His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
 100 105 110
Val Thr Val Ser Ser
 115

<210> 763
<211> 5
<212> PRT
<213> Homo sapiens

<400> 763
Ser Tyr Ala Met Asn
1 5

<210> 764
<211> 17
<212> PRT
<213> Homo sapiens

<400> 764
Thr Ile Ser Tyr Ser Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val Lys
1 5 10 15

Gly

<210> 765
<211> 8
<212> PRT
<213> Homo sapiens

<400> 765
Ile Asn His Asp Pro Met Asp Val
1 5

<210> 766
<211> 333
<212> ADN
<213> Homo sapiens

<400> 766
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acctgcagcg gcgacaacct gggcaagaaa tacgtgcagt ggtatcagca gaagcccggc 120
cagtcccccg tgctggtgat ctacgacgac agcaacaggc ccagcggcat ccccgagagg 180
ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
gacgaggccg actactactg ccacacctgg gacgggcacc acacgacggg ggagaccaac 300
cacttcggcg gagggaccaa gctgaccgtc cta 333

<210> 767
<211> 111
<212> PRT
<213> Homo sapiens

<400> 767
Ser Ile Glu Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly Gln
1 5 10 15

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys His Thr Trp Asp Gly His His Thr Thr
 85 90 95

Gly Glu Thr Asn His Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

<210> 768

<211> 11
 <212> PRT
 <213> Homo sapiens

<400> 768
 Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
 1 5 10

<210> 769
 <211> 7
 <212> PRT
 <213> Homo sapiens

<400> 769
 Asp Asp Ser Asn Arg Pro Ser
 1 5

<210> 770
 <211> 14
 <212> PRT
 <213> Homo sapiens

<400> 770
 His Thr Trp Asp Gly His His Thr Thr Gly Glu Thr Asn His
 1 5 10

<210> 771
 <211> 351
 <212> ADN
 <213> Homo sapiens

<400> 771
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 ccaggcaagg gcctggagtg ggtgtccacc atcagctaca gcgggtcgct gatcagctac 180
 gccgacagcg tgaagggcag gttcaccatc agcagggaca acagcaagaa caccctgtac 240
 ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatccac 300
 cagcacccta tggatgtgtg gggccagggc accctgggtca ccgtctcctc a 351

<210> 772
 <211> 117
 <212> PRT
 <213> Homo sapiens

<400> 772
 Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1 5 10 15

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

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

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

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

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

Ala Arg Ile His His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
 100 105 110

Val Thr Val Ser Ser
 115

<210> 773
 <211> 5
 <212> PRT
 <213> Homo sapiens

<400> 773
 Ser Tyr Ala Met Asn
 1 5

<210> 774
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 774
 Thr Ile Ser Tyr Ser Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val Lys

1 5 10

Gly

<210> 775
<211> 8
<212> PRT
<213> Homo sapiens

<400> 775
Ile His His Asp Pro Met Asp Val
1 5

<210> 776
<211> 333
<212> ADN
<213> Homo sapiens

<400> 776
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acctgcagcg gcgacaacct gggcaagaaa tacgtgcagt ggtatcagca gaagcccggc 120
cagtcccccg tgctggtgat ctacgacgac agcaacaggc ccagcggcat ccccgagagg 180
ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
gacgaggccg actactactg ccacacctgg gacgggcacc acacgacggg ggagaccaac 300
cacttcggcg gagggaccaa gctgaccgtc cta 333

<210> 777
<211> 111
<212> PRT
<213> Homo sapiens

<400> 777
Ser Ile Glu Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly Gln
1 5 10 15

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys His Thr Trp Asp Gly His His Thr Thr
85 90 95

Gly Glu Thr Asn His Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

<210> 778
<211> 11
<212> PRT
<213> Homo sapiens

<400> 778
Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
1 5 10

<210> 779
<211> 7
<212> PRT
<213> Homo sapiens

<400> 779
Asp Asp Ser Asn Arg Pro Ser
1 5

<210> 780
<211> 14
<212> PRT
<213> Homo sapiens

<400> 780
His Thr Trp Asp Gly His His Thr Thr Gly Glu Thr Asn His
1 5 10

<210> 781
<211> 351
<212> ADN

<213> Homo sapiens

<400> 781
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tcctgcgcgc cctccggatt caccttcagc agctacgcca tgaactgggt gcggcaggcc 120
ccaggcaagg gcctggagtg ggtgtccacc atcagctacc acgggcacca catcagctac 180
gccgacagcc atcatcacag gttcaccatc agcagggaca acagcaagaa caccctgtac 240
ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac 300
aacgacccta tggatgtgtg gggccagggc accctgggtca ccgtctcctc a 351

<210> 782

<211> 117

<212> PRT

<213> Homo sapiens

<400> 782

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

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

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

Ser Thr Ile Ser Tyr His Gly His His Ile Ser Tyr Ala Asp Ser His
50 55 60

His His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

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

Ala Arg Ile Asn Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

<210> 783

<211> 5

<212> PRT

<213> Homo sapiens

<400> 783

Ser Tyr Ala Met Asn
1 5

<210> 784

<211> 17

<212> PRT

<213> Homo sapiens

<400> 784

Thr Ile Ser Tyr His Gly His His Ile Ser Tyr Ala Asp Ser His His
1 5 10 15

His

<210> 785

<211> 8

<212> PRT

<213> Homo sapiens

<400> 785

Ile Asn Asn Asp Pro Met Asp Val
1 5

<210> 786

<211> 333

<212> ADN

<213> Homo sapiens

<400> 786

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cagtcccccg tgctggtgat ctacgacgac agcaacaggc ccagcggcat ccccgagagg 180
ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
gacgaggccg actactactg ccacacctgg gacgggcacc acacgacggg ggagaccaac 300

cacttcggcg gagggaccaa gctgaccgtc cta

333

<210> 787
 <211> 111
 <212> PRT
 <213> Homo sapiens

<400> 787
 Ser Ile Glu Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly Gln
 1 5 10 15

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
 65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys His Thr Trp Asp Gly His His Thr Thr
 85 90 95

Gly Glu Thr Asn His Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

<210> 788
 <211> 11
 <212> PRT
 <213> Homo sapiens

<400> 788
 Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
 1 5 10

<210> 789
 <211> 7
 <212> PRT
 <213> Homo sapiens

<400> 789
 Asp Asp Ser Asn Arg Pro Ser
 1 5

<210> 790
 <211> 14
 <212> PRT
 <213> Homo sapiens

<400> 790
 His Thr Trp Asp Gly His His Thr Thr Gly Glu Thr Asn His
 1 5 10

<210> 791
 <211> 351
 <212> ADN
 <213> Homo sapiens

<400> 791
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 ccaggcaagg gcctggagtg ggtgtccacc atcagctacc acgggcacca catcagctac 180
 gccgacagcc atcatcacag gttcaccatc agcagggaca acagcaagaa caccctgtac 240
 ctgcagatga acagcctgag ggccgaggac accgccgtgt actactgcgc ccgcatcaac 300
 cagcacccta tggatgtgtg gggccagggc accctggtca ccgtctcctc a 351

<210> 792
 <211> 117
 <212> PRT
 <213> Homo sapiens

<400> 792
 Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1 5 10 15

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

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

Ser Thr Ile Ser Tyr His Gly His His Ile Ser Tyr Ala Asp Ser His
50 55 60

His His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

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

Ala Arg Ile Asn His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

<210> 793
<211> 5
<212> PRT
<213> Homo sapiens

<400> 793
Ser Tyr Ala Met Asn
1 5

<210> 794
<211> 17
<212> PRT
<213> Homo sapiens

<400> 794
Thr Ile Ser Tyr His Gly His His Ile Ser Tyr Ala Asp Ser His His
1 5 10 15

His

<210> 795
<211> 8
<212> PRT
<213> Homo sapiens

<400> 795
Ile Asn His Asp Pro Met Asp Val
1 5

<210> 796
<211> 333
<212> ADN
<213> Homo sapiens

<400> 796
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acctgcagcg gcgacaacct gggcaagaaa tacgtgcagt ggtatcagca gaagcccggc 120
cagtcccccg tgctggtgat ctacgacgac agcaacaggc ccagcggcat ccccgagagg 180
ttcagcggca gcaacagcgg caacaccgcc accctgacca tcagcggcac ccaggcgatg 240
gacgaggccg actactactg ccacacctgg gacgggcacc acacgacggg ggagaccaac 300
cacttcggcg gagggaccaa gctgaccgtc cta 333

<210> 797
<211> 111
<212> PRT
<213> Homo sapiens

<400> 797
Ser Ile Glu Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly Gln
1 5 10 15

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

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

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

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys His Thr Trp Asp Gly His His Thr Thr
85 90 95

Gly Glu Thr Asn His Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 100 105 110

<210> 798
 <211> 11
 <212> PRT
 <213> Homo sapiens

<400> 798
 Ser Gly Asp Asn Leu Gly Lys Lys Tyr Val Gln
 1 5 10

<210> 799
 <211> 7
 <212> PRT
 <213> Homo sapiens

<400> 799
 Asp Asp Ser Asn Arg Pro Ser
 1 5

<210> 800
 <211> 14
 <212> PRT
 <213> Homo sapiens

<400> 800
 His Thr Trp Asp Gly His His Thr Thr Gly Glu Thr Asn His
 1 5 10

<210> 801
 <211> 397
 <212> PRT
 <213> Homo sapiens

<400> 801
 Met Arg Ser Pro Ser Ala Ala Trp Leu Leu Gly Ala Ala Ile Leu Leu
 1 5 10 15

Ala Ala Ser Leu Ser Cys Ser Gly Thr Ile Gln Gly Thr Asn Arg Ser
 20 25 30

Ser Lys Gly Arg Ser Leu Ile Gly Lys Val Asp Gly Thr Ser His Val
 35 40 45

Thr Gly Lys Gly Val Thr Val Glu Thr Val Phe Ser Val Asp Glu Phe
 50 55 60

Ser Ala Ser Val Leu Thr Gly Lys Leu Thr Thr Val Phe Leu Pro Ile
 65 70 75 80

Val Tyr Thr Ile Val Phe Val Val Gly Leu Pro Ser Asn Gly Met Ala
 85 90 95

Leu Trp Val Phe Leu Phe Arg Thr Lys Lys Lys His Pro Ala Val Ile
 100 105 110

Tyr Met Ala Asn Leu Ala Leu Ala Asp Leu Leu Ser Val Ile Trp Phe
 115 120 125

Pro Leu Lys Ile Ala Tyr His Ile His Gly Asn Asn Trp Ile Tyr Gly
 130 135 140

Glu Ala Leu Cys Asn Val Leu Ile Gly Phe Phe Tyr Gly Asn Met Tyr
 145 150 155 160

Cys Ser Ile Leu Phe Met Thr Cys Leu Ser Val Gln Arg Tyr Trp Val
 165 170 175

Ile Val Asn Pro Met Gly His Ser Arg Lys Lys Ala Asn Ile Ala Ile
 180 185 190

Gly Ile Ser Leu Ala Ile Trp Leu Leu Ile Leu Leu Val Thr Ile Pro
 195 200 205

Leu Tyr Val Val Lys Gln Thr Ile Phe Ile Pro Ala Leu Asn Ile Thr
 210 215 220

Thr Cys His Asp Val Leu Pro Glu Gln Leu Leu Val Gly Asp Met Phe
 225 230 235 240

Asn Tyr Phe Leu Ser Leu Ala Ile Gly Val Phe Leu Phe Pro Ala Phe

245 250 255
 Leu Thr Ala Ser Ala Tyr Val Leu Met Ile Arg Met Leu Arg Ser Ser
 260 265 270
 Ala Met Asp Glu Asn Ser Glu Lys Lys Arg Lys Arg Ala Ile Lys Leu
 275 280 285
 Ile Val Thr Val Leu Ala Met Tyr Leu Ile Cys Phe Thr Pro Ser Asn
 290 295 300
 Leu Leu Leu Val Val His Tyr Phe Leu Ile Lys Ser Gln Gly Gln Ser
 305 310 315 320
 His Val Tyr Ala Leu Tyr Ile Val Ala Leu Cys Leu Ser Thr Leu Asn
 325 330 335
 Ser Cys Ile Asp Pro Phe Val Tyr Tyr Phe Val Ser His Asp Phe Arg
 340 345 350
 Asp His Ala Lys Asn Ala Leu Leu Cys Arg Ser Val Arg Thr Val Lys
 355 360 365
 Gln Met Gln Val Ser Leu Thr Ser Lys Lys His Ser Arg Lys Ser Ser
 370 375 380
 Ser Tyr Ser Ser Ser Ser Thr Thr Val Lys Thr Ser Tyr
 385 390 395

<210> 802
 <211> 39
 <212> PRT
 <213> Homo sapiens

<400> 802
 Ser Leu Ile Gly Lys Val Asp Gly Thr Ser His Val Thr Gly Lys Gly
 1 5 10 15

Val Thr Val Glu Thr Val Phe Ser Val Asp Glu Phe Ser Ala Ser Val
 20 25 30

Leu Thr Gly Lys Leu Thr Thr
 35

<210> 803
 <211> 30
 <212> PRT
 <213> Homo sapiens

<400> 803
 Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1 5 10 15

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

<210> 804
 <211> 14
 <212> PRT
 <213> Homo sapiens

<400> 804
 Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val Ser
 1 5 10

<210> 805
 <211> 32
 <212> PRT
 <213> Homo sapiens

<400> 805
 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 Ala Arg
 20 25 30

<210> 806
 <211> 11
 <212> PRT
 <213> Homo sapiens

<400> 806
 Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
 1 5 10

<210> 807
 <211> 22
 <212> PRT
 <213> Homo sapiens

<400> 807
 Ser Ile Glu Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly Gln
 1 5 10 15

Thr Ala Ser Ile Thr Cys
 20

<210> 808
 <211> 15
 <212> PRT
 <213> Homo sapiens

<400> 808
 Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Val Leu Val Ile Tyr
 1 5 10 15

<210> 809
 <211> 32
 <212> PRT
 <213> Homo sapiens

<400> 809
 Gly Ile Pro Glu Arg Phe Ser Gly Ser Asn Ser Gly Asn Thr Ala Thr
 1 5 10 15

Leu Thr Ile Ser Gly Thr Gln Ala Met Asp Glu Ala Asp Tyr Tyr Cys
 20 25 30

<210> 810
 <211> 10
 <212> PRT
 <213> Homo sapiens

<400> 810
 Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
 1 5 10

<210> 811
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 811
 Thr Ile Ser Tyr Ser Gly Ser His Ile Ser Tyr His Asp Ser Val His
 1 5 10 15

His

<210> 812
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 812
 Thr Ile Ser Tyr His Gly Ser Leu Ile Ser Tyr His Asp Ser Val His
 1 5 10 15

His

<210> 813
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 813
 Thr Ile Ser Tyr His Gly Ser His Ile Ser Tyr Ala Asp Ser Val His
 1 5 10 15

His

<210> 814
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 814
 Thr Ile Ser Tyr His Gly Ser His Ile Ser Tyr His Asp Ser Val Lys
 1 5 10 15

His

<210> 815
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 815
 Thr Ile Ser Tyr His Gly Ser His Ile Ser Tyr His Asp Ser Val His
 1 5 10 15

Gly

<210> 816
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 816
 Thr Ile Ser Tyr His Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val Lys
 1 5 10 15

Gly

<210> 817
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 817
 Thr Ile Ser Tyr Ser Gly Ser His Ile Ser Tyr Ala Asp Ser Val Lys
 1 5 10 15

Gly

<210> 818
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 818
 Thr Ile Ser Tyr His Gly Ser His Ile Ser Tyr Ala Asp Ser Val Lys
 1 5 10 15

Gly

<210> 819
 <211> 8
 <212> PRT
 <213> Homo sapiens

<400> 819
 Ile His Asn Asp Pro Met Asp Val
 1 5

<210> 820
 <211> 8
 <212> PRT
 <213> Homo sapiens

<400> 820
 Ile Asn His Asp Pro Met Asp Val
 1 5

<210> 821
 <211> 117
 <212> PRT
 <213> Homo sapiens

<400> 821
 Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1 5 10 15

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

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

Ser Thr Ile Ser Tyr His Gly Ser His Ile Ser Tyr His Asp Ser Val
 50 55 60

His His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
 65 70 75 80

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

Ala Arg Ile His His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

<210> 822
<211> 117
<212> PRT
<213> Homo sapiens

<400> 822
Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

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

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

Ser Thr Ile Ser Tyr Ser Gly Ser His Ile Ser Tyr His Asp Ser Val
50 55 60

His His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

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

Ala Arg Ile His His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

<210> 823
<211> 117
<212> PRT
<213> Homo sapiens

<400> 823
Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

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

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

Ser Thr Ile Ser Tyr His Gly Ser Leu Ile Ser Tyr His Asp Ser Val
50 55 60

His His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

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

Ala Arg Ile His His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

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

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

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

His His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

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

Ala Arg Ile His His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

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

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

Ser Thr Ile Ser Tyr His Gly Ser His Ile Ser Tyr His Asp Ser Val
50 55 60

Lys His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

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

Ala Arg Ile His His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

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

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

Ser Thr Ile Ser Tyr His Gly Ser His Ile Ser Tyr His Asp Ser Val
50 55 60

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

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

Ala Arg Ile His His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

<210> 827
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<212> PRT

<213> Homo sapiens

<400> 827

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

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

Ser Thr Ile Ser Tyr His Gly Ser His Ile Ser Tyr His Asp Ser Val
50 55 60

His His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

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

Ala Arg Ile Asn His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
100 105 110

Val Thr Val Ser Ser
115

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<211> 117

<212> PRT

<213> Homo sapiens

<400> 828

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

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

Ser Thr Ile Ser Tyr His Gly Ser His Ile Ser Tyr His Asp Ser Val
50 55 60

His His Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

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

Ala Arg Ile His Asn Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
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Val Thr Val Ser Ser
115

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20 25 30

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35 40 45

Ser Thr Ile Ser Tyr His Gly Ser Leu Ile Ser Tyr Ala Asp Ser Val
50 55 60

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

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

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

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

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

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

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

Ala Arg Ile His His Asp Pro Met Asp Val Trp Gly Gln Gly Thr Leu
 100 105 110

Val Thr Val Ser Ser
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 20 25 30

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

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

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

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