SEQUENCE LISTING

<110> Inviragen, Inc.
THE GOVERNMENT OF THE UNITED STATES OF AMERICA AS REPRESENTED BY THE SECRETARY OF THE DEPARTMENT OF HEALTH AND HUMAN SERVICES
Stinchcomb, Dan
Huang, Claire Y.
Kinney, Richard M.
Livengood, Jill A.

<120> COMPOSITIONS AND METHODS FOR DENGUE VIRUS CHIMERIC CONSTRUCTS IN VACCINES

<130> 478484.85

<150> 61/800,204
<151> 2013-03-15

<160> 12

<170> PatentIn version 3.5

<210> 1
<211> 10723
<212> DNA
<213> Artificial Sequence

<220>
<223> Dengue virus serotype 1, BVS

<400> 1
agttgttagt ctacgtggac cgacaaagac agattctttg agggagctaa gctcaatgta 60

gttctaacag ttttttaatt agagagcaga tctctgatga ataaccaacg gaaaaaggcg 120

aaaaacacgc ctttcaatat gctgaaacgc gagagaaacc gcgtgtcgac tgtgcaacag 180

ctgacaaaga gattctcact tggaatgctg cagggacgag gaccattaaa actgttcatg 240

gccctggtgg cgttccttcg tttcctaaca atcccaccaa cagcagggat attgaagaga 300

tggggaacaa ttaaaaaatc aaaagctatt aatgttttga gagggttcag gaaagagatt 360

ggaaggatgc tgaacatctt gaataggaga cgcagatctg caggcatgat cattatgctg 420

attccaacag tgatggcgtt ccatttaacc acgcgtgggg gagagccgca tatgatagtt 480

agcaagcagg aaagaggaaa gtcacttttg ttcaagacct ctgcaggtgt caacatgtgc 540

accctcattg cgatggattt gggagagttg tgtgaggaca cgatgaccta caaatgcccc 600

cggatcactg aggcggaacc agatgacgtt gactgttggt gcaatgccac ggacacatgg 660

gtgacctatg gaacgtgctc tcaaactggc gaacaccgac gagacaaacg ttccgtcgca 720

ttggccccac acgtggggct tggcctagaa acaagagccg aaacgtggat gtcctctgaa 780

ggtgcttgga aacagataca aaaagtagag acttgggctc tgagacatcc aggattcacg 840

gtgatagccc tttttctagc acatgccata ggaacatcca tcacccagaa agggatcatt 900

ttcattttgc tgatgctggt aacaccatct atggccatgc gatgcgtggg aataggcaac 960

agagacttcg tggaaggact gtcaggagca acatgggtgg atgtggtact ggagcatgga 1020

agttgcgtca ccaccatggc aaaaaacaaa ccaacactgg acattgaact cttgaagacg 1080

gaggtcacaa accctgcagt tctgcgtaaa ttgtgcattg aagctaaaat atcaaacacc 1140

accaccgatt cgagatgtcc aacacaagga gaagccacac tggtggaaga acaagacgcg 1200

aactttgtgt gccgacgaac gttcgtggac agaggctggg gcaatggctg tgggctattc 1260

ggaaaaggta gtctaataac gtgtgccaag tttaagtgtg tgacaaaact agaaggaaag 1320

atagttcaat atgaaaacct aaaatattca gtgatagtca ccgtccacac tggagatcag 1380

caccaggtgg gaaatgagac tacagaacat ggaacaactg caaccataac acctcaagct 1440

cctacgtcgg aaatacagct gaccgactac ggaaccctta cattagattg ttcacctagg 1500

acagggctag attttaacga gatggtgttg ctgacaatga aagaaagatc atggcttgtc 1560

cacaaacaat ggttcctaga cttaccactg ccttggacct ctggggcttc aacatcccaa 1620

gagacttgga acagacaaga tttactggtc acatttaaga cagctcatgc aaagaagcag 1680

gaagtagtcg tactaggatc acaagaagga gcaatgcaca ctgcgctgac tggagcgaca 1740

gaaatccaaa cgtcaggaac gacaacaatt ttcgcaggac acctaaaatg cagactaaaa 1800

atggacaaac taactttaaa agggatgtca tatgtgatgt gcacaggctc attcaagtta 1860

gagaaagaag tggctgagac ccagcatgga actgttctgg tgcaggttaa atatgaagga 1920

acagacgcac catgcaagat tccctttcg acccaagatg agaaaggagc aacccagaat 1980

gggagattaa taacagccaa ccccatagtc actgacaaag aaaaaccagt caatattgag 2040

gcagaaccac cctttggtga gagctacatc gtggtaggag caggtgaaaa agctttgaaa 2100

ctaagctggt tcaagaaagg aagcagcata gggaaaatgt ttgaagcaac tgcccgagga 2160

gcacgaagga tggccattct gggagacacc gcatgggact tcggttctat aggaggagtg 2220

ttcacgtcta tgggaaaact ggtacaccag gtttttggaa ctgcatatgg agttttgttt 2280

agcggagttt cttggaccat gaaaatagga atagggattc tgctgacatg gctaggatta 2340

aattcaagga acacgtccct ttcgatgatg tgcatcgcag ccgccattgt gacactgtat 2400

ttgggagtca tggtgcaggc cgatagtggt tgcgttgtga gctggaaaaa caaagaactg 2460

aaatgtggca gtgggatttt catcacagac aacgtgcaca catggacaga acaatacaag 2520

ttccaaccag aatccccttc aaaactagct tcagctatcc agaaagccca tgaagaggac 2580

atttgtggaa tccgctcagt aacaagactg gagaatctga tgtggaaaca aataacacca 2640

gaattgaatc acattctatc agaaaatgag gtgaagttaa ctattatgac aggagacatc 2700

aaaggaatca tgcaggcagg aaaacgatct ctgcggcctc agcccactga gctgaagtat 2760

tcatggaaaa catggggcaa agcaaaaatg ctctctacag agtctcataa ccagaccttt 2820

ctcattgatg gccccgaaac agcagaatgc cccaacacaa atagagcttg gaattcgttg 2880

gaagttgaag actatggctt tggagtattc accaccaata tatggctaaa attgaaagaa 2940

aaacaggatg tattctgcga ctcaaaactc atgtcagcgg ccataaaaga caacagagcc 3000

gtccatgccg atatgggtta ttggatagaa agtgcactca atgacacatg gaagatagag 3060

aaagcctctt tcattgaagt taaaaactgc cactggccaa aatcacacac cctctggagc 3120

aatggagtgc tagaaagtga gatgataatt ccaaagaatc tcgctggacc agtgtctcaa 3180

cacaactata gaccaggcta ccatacacaa ataacaggac catggcatct aggtaagctt 3240

gagatggact ttgatttctg tgatggaaca acagtggtag tgactgagga ctgcggaaat 3300

agaggaccct ctttgagaac aaccactgcc tctggaaaac tcataacaga atggtgctgc 3360

cgatcttgca cattaccacc gctaagatac agaggtgagg atgggtgctg gtacgggatg 3420

gaaatcagac cattgaagga gaaagaagag aatttggtca actccttggt cacagctgga 3480

catgggcagg tcgacaactt ttcactagga gtcttgggaa tggcattgtt cctggaggaa 3540

atgcttagga cccgagtagg aacgaaacat gcaatactac tagttgcagt ttcttttgtg 3600

acattgatca cagggaacat gtcctttaga gacctgggaa gagtgatggt tatggtaggc 3660

gccactatga cggatgacat aggtatgggc gtgacttatc ttgccctact agcagccttc 3720

aaagtcagac caacttttgc agctggacta ctcttgagaa agctgacctc caaggaattg 3780

atgatgacta ctataggaat tgtactcctc tcccagagca ccctaccaga gaccattctt 3840

gagttgactg atgcgttagc cttaggcatg atggtcctca aaatggtgag aaatatggaa 3900

aagtatcaat tggcagtgac tatcatggct atcttgtgcg tcccaaacgc agtgatatta 3960

caaaacgcat ggaaagtgag ttgcacaata ttggcagtgg tgtccgtttc cccactgttc 4020

ttaacatcct cacagcaaaa aacagattgg ataccattag cattgacgat caaaggtctc 4080

aatccaacag ctatttttct aacaaccctc tcaagaacca gcaagaaaag gagctggcca 4140

ttaaatgagg ctatcatggc agtcgggatg gtgagcattt tagccagttc tctcctaaaa 4200

aatgatattc ccatgacagg accattagtg gctggagggc tcctcactgt gtgctacgtg 4260

ctcactggac gatcggccga tttggaactg gagagagcag ccgatgtcaa atgggaagac 4320

caggcagaga tatcaggaag cagtccaatc ctgtcaataa caatatcaga agatggtagc 4380

atgtcgataa aaaatgaaga ggaagatcaa acactgacca tactcattag aacaggattg 4440

ctggtgatct caggactttt tcctgtatca ataccaatca cggcagcagc atggtacctg 4500

tgggaagtga agaaacaacg ggccggagta ttgtgggatg ttccttcacc cccacccatg 4560

ggaaaggctg aactggaaga tggagcctat agaattaagc aaaaagggat tcttggatat 4620

tcccagatcg gagccggagt ttacaaagaa ggaacattcc atacaatgtg gcatgtcaca 4680

cgtggcgctg ttctaatgca taaaggaaag aggattgaac catcatgggc ggacgtcaag 4740

aaagacctaa tatcatatgg aggaggctgg aagttagaag gagaatggaa ggaaggagaa 4800

gaagtccagg tattggcact ggagcctgga aaaaatccaa gagccgtcca aacgaaacct 4860

ggtcttttca aaaccaacgc cggaacaata ggtgctgtat ctctggactt ttctcctgga 4920

acgtcaggat ctccaattat cgacaaaaaa ggaaaagttg tgggtcttta tggtaatggt 4980

gttgttacaa ggagtggagc atatgtgagt gctatagccc agactgaaaa aagcattgaa 5040

gacaacccag agatcgaaga tgacattttc cgaaagagaa gactgaccat catggacctc 5100

cacccaggag cgggaaagac gaagagatac cttccggcca tagtcagaga agctataaaa 5160

cggggtttga gaacattaat cttggccccc actagagttg tggcagctga aatggaggaa 5220

gcccttagag gacttccaat aagataccag accccagcca tcagagctgt gcacaccggg 5280

cgggagattg tggacctaat gtgtcatgcc acatttacca tgaggctgct atcaccagtt 5340

agagtgccaa actacaacct gattatcatg gacgaagccc atttcacaga cccagcaagt 5400

atagcagcta gaggatacat ctcaactcga gtggagatgg gtgaggcagc tgggattttt 5460

atgacagcca ctcccccggg aagcagagac ccatttcctc agagcaatgc accaatcata 5520

gatgaagaaa gagaaatccc tgaacgctcg tggaattccg gacatgaatg ggtcacggat 5580

tttaaaggga agactgtttg gttcgttcca agtataaaag caggaaatga tatagcagct 5640

tgcctgagga aaaatggaaa gaaagtgata caactcagta ggaagacctt tgattctgag 5700

tatgtcaaga ctagaaccaa tgattgggac ttcgtggtta caactgacat ttcagaaatg 5760

ggtgccaatt tcaaggctga gagggttata gaccccagac gctgcatgaa accagtcata 5820

ctaacagatg gtgaagagcg ggtgattctg gcaggaccta tgccagtgac ccactctagt 5880

gcagcacaaa gaagagggag aataggaaga aatccaaaaa atgagaatga ccagtacata 5940

tacatggggg aacctctgga aaatgatgaa gactgtgcac actggaaaga agctaaaatg 6000

ctcctagata acatcaacac gccagaagga atcattccta gcatgttcga accagagcgt 6060

gaaaaggtgg atgccattga tggcgaatac cgcttgagag gagaagcaag gaaaaccttt 6120

gtagacttaa tgagaagagg agacctacca gtctggttgg cctacagagt ggcagctgaa 6180

ggcatcaact acgcagacag aaggtggtgt tttgatggag tcaagaacaa ccaaatccta 6240

gaagaaaacg tggaagttga aatctggaca aaagaagggg aaaggaagaa attgaaaccc 6300

agatggttgg atgctaggat ctattctgac ccactggcgc taaaagaatt taaggaattt 6360

gcagccggaa gaaagtctct gaccctgaac ctaatcacag aaatgggtag gctcccaacc 6420

ttcatgactc agaaggcaag agacgcactg gacaacttag cagtgctgca cacggctgag 6480

gcaggtggaa gggcgtacaa ccatgctctc agtgaactgc cggagaccct ggagacattg 6540

cttttactga cacttctggc tacagtcacg ggagggatct ttttattctt gatgagcgca 6600

aggggcatag ggaagatgac cctgggaatg tgctgcataa tcacggctag catcctccta 6660

tggtacgcac aaatacagcc acactggata gcagcttcaa taatactgga gtttttttctc 6720

atagttttgc ttattccaga acctgaaaaa cagagaacac cccaagacaa ccaactgacc 6780

tacgttgtca tagccatcct cacagtggtg gccgcaacca tggcaaacga gatgggtttc 6840

ctagaaaaaa cgaagaaaga tctcggattg ggaagcattg caacccagca acccgagagc 6900

aacatcctgg acatagatct acgtcctgca tcagcatgga cgctgtatgc cgtggccaca 6960

acatttgtta caccaatgtt gagacatagc attgaaaatt cctcagtgaa tgtgtcccta 7020

acagctatag ccaaccaagc cacagtgtta atgggtctcg ggaaaggatg gccattgtca 7080

aagatggaca tcggagttcc ccttctcgcc attggatgct actcacaagt caaccccata 7140

actctcatag cagctctttt cttattggta gcacattatg ccatcatagg gccaggactc 7200

caagcaaaag caaccagaga agctcagaaa agagcagcgg cgggcatcat gaaaaaccca 7260

actgtcgatg gaataacagt gattgaccta gatccaatac cttatgatcc gaagtttgaa 7320

aagcagttgg gacaagtaat gctcctagtc ctctgcgtga ctcaagtatt gatgatgagg 7380

actacatggg ctctgtgtga ggctttaacc ttagctaccg ggcccatctc cacattgtgg 7440

gaaggaaatc cagggaggtt ttggaacact accattgcgg tgtcaatggc taacattttt 7500

agagggagtt acttggccgg agctggactt ctcttttcta ttatgaagaa cacaaccaac 7560

acaagaaggg gaactggcaa cataggagag acgcttggag agaaatggaa aagccgattg 7620

aacgcattgg gaaaaagtga attccagatc tacaagaaaa gtggaatcca ggaagtggat 7680

agaaccttag caaaagaagg cattaaaaga ggagaaacgg accatcacgc tgtgtcgcga 7740

ggctcagcaa aactgagatg gttcgttgag agaaacatgg tcacaccaga agggaaagta 7800

gtggacctcg gttgtggcag aggaggctgg tcatactatt gtggaggact aaagaatgta 7860

agagaagtca aaggcctaac aaaaggagga ccaggacacg aagaacccat ccccatgtca 7920

acatatgggt ggaatctagt gcgtcttcaa agtggagttg acgttttctt catcccgcca 7980

gaaaagtgtg acacattatt gtgtgacata ggggagtcat caccaaatcc cacagtggaa 8040

gcaggacgaa cactcagagt ccttaactta gtagaaaatt ggttgaacaa caacactcaa 8100

ttttgcataa aggttctcaa cccatatatg ccctcagtca tagaaaaaat ggaagcacta 8160

caaaggaaat atggaggagc cttagtgagg aatccactct cacgaaactc cacacatgag 8220

atgtactggg tatccaatgc ttccgggaac atagtgtcat cagtgaacat gatttcaagg 8280

atgttgatca acagatttac aatgagatac aagaaagcca cttacgagcc ggatgttgac 8340

ctcggaagcg gaacccgtaa catcgggatt gaaagtgaga taccaaacct agatataatt 8400

gggaaaagaa tagaaaaaat aaagcaagag catgaaacat catggcacta tgaccaagac 8460

cacccataca aaacgtgggc ataccatggt agctatgaaa caaaacagac tggatcagca 8520

tcatccatgg tcaacggagt ggtcaggctg ctgacaaaac cttgggacgt cgtccccatg 8580

gtgacacaga tggcaatgac agacacgact ccatttggac aacagcgcgt ttttaaagag 8640

aaagtggaca cgagaaccca agaaccgaaa gaaggcacga agaaactaat gaaaataaca 8700

gcagagtggc tttggaaaga attagggaag aaaaagacac ccaggatgtg caccagagaa 8760

gaattcacaa gaaaggtgag aagcaatgca gccttggggg ccatattcac tgatgagaac 8820

aagtggaagt cggcacgtga ggctgttgaa gatagtaggt tttgggagct ggttgacaag 8880

gaaaggaatc tccatcttga aggaaagtgt gaaacatgtg tgtacaacat gatgggaaaa 8940

agagagaaga agctagggga attcggcaag gcaaaaggca gcagagccat atggtacatg 9000

tggcttggag cacgcttctt agagtttgaa gccctaggat tcttaaatga agatcactgg 9060

ttctccagag agaactccct gagtggagtg gaaggagaag ggctgcacaa gctaggttac 9120

attctaagag acgtgagcaa gaaagaggga ggagcaatgt atgccgatga caccgcagga 9180

tgggatacaa gaatcacact agaagaccta aaaaatgaag aaatggtaac aaaccacatg 9240

gaaggagaac acaagaaact agccgaggcc attttcaaac taacgtacca aaacaaggtg 9300

gtgcgtgtgc aaagaccaac accaagaggc acagtaatgg acatcatatc gagaagagac 9360

caaagaggta gtggacaagt tggcacctat ggactcaata ctttcaccaa tatggaagcc 9420

caactaatca gacagatgga gggagaagga gtctttaaaa gcattcagca cctaacaatc 9480

acagaagaaa tcgctgtgca aaactggtta gcaagagtgg ggcgcgaaag gttatcaaga 9540

atggccatca gtggagatga ttgtgttgtg aaacctttag atgacaggtt cgcaagcgct 9600

ttaacagctc taaatgacat gggaaagatt aggaaagaca tacaacaatg ggaaccttca 9660

agaggatgga atgattggac acaagtgccc ttctgttcac accatttcca tgagttaatc 9720

atgaaagacg gtcgcgtact cgttgttcca tgtagaaacc aagatgaact gattggcaga 9780

gcccgaatct cccaaggagc agggtggtct ttgcgggaga cggcctgttt ggggaagtct 9840

tacgcccaaa tgtggagctt gatgtacttc cacagacgcg acctcaggct ggcggcaaat 9900

gctatttgct cggcagtacc atcacattgg gttccaacaa gtcgaacaac ctggtccata 9960

catgctaaac atgaatggat gacaacggaa gacatgctga cagtctggaa cagggtgtgg 10020

attcaagaaa acccatggat ggaagacaaa actccagtgg aatcatggga ggaaatccca 10080

tacttgggga aaagagaaga ccaatggtgc ggctcattga ttgggttaac aagcagggcc 10140

acctgggcaa agaacatcca agcagcaata aatcaagtta gatcccttat aggcaatgaa 10200

gaatacacag attacatgcc atccatgaaa agattcagaa gagaagagga agaagcagga 10260

gttctgtggt agaaagcaaa actaacatga aacaaggcta gaagtcaggt cggattaagc 10320

catagtacgg aaaaaactat gctacctgtg agccccgtcc aaggacgtta aaagaagtca 10380

ggccatcata aatgccatag cttgagtaaa ctatgcagcc tgtagctcca cctgagaagg 10440

tgtaaaaaat ccgggaggcc acaaaccatg gaagctgtac gcatggcgta gtggactagc 10500

ggttagagga gacccctccc ttacaaatcg cagcaacaat gggggcccaa ggcgagatga 10560

agctgtagtc tcgctggaag gactagaggt tagaggagac ccccccgaaa caaaaaacag 10620

catattgacg ctgggaaaga ccagagatcc tgctgtctcc tcagcatcat tccaggcaca 10680

gaacgccaga aaatggaatg gtgctgttga atcaacaggt tct 10723

<210> 2
<211> 3464
<212> PRT
<213> Artificial Sequence

<220>
<223> Dengue virus serotype 1, BVS

<400> 2

Met Asn Asn Gln Arg Lys Lys Ala Lys Asn Thr Pro Phe Asn Met Leu
1 5 10 15

Lys Arg Glu Arg Asn Arg Val Ser Thr Val Gln Gln Leu Thr Lys Arg
20 25 30

```
Phe Ser Leu Gly Met Leu Gln Gly Arg Gly Pro Leu Lys Leu Phe Met
        35                  40                  45

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

Ile Leu Lys Arg Trp Gly Thr Ile Lys Lys Ser Lys Ala Ile Asn Val
65                  70                  75                  80

Leu Arg Gly Phe Arg Lys Glu Ile Gly Arg Met Leu Asn Ile Leu Asn
                85                  90                  95

Arg Arg Arg Arg Ser Ala Gly Met Ile Ile Met Leu Ile Pro Thr Val
            100                 105                 110

Met Ala Phe His Leu Thr Thr Arg Gly Gly Glu Pro His Met Ile Val
        115                 120                 125

Ser Lys Gln Glu Arg Gly Lys Ser Leu Leu Phe Lys Thr Ser Ala Gly
    130                 135                 140

Val Asn Met Cys Thr Leu Ile Ala Met Asp Leu Gly Glu Leu Cys Glu
145                 150                 155                 160

Asp Thr Met Thr Tyr Lys Cys Pro Arg Ile Thr Glu Ala Glu Pro Asp
                165                 170                 175

Asp Val Asp Cys Trp Cys Asn Ala Thr Asp Thr Trp Val Thr Tyr Gly
            180                 185                 190

Thr Cys Ser Gln Thr Gly Glu His Arg Arg Asp Lys Arg Ser Val Ala
        195                 200                 205

Leu Ala Pro His Val Gly Leu Gly Leu Glu Thr Arg Ala Glu Thr Trp
    210                 215                 220

Met Ser Ser Glu Gly Ala Trp Lys Gln Ile Gln Lys Val Glu Thr Trp
225                 230                 235                 240

Ala Leu Arg His Pro Gly Phe Thr Val Ile Ala Leu Phe Leu Ala His
                245                 250                 255

Ala Ile Gly Thr Ser Ile Thr Gln Lys Gly Ile Ile Phe Ile Leu Leu
            260                 265                 270

Met Leu Val Thr Pro Ser Met Ala Met Arg Cys Val Gly Ile Gly Asn
        275                 280                 285

Arg Asp Phe Val Glu Gly Leu Ser Gly Ala Thr Trp Val Asp Val Val
    290                 295                 300
```

```
Leu Glu His Gly Ser Cys Val Thr Thr Met Ala Lys Asn Lys Pro Thr
305                 310                 315                 320

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

Arg Lys Leu Cys Ile Glu Ala Lys Ile Ser Asn Thr Thr Thr Asp Ser
            340                 345                 350

Arg Cys Pro Thr Gln Gly Glu Ala Thr Leu Val Glu Glu Gln Asp Ala
        355                 360                 365

Asn Phe Val Cys Arg Arg Thr Phe Val Asp Arg Gly Trp Gly Asn Gly
    370                 375                 380

Cys Gly Leu Phe Gly Lys Gly Ser Leu Ile Thr Cys Ala Lys Phe Lys
385                 390                 395                 400

Cys Val Thr Lys Leu Glu Gly Lys Ile Val Gln Tyr Glu Asn Leu Lys
                405                 410                 415

Tyr Ser Val Ile Val Thr Val His Thr Gly Asp Gln His Gln Val Gly
            420                 425                 430

Asn Glu Thr Thr Glu His Gly Thr Thr Ala Thr Ile Thr Pro Gln Ala
        435                 440                 445

Pro Thr Ser Glu Ile Gln Leu Thr Asp Tyr Gly Thr Leu Thr Leu Asp
    450                 455                 460

Cys Ser Pro Arg Thr Gly Leu Asp Phe Asn Glu Met Val Leu Leu Thr
465                 470                 475                 480

Met Lys Glu Arg Ser Trp Leu Val His Lys Gln Trp Phe Leu Asp Leu
                485                 490                 495

Pro Leu Pro Trp Thr Ser Gly Ala Ser Thr Ser Gln Glu Thr Trp Asn
            500                 505                 510

Arg Gln Asp Leu Leu Val Thr Phe Lys Thr Ala His Ala Lys Lys Gln
        515                 520                 525

Glu Val Val Val Leu Gly Ser Gln Glu Gly Ala Met His Thr Ala Leu
    530                 535                 540

Thr Gly Ala Thr Glu Ile Gln Thr Ser Gly Thr Thr Thr Ile Phe Ala
545                 550                 555                 560

Gly His Leu Lys Cys Arg Leu Lys Met Asp Lys Leu Thr Leu Lys Gly
                565                 570                 575
```

```
Met Ser Tyr Val Met Cys Thr Gly Ser Phe Lys Leu Glu Lys Glu Val
            580                 585                 590

Ala Glu Thr Gln His Gly Thr Val Leu Val Gln Val Lys Tyr Glu Gly
        595                 600                 605

Thr Asp Ala Pro Cys Lys Ile Pro Phe Ser Thr Gln Asp Glu Lys Gly
    610                 615                 620

Ala Thr Gln Asn Gly Arg Leu Ile Thr Ala Asn Pro Ile Val Thr Asp
625                 630                 635                 640

Lys Glu Lys Pro Val Asn Ile Glu Ala Glu Pro Pro Phe Gly Glu Ser
                645                 650                 655

Tyr Ile Val Val Gly Ala Gly Glu Lys Ala Leu Lys Leu Ser Trp Phe
            660                 665                 670

Lys Lys Gly Ser Ser Ile Gly Lys Met Phe Glu Ala Thr Ala Arg Gly
        675                 680                 685

Ala Arg Arg Met Ala Ile Leu Gly Asp Thr Ala Trp Asp Phe Gly Ser
    690                 695                 700

Ile Gly Gly Val Phe Thr Ser Met Gly Lys Leu Val His Gln Val Phe
705                 710                 715                 720

Gly Thr Ala Tyr Gly Val Leu Phe Ser Gly Val Ser Trp Thr Met Lys
                725                 730                 735

Ile Gly Ile Gly Ile Leu Leu Thr Trp Leu Gly Leu Asn Ser Arg Asn
            740                 745                 750

Thr Ser Leu Ser Met Met Cys Ile Ala Ala Ala Ile Val Thr Leu Tyr
        755                 760                 765

Leu Gly Val Met Val Gln Ala Asp Ser Gly Cys Val Val Ser Trp Lys
    770                 775                 780

Asn Lys Glu Leu Lys Cys Gly Ser Gly Ile Phe Ile Thr Asp Asn Val
785                 790                 795                 800

His Thr Trp Thr Glu Gln Tyr Lys Phe Gln Pro Glu Ser Pro Ser Lys
                805                 810                 815

Leu Ala Ser Ala Ile Gln Lys Ala His Glu Glu Asp Ile Cys Gly Ile
            820                 825                 830

Arg Ser Val Thr Arg Leu Glu Asn Leu Met Trp Lys Gln Ile Thr Pro
        835                 840                 845
```

```
Glu Leu Asn His Ile Leu Ser Glu Asn Glu Val Lys Leu Thr Ile Met
    850                 855                 860

Thr Gly Asp Ile Lys Gly Ile Met Gln Ala Gly Lys Arg Ser Leu Arg
865                 870                 875                 880

Pro Gln Pro Thr Glu Leu Lys Tyr Ser Trp Lys Thr Trp Gly Lys Ala
                885                 890                 895

Lys Met Leu Ser Thr Glu Ser His Asn Gln Thr Phe Leu Ile Asp Gly
            900                 905                 910

Pro Glu Thr Ala Glu Cys Pro Asn Thr Asn Arg Ala Trp Asn Ser Leu
        915                 920                 925

Glu Val Glu Asp Tyr Gly Phe Gly Val Phe Thr Thr Asn Ile Trp Leu
    930                 935                 940

Lys Leu Lys Glu Lys Gln Asp Val Phe Cys Asp Ser Lys Leu Met Ser
945                 950                 955                 960

Ala Ala Ile Lys Asp Asn Arg Ala Val His Ala Asp Met Gly Tyr Trp
                965                 970                 975

Ile Glu Ser Ala Leu Asn Asp Thr Trp Lys Ile Glu Lys Ala Ser Phe
            980                 985                 990

Ile Glu Val Lys Asn Cys His Trp  Pro Lys Ser His Thr  Leu Trp Ser
        995                 1000                1005

Asn Gly  Val Leu Glu Ser Glu  Met Ile Ile Pro Lys  Asn Leu Ala
    1010                1015                1020

Gly Pro  Val Ser Gln His Asn  Tyr Arg Pro Gly Tyr  His Thr Gln
    1025                1030                1035

Ile Thr  Gly Pro Trp His Leu  Gly Lys Leu Glu Met  Asp Phe Asp
    1040                1045                1050

Phe Cys  Asp Gly Thr Thr Val  Val Val Thr Glu Asp  Cys Gly Asn
    1055                1060                1065

Arg Gly  Pro Ser Leu Arg Thr  Thr Thr Ala Ser Gly  Lys Leu Ile
    1070                1075                1080

Thr Glu  Trp Cys Cys Arg Ser  Cys Thr Leu Pro Pro  Leu Arg Tyr
    1085                1090                1095

Arg Gly  Glu Asp Gly Cys Trp  Tyr Gly Met Glu Ile  Arg Pro Leu
    1100                1105                1110
```

```
Lys Glu  Lys Glu Glu Asn Leu  Val Asn Ser Leu Val  Thr Ala Gly
    1115                 1120                 1125

His Gly  Gln Val Asp Asn Phe  Ser Leu Gly Val Leu  Gly Met Ala
    1130                 1135                 1140

Leu Phe  Leu Glu Glu Met Leu  Arg Thr Arg Val Gly  Thr Lys His
    1145                 1150                 1155

Ala Ile  Leu Leu Val Ala Val  Ser Phe Val Thr Leu  Ile Thr Gly
    1160                 1165                 1170

Asn Met  Ser Phe Arg Asp Leu  Gly Arg Val Met Val  Met Val Gly
    1175                 1180                 1185

Ala Thr  Met Thr Asp Asp Ile  Gly Met Gly Val Thr  Tyr Leu Ala
    1190                 1195                 1200

Leu Leu  Ala Ala Phe Lys Val  Arg Pro Thr Phe Ala  Ala Gly Leu
    1205                 1210                 1215

Leu Leu  Arg Lys Leu Thr Ser  Lys Glu Leu Met Met  Thr Thr Ile
    1220                 1225                 1230

Gly Ile  Val Leu Leu Ser Gln  Ser Thr Leu Pro Glu  Thr Ile Leu
    1235                 1240                 1245

Glu Leu  Thr Asp Ala Leu Ala  Leu Gly Met Met Val  Leu Lys Met
    1250                 1255                 1260

Val Arg  Asn Met Glu Lys Tyr  Gln Leu Ala Val Thr  Ile Met Ala
    1265                 1270                 1275

Ile Leu  Cys Val Pro Asn Ala  Val Ile Leu Gln Asn  Ala Trp Lys
    1280                 1285                 1290

Val Ser  Cys Thr Ile Leu Ala  Val Val Ser Val Ser  Pro Leu Phe
    1295                 1300                 1305

Leu Thr  Ser Ser Gln Gln Lys  Thr Asp Trp Ile Pro  Leu Ala Leu
    1310                 1315                 1320

Thr Ile  Lys Gly Leu Asn Pro  Thr Ala Ile Phe Leu  Thr Thr Leu
    1325                 1330                 1335

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

Met Ala  Val Gly Met Val Ser  Ile Leu Ala Ser Ser  Leu Leu Lys
    1355                 1360                 1365
```

```
Asn Asp  Ile Pro Met Thr Gly  Pro Leu Val Ala Gly  Gly Leu Leu
    1370                 1375                 1380

Thr Val  Cys Tyr Val Leu Thr  Gly Arg Ser Ala Asp  Leu Glu Leu
    1385                 1390                 1395

Glu Arg  Ala Ala Asp Val Lys  Trp Glu Asp Gln Ala  Glu Ile Ser
    1400                 1405                 1410

Gly Ser  Ser Pro Ile Leu Ser  Ile Thr Ile Ser Glu  Asp Gly Ser
    1415                 1420                 1425

Met Ser  Ile Lys Asn Glu Glu  Glu Asp Gln Thr Leu  Thr Ile Leu
    1430                 1435                 1440

Ile Arg  Thr Gly Leu Leu Val  Ile Ser Gly Leu Phe  Pro Val Ser
    1445                 1450                 1455

Ile Pro  Ile Thr Ala Ala Ala  Trp Tyr Leu Trp Glu  Val Lys Lys
    1460                 1465                 1470

Gln Arg  Ala Gly Val Leu Trp  Asp Val Pro Ser Pro  Pro Pro Met
    1475                 1480                 1485

Gly Lys  Ala Glu Leu Glu Asp  Gly Ala Tyr Arg Ile  Lys Gln Lys
    1490                 1495                 1500

Gly Ile  Leu Gly Tyr Ser Gln  Ile Gly Ala Gly Val  Tyr Lys Glu
    1505                 1510                 1515

Gly Thr  Phe His Thr Met Trp  His Val Thr Arg Gly  Ala Val Leu
    1520                 1525                 1530

Met His  Lys Gly Lys Arg Ile  Glu Pro Ser Trp Ala  Asp Val Lys
    1535                 1540                 1545

Lys Asp  Leu Ile Ser Tyr Gly  Gly Gly Trp Lys Leu  Glu Gly Glu
    1550                 1555                 1560

Trp Lys  Glu Gly Glu Glu Val  Gln Val Leu Ala Leu  Glu Pro Gly
    1565                 1570                 1575

Lys Asn  Pro Arg Ala Val Gln  Thr Lys Pro Gly Leu  Phe Lys Thr
    1580                 1585                 1590

Asn Ala  Gly Thr Ile Gly Ala  Val Ser Leu Asp Phe  Ser Pro Gly
    1595                 1600                 1605

Thr Ser  Gly Ser Pro Ile Ile  Asp Lys Lys Gly Lys  Val Val Gly
    1610                 1615                 1620
```

```
Leu Tyr  Gly Asn Gly Val Val  Thr Arg Ser Gly Ala  Tyr Val Ser
    1625                1630                1635

Ala Ile  Ala Gln Thr Glu Lys  Ser Ile Glu Asp Asn  Pro Glu Ile
    1640                1645                1650

Glu Asp  Asp Ile Phe Arg Lys  Arg Arg Leu Thr Ile  Met Asp Leu
    1655                1660                1665

His Pro  Gly Ala Gly Lys Thr  Lys Arg Tyr Leu Pro  Ala Ile Val
    1670                1675                1680

Arg Glu  Ala Ile Lys Arg Gly  Leu Arg Thr Leu Ile  Leu Ala Pro
    1685                1690                1695

Thr Arg  Val Val Ala Ala Glu  Met Glu Glu Ala Leu  Arg Gly Leu
    1700                1705                1710

Pro Ile  Arg Tyr Gln Thr Pro  Ala Ile Arg Ala Val  His Thr Gly
    1715                1720                1725

Arg Glu  Ile Val Asp Leu Met  Cys His Ala Thr Phe  Thr Met Arg
    1730                1735                1740

Leu Leu  Ser Pro Val Arg Val  Pro Asn Tyr Asn Leu  Ile Ile Met
    1745                1750                1755

Asp Glu  Ala His Phe Thr Asp  Pro Ala Ser Ile Ala  Ala Arg Gly
    1760                1765                1770

Tyr Ile  Ser Thr Arg Val Glu  Met Gly Glu Ala Ala  Gly Ile Phe
    1775                1780                1785

Met Thr  Ala Thr Pro Pro Gly  Ser Arg Asp Pro Phe  Pro Gln Ser
    1790                1795                1800

Asn Ala  Pro Ile Ile Asp Glu  Glu Arg Glu Ile Pro  Glu Arg Ser
    1805                1810                1815

Trp Asn  Ser Gly His Glu Trp  Val Thr Asp Phe Lys  Gly Lys Thr
    1820                1825                1830

Val Trp  Phe Val Pro Ser Ile  Lys Ala Gly Asn Asp  Ile Ala Ala
    1835                1840                1845

Cys Leu  Arg Lys Asn Gly Lys  Lys Val Ile Gln Leu  Ser Arg Lys
    1850                1855                1860

Thr Phe  Asp Ser Glu Tyr Val  Lys Thr Arg Thr Asn  Asp Trp Asp
    1865                1870                1875
```

```
Phe Val  Val Thr Thr Asp Ile  Ser Glu Met Gly Ala  Asn Phe Lys
    1880                 1885                 1890

Ala Glu  Arg Val Ile Asp Pro  Arg Arg Cys Met Lys  Pro Val Ile
    1895                 1900                 1905

Leu Thr  Asp Gly Glu Glu Arg  Val Ile Leu Ala Gly  Pro Met Pro
    1910                 1915                 1920

Val Thr  His Ser Ser Ala Ala  Gln Arg Arg Gly Arg  Ile Gly Arg
    1925                 1930                 1935

Asn Pro  Lys Asn Glu Asn Asp  Gln Tyr Ile Tyr Met  Gly Glu Pro
    1940                 1945                 1950

Leu Glu  Asn Asp Glu Asp Cys  Ala His Trp Lys Glu  Ala Lys Met
    1955                 1960                 1965

Leu Leu  Asp Asn Ile Asn Thr  Pro Glu Gly Ile Ile  Pro Ser Met
    1970                 1975                 1980

Phe Glu  Pro Glu Arg Glu Lys  Val Asp Ala Ile Asp  Gly Glu Tyr
    1985                 1990                 1995

Arg Leu  Arg Gly Glu Ala Arg  Lys Thr Phe Val Asp  Leu Met Arg
    2000                 2005                 2010

Arg Gly  Asp Leu Pro Val Trp  Leu Ala Tyr Arg Val  Ala Ala Glu
    2015                 2020                 2025

Gly Ile  Asn Tyr Ala Asp Arg  Arg Trp Cys Phe Asp  Gly Val Lys
    2030                 2035                 2040

Asn Asn  Gln Ile Leu Glu Glu  Asn Val Glu Val Glu  Ile Trp Thr
    2045                 2050                 2055

Lys Glu  Gly Glu Arg Lys Lys  Leu Lys Pro Arg Trp  Leu Asp Ala
    2060                 2065                 2070

Arg Ile  Tyr Ser Asp Pro Leu  Ala Leu Lys Glu Phe  Lys Glu Phe
    2075                 2080                 2085

Ala Ala  Gly Arg Lys Ser Leu  Thr Leu Asn Leu Ile  Thr Glu Met
    2090                 2095                 2100

Gly Arg  Leu Pro Thr Phe Met  Thr Gln Lys Ala Arg  Asp Ala Leu
    2105                 2110                 2115

Asp Asn  Leu Ala Val Leu His  Thr Ala Glu Ala Gly  Gly Arg Ala
    2120                 2125                 2130
```

```
Tyr Asn  His Ala Leu Ser Glu  Leu Pro Glu Thr Leu  Glu Thr Leu
    2135                 2140                 2145

Leu Leu  Leu Thr Leu Leu Ala  Thr Val Thr Gly Gly  Ile Phe Leu
    2150                 2155                 2160

Phe Leu  Met Ser Ala Arg Gly  Ile Gly Lys Met Thr  Leu Gly Met
    2165                 2170                 2175

Cys Cys  Ile Ile Thr Ala Ser  Ile Leu Leu Trp Tyr  Ala Gln Ile
    2180                 2185                 2190

Gln Pro  His Trp Ile Ala Ala  Ser Ile Ile Leu Glu  Phe Phe Leu
    2195                 2200                 2205

Ile Val  Leu Leu Ile Pro Glu  Pro Glu Lys Gln Arg  Thr Pro Gln
    2210                 2215                 2220

Asp Asn  Gln Leu Thr Tyr Val  Val Ile Ala Ile Leu  Thr Val Val
    2225                 2230                 2235

Ala Ala  Thr Met Ala Asn Glu  Met Gly Phe Leu Glu  Lys Thr Lys
    2240                 2245                 2250

Lys Asp  Leu Gly Leu Gly Ser  Ile Ala Thr Gln Gln  Pro Glu Ser
    2255                 2260                 2265

Asn Ile  Leu Asp Ile Asp Leu  Arg Pro Ala Ser Ala  Trp Thr Leu
    2270                 2275                 2280

Tyr Ala  Val Ala Thr Thr Phe  Val Thr Pro Met Leu  Arg His Ser
    2285                 2290                 2295

Ile Glu  Asn Ser Ser Val Asn  Val Ser Leu Thr Ala  Ile Ala Asn
    2300                 2305                 2310

Gln Ala  Thr Val Leu Met Gly  Leu Gly Lys Gly Trp  Pro Leu Ser
    2315                 2320                 2325

Lys Met  Asp Ile Gly Val Pro  Leu Leu Ala Ile Gly  Cys Tyr Ser
    2330                 2335                 2340

Gln Val  Asn Pro Ile Thr Leu  Ile Ala Ala Leu Phe  Leu Leu Val
    2345                 2350                 2355

Ala His  Tyr Ala Ile Ile Gly  Pro Gly Leu Gln Ala  Lys Ala Thr
    2360                 2365                 2370

Arg Glu  Ala Gln Lys Arg Ala  Ala Ala Gly Ile Met  Lys Asn Pro
    2375                 2380                 2385
```

```
Thr Val  Asp Gly Ile Thr Val  Ile Asp Leu Asp Pro  Ile Pro Tyr
    2390                 2395                 2400

Asp Pro  Lys Phe Glu Lys Gln  Leu Gly Gln Val Met  Leu Leu Val
    2405                 2410                 2415

Leu Cys  Val Thr Gln Val Leu  Met Met Arg Thr Thr  Trp Ala Leu
    2420                 2425                 2430

Cys Glu  Ala Leu Thr Leu Ala  Thr Gly Pro Ile Ser  Thr Leu Trp
    2435                 2440                 2445

Glu Gly  Asn Pro Gly Arg Phe  Trp Asn Thr Thr Ile  Ala Val Ser
    2450                 2455                 2460

Met Ala  Asn Ile Phe Arg Gly  Ser Tyr Leu Ala Gly  Ala Gly Leu
    2465                 2470                 2475

Leu Phe  Ser Ile Met Lys Asn  Thr Thr Asn Thr Arg  Arg Gly Thr
    2480                 2485                 2490

Gly Asn  Ile Gly Glu Thr Leu  Gly Glu Lys Trp Lys  Ser Arg Leu
    2495                 2500                 2505

Asn Ala  Leu Gly Lys Ser Glu  Phe Gln Ile Tyr Lys  Lys Ser Gly
    2510                 2515                 2520

Ile Gln  Glu Val Asp Arg Thr  Leu Ala Lys Glu Gly  Ile Lys Arg
    2525                 2530                 2535

Gly Glu  Thr Asp His His Ala  Val Ser Arg Gly Ser  Ala Lys Leu
    2540                 2545                 2550

Arg Trp  Phe Val Glu Arg Asn  Met Val Thr Pro Glu  Gly Lys Val
    2555                 2560                 2565

Val Asp  Leu Gly Cys Gly Arg  Gly Gly Trp Ser Tyr  Tyr Cys Gly
    2570                 2575                 2580

Gly Leu  Lys Asn Val Arg Glu  Val Lys Gly Leu Thr  Lys Gly Gly
    2585                 2590                 2595

Pro Gly  His Glu Glu Pro Ile  Pro Met Ser Thr Tyr  Gly Trp Asn
    2600                 2605                 2610

Leu Val  Arg Leu Gln Ser Gly  Val Asp Val Phe Phe  Ile Pro Pro
    2615                 2620                 2625

Glu Lys  Cys Asp Thr Leu Leu  Cys Asp Ile Gly Glu  Ser Ser Pro
    2630                 2635                 2640
```

```
Asn Pro  Thr Val Glu Ala Gly  Arg Thr Leu Arg Val  Leu Asn Leu
    2645                 2650                 2655

Val Glu  Asn Trp Leu Asn Asn  Asn Thr Gln Phe Cys  Ile Lys Val
    2660                 2665                 2670

Leu Asn  Pro Tyr Met Pro Ser  Val Ile Glu Lys Met  Glu Ala Leu
    2675                 2680                 2685

Gln Arg  Lys Tyr Gly Gly Ala  Leu Val Arg Asn Pro  Leu Ser Arg
    2690                 2695                 2700

Asn Ser  Thr His Glu Met Tyr  Trp Val Ser Asn Ala  Ser Gly Asn
    2705                 2710                 2715

Ile Val  Ser Ser Val Asn Met  Ile Ser Arg Met Leu  Ile Asn Arg
    2720                 2725                 2730

Phe Thr  Met Arg Tyr Lys Lys  Ala Thr Tyr Glu Pro  Asp Val Asp
    2735                 2740                 2745

Leu Gly  Ser Gly Thr Arg Asn  Ile Gly Ile Glu Ser  Glu Ile Pro
    2750                 2755                 2760

Asn Leu  Asp Ile Ile Gly Lys  Arg Ile Glu Lys Ile  Lys Gln Glu
    2765                 2770                 2775

His Glu  Thr Ser Trp His Tyr  Asp Gln Asp His Pro  Tyr Lys Thr
    2780                 2785                 2790

Trp Ala  Tyr His Gly Ser Tyr  Glu Thr Lys Gln Thr  Gly Ser Ala
    2795                 2800                 2805

Ser Ser  Met Val Asn Gly Val  Val Arg Leu Leu Thr  Lys Pro Trp
    2810                 2815                 2820

Asp Val  Val Pro Met Val Thr  Gln Met Ala Met Thr  Asp Thr Thr
    2825                 2830                 2835

Pro Phe  Gly Gln Gln Arg Val  Phe Lys Glu Lys Val  Asp Thr Arg
    2840                 2845                 2850

Thr Gln  Glu Pro Lys Glu Gly  Thr Lys Lys Leu Met  Lys Ile Thr
    2855                 2860                 2865

Ala Glu  Trp Leu Trp Lys Glu  Leu Gly Lys Lys Lys  Thr Pro Arg
    2870                 2875                 2880

Met Cys  Thr Arg Glu Glu Phe  Thr Arg Lys Val Arg  Ser Asn Ala
    2885                 2890                 2895
```

```
Ala Leu  Gly Ala Ile Phe Thr  Asp Glu Asn Lys Trp  Lys Ser Ala
    2900                 2905                 2910

Arg Glu  Ala Val Glu Asp Ser  Arg Phe Trp Glu Leu  Val Asp Lys
    2915                 2920                 2925

Glu Arg  Asn Leu His Leu Glu  Gly Lys Cys Glu Thr  Cys Val Tyr
    2930                 2935                 2940

Asn Met  Met Gly Lys Arg Glu  Lys Lys Leu Gly Glu  Phe Gly Lys
    2945                 2950                 2955

Ala Lys  Gly Ser Arg Ala Ile  Trp Tyr Met Trp Leu  Gly Ala Arg
    2960                 2965                 2970

Phe Leu  Glu Phe Glu Ala Leu  Gly Phe Leu Asn Glu  Asp His Trp
    2975                 2980                 2985

Phe Ser  Arg Glu Asn Ser Leu  Ser Gly Val Glu Gly  Glu Gly Leu
    2990                 2995                 3000

His Lys  Leu Gly Tyr Ile Leu  Arg Asp Val Ser Lys  Lys Glu Gly
    3005                 3010                 3015

Gly Ala  Met Tyr Ala Asp Asp  Thr Ala Gly Trp Asp  Thr Arg Ile
    3020                 3025                 3030

Thr Leu  Glu Asp Leu Lys Asn  Glu Glu Met Val Thr  Asn His Met
    3035                 3040                 3045

Glu Gly  Glu His Lys Lys Leu  Ala Glu Ala Ile Phe  Lys Leu Thr
    3050                 3055                 3060

Tyr Gln  Asn Lys Val Val Arg  Val Gln Arg Pro Thr  Pro Arg Gly
    3065                 3070                 3075

Thr Val  Met Asp Ile Ile Ser  Arg Arg Asp Gln Arg  Gly Ser Gly
    3080                 3085                 3090

Gln Val  Gly Thr Tyr Gly Leu  Asn Thr Phe Thr Asn  Met Glu Ala
    3095                 3100                 3105

Gln Leu  Ile Arg Gln Met Glu  Gly Glu Gly Val Phe  Lys Ser Ile
    3110                 3115                 3120

Gln His  Leu Thr Ile Thr Glu  Glu Ile Ala Val Gln  Asn Trp Leu
    3125                 3130                 3135

Ala Arg  Val Gly Arg Glu Arg  Leu Ser Arg Met Ala  Ile Ser Gly
    3140                 3145                 3150
```

```
Asp Asp  Cys Val Val Lys Pro  Leu Asp Asp Arg Phe  Ala Ser Ala
    3155                 3160                 3165

Leu Thr  Ala Leu Asn Asp Met  Gly Lys Ile Arg Lys  Asp Ile Gln
    3170                 3175                 3180

Gln Trp  Glu Pro Ser Arg Gly  Trp Asn Asp Trp Thr  Gln Val Pro
    3185                 3190                 3195

Phe Cys  Ser His His Phe His  Glu Leu Ile Met Lys  Asp Gly Arg
    3200                 3205                 3210

Val Leu  Val Val Pro Cys Arg  Asn Gln Asp Glu Leu  Ile Gly Arg
    3215                 3220                 3225

Ala Arg  Ile Ser Gln Gly Ala  Gly Trp Ser Leu Arg  Glu Thr Ala
    3230                 3235                 3240

Cys Leu  Gly Lys Ser Tyr Ala  Gln Met Trp Ser Leu  Met Tyr Phe
    3245                 3250                 3255

His Arg  Arg Asp Leu Arg Leu  Ala Ala Asn Ala Ile  Cys Ser Ala
    3260                 3265                 3270

Val Pro  Ser His Trp Val Pro  Thr Ser Arg Thr Thr  Trp Ser Ile
    3275                 3280                 3285

His Ala  Lys His Glu Trp Met  Thr Thr Glu Asp Met  Leu Thr Val
    3290                 3295                 3300

Trp Asn  Arg Val Trp Ile Gln  Glu Asn Pro Trp Met  Glu Asp Lys
    3305                 3310                 3315

Thr Pro  Val Glu Ser Trp Glu  Glu Ile Pro Tyr Leu  Gly Lys Arg
    3320                 3325                 3330

Glu Asp  Gln Trp Cys Gly Ser  Leu Ile Gly Leu Thr  Ser Arg Ala
    3335                 3340                 3345

Thr Trp  Ala Lys Asn Ile Gln  Ala Ala Ile Asn Gln  Val Arg Ser
    3350                 3355                 3360

Leu Ile  Gly Asn Glu Glu Tyr  Thr Asp Tyr Met Pro  Ser Met Lys
    3365                 3370                 3375

Arg Phe  Arg Arg Glu Glu Glu  Glu Ala Gly Val Leu  Trp Lys Ala
    3380                 3385                 3390

Lys Leu  Thr Asn Lys Ala Arg  Ser Gln Val Gly Leu  Ser His Ser
    3395                 3400                 3405
```

Thr Glu Lys Thr Met Leu Pro Val Ser Pro Val Gln Gly Arg Lys
3410 3415 3420

Lys Ser Gly His His Lys Cys His Ser Leu Ser Lys Leu Cys Ser
3425 3430 3435

Leu Leu His Leu Arg Arg Cys Lys Lys Ser Gly Arg Pro Gln Thr
3440 3445 3450

Met Glu Ala Val Arg Met Ala Trp Thr Ser Gly
3455 3460

<210> 3
<211> 3391
<212> PRT
<213> Artificial Sequence

<220>
<223> Dengue virus serotype 1, MVS

<400> 3

Met Asn Asn Gln Arg Lys Lys Ala Lys Asn Thr Pro Phe Asn Met Leu
1 5 10 15

Lys Arg Glu Arg Asn Arg Val Ser Thr Val Gln Gln Leu Thr Lys Arg
20 25 30

Phe Ser Leu Gly Met Leu Gln Gly Arg Gly Pro Leu Lys Leu Phe Met
35 40 45

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

Ile Leu Lys Arg Trp Gly Thr Ile Lys Lys Ser Lys Ala Ile Asn Val
65 70 75 80

Leu Arg Gly Phe Arg Lys Glu Ile Gly Arg Met Leu Asn Ile Leu Asn
85 90 95

Arg Arg Arg Arg Ser Ala Gly Met Ile Ile Met Leu Ile Pro Thr Val
100 105 110

Met Ala Phe His Leu Thr Thr Arg Gly Gly Glu Pro His Met Ile Val
115 120 125

Ser Lys Gln Glu Arg Gly Lys Ser Leu Leu Phe Lys Thr Ser Ala Gly
130 135 140

Val Asn Met Cys Thr Leu Ile Ala Met Asp Leu Gly Glu Leu Cys Glu
145 150 155 160

Asp Thr Met Thr Tyr Lys Cys Pro Arg Ile Thr Glu Ala Glu Pro Asp

```
                165                 170                 175

Asp Val Asp Cys Trp Cys Asn Ala Thr Asp Thr Trp Val Thr Tyr Gly
            180                 185                 190

Thr Cys Ser Gln Thr Gly Glu His Arg Arg Asp Lys Arg Ser Val Ala
        195                 200                 205

Leu Ala Pro His Val Gly Leu Gly Leu Glu Thr Arg Ala Glu Thr Trp
    210                 215                 220

Met Ser Ser Glu Gly Ala Trp Lys Gln Ile Gln Lys Val Glu Thr Trp
225                 230                 235                 240

Ala Leu Arg His Pro Gly Phe Thr Val Ile Ala Leu Phe Leu Ala His
                245                 250                 255

Ala Ile Gly Thr Ser Ile Thr Gln Lys Gly Ile Ile Phe Ile Leu Leu
            260                 265                 270

Met Leu Val Thr Pro Ser Met Ala Met Arg Cys Val Gly Ile Gly Asn
        275                 280                 285

Arg Asp Phe Val Glu Gly Leu Ser Gly Ala Thr Trp Val Asp Val Val
    290                 295                 300

Leu Glu His Gly Ser Cys Val Thr Thr Met Ala Lys Asn Lys Pro Thr
305                 310                 315                 320

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

Arg Lys Leu Cys Ile Glu Ala Lys Ile Ser Asn Thr Thr Thr Asp Ser
            340                 345                 350

Arg Cys Pro Thr Gln Gly Glu Ala Thr Leu Val Glu Glu Gln Asp Ala
        355                 360                 365

Asn Phe Val Cys Arg Arg Thr Phe Val Asp Arg Gly Trp Gly Asn Gly
    370                 375                 380

Cys Gly Leu Phe Gly Lys Gly Ser Leu Ile Thr Cys Ala Lys Phe Lys
385                 390                 395                 400

Cys Val Thr Lys Leu Glu Gly Lys Ile Val Gln Tyr Glu Asn Leu Lys
                405                 410                 415

Tyr Ser Val Ile Val Thr Val His Thr Gly Asp Gln His Gln Val Gly
            420                 425                 430

Asn Glu Thr Thr Glu His Gly Thr Thr Ala Thr Ile Thr Pro Gln Ala
```

435 440 445

Pro Thr Ser Glu Ile Gln Leu Thr Asp Tyr Gly Thr Leu Thr Leu Asp
450 455 460

Cys Ser Pro Arg Thr Gly Leu Asp Phe Asn Glu Met Val Leu Leu Thr
465 470 475 480

Met Lys Glu Arg Ser Trp Leu Val His Lys Gln Trp Phe Leu Asp Leu
485 490 495

Pro Leu Pro Trp Thr Ser Gly Ala Ser Thr Ser Gln Glu Thr Trp Asn
500 505 510

Arg Gln Asp Leu Leu Val Thr Phe Lys Thr Ala His Ala Lys Lys Gln
515 520 525

Glu Val Val Val Leu Gly Ser Gln Glu Gly Ala Met His Thr Ala Leu
530 535 540

Thr Gly Ala Thr Glu Ile Gln Thr Ser Gly Thr Thr Thr Ile Phe Ala
545 550 555 560

Gly His Leu Lys Cys Arg Leu Lys Met Asp Lys Leu Thr Leu Lys Gly
565 570 575

Met Ser Tyr Val Met Cys Thr Gly Ser Phe Lys Leu Glu Lys Glu Val
580 585 590

Ala Glu Thr Gln His Gly Thr Val Leu Val Gln Val Lys Tyr Glu Gly
595 600 605

Thr Asp Ala Pro Cys Lys Ile Pro Phe Ser Thr Gln Asp Glu Lys Gly
610 615 620

Ala Thr Gln Asn Gly Arg Leu Ile Thr Ala Asn Pro Ile Val Thr Asp
625 630 635 640

Lys Glu Lys Pro Val Asn Ile Glu Ala Glu Pro Pro Phe Gly Glu Ser
645 650 655

Tyr Ile Val Val Gly Ala Gly Glu Lys Ala Leu Lys Leu Ser Trp Phe
660 665 670

Lys Lys Gly Ser Ser Ile Gly Lys Met Phe Glu Ala Thr Ala Arg Gly
675 680 685

Ala Arg Arg Met Ala Ile Leu Gly Asp Thr Ala Trp Asp Phe Gly Ser
690 695 700

Ile Gly Gly Val Phe Thr Ser Met Gly Lys Leu Val His Gln Val Phe

705 710 715 720

Gly Thr Ala Tyr Gly Val Leu Phe Ser Gly Val Ser Trp Thr Met Lys
725 730 735

Ile Gly Ile Gly Ile Leu Leu Thr Trp Leu Gly Leu Asn Ser Arg Asn
740 745 750

Thr Ser Leu Ser Met Met Cys Ile Ala Ala Gly Ile Val Thr Leu Tyr
755 760 765

Leu Gly Val Met Val Gln Ala Asp Ser Gly Cys Val Val Ser Trp Lys
770 775 780

Asn Lys Glu Leu Lys Cys Gly Ser Gly Ile Phe Ile Thr Asp Asn Val
785 790 795 800

His Thr Trp Thr Glu Gln Tyr Lys Phe Gln Pro Glu Ser Pro Ser Lys
805 810 815

Leu Ala Ser Ala Ile Gln Lys Ala His Glu Glu Asp Ile Cys Gly Ile
820 825 830

Arg Ser Val Thr Arg Leu Glu Asn Leu Met Trp Lys Gln Ile Thr Pro
835 840 845

Glu Leu Asn His Ile Leu Ser Glu Asn Glu Val Lys Leu Thr Ile Met
850 855 860

Thr Gly Asp Ile Lys Gly Ile Met Gln Ala Gly Lys Arg Ser Leu Arg
865 870 875 880

Pro Gln Pro Thr Glu Leu Lys Tyr Ser Trp Lys Thr Trp Gly Lys Ala
885 890 895

Lys Met Leu Ser Thr Glu Ser His Asn Gln Thr Phe Leu Ile Asp Gly
900 905 910

Pro Glu Thr Ala Glu Cys Pro Asn Thr Asn Arg Ala Trp Asn Ser Leu
915 920 925

Glu Val Glu Asp Tyr Gly Phe Gly Val Phe Thr Thr Asn Ile Trp Leu
930 935 940

Lys Leu Lys Glu Lys Gln Asp Val Phe Cys Asp Ser Lys Leu Met Ser
945 950 955 960

Ala Ala Ile Lys Asp Asn Arg Ala Val His Ala Asp Met Gly Tyr Trp
965 970 975

Ile Glu Ser Ala Leu Asn Asp Thr Trp Lys Ile Glu Lys Ala Ser Phe

980 985 990

Ile Glu Val Lys Asn Cys His Trp Pro Lys Ser His Thr Leu Trp Ser
995 1000 1005

Asn Gly Val Leu Glu Ser Glu Met Ile Ile Pro Lys Asn Leu Ala
1010 1015 1020

Gly Pro Val Ser Gln His Asn Tyr Arg Pro Gly Tyr His Thr Gln
1025 1030 1035

Ile Thr Gly Pro Trp His Leu Gly Lys Leu Glu Met Asp Phe Asp
1040 1045 1050

Phe Cys Asp Gly Thr Thr Val Val Val Thr Glu Asp Cys Gly Asn
1055 1060 1065

Arg Gly Pro Ser Leu Arg Thr Thr Thr Ala Ser Gly Lys Leu Ile
1070 1075 1080

Thr Glu Trp Cys Cys Arg Ser Cys Thr Leu Pro Pro Leu Arg Tyr
1085 1090 1095

Arg Gly Glu Asp Gly Cys Trp Tyr Gly Met Glu Ile Arg Pro Leu
1100 1105 1110

Lys Glu Lys Glu Glu Asn Leu Val Asn Ser Leu Val Thr Ala Gly
1115 1120 1125

His Gly Gln Val Asp Asn Phe Ser Leu Gly Val Leu Gly Met Ala
1130 1135 1140

Leu Phe Leu Glu Glu Met Leu Arg Thr Arg Val Gly Thr Lys His
1145 1150 1155

Ala Ile Leu Leu Val Ala Val Ser Phe Val Thr Leu Ile Thr Gly
1160 1165 1170

Asn Met Ser Phe Arg Asp Leu Gly Arg Val Met Val Met Val Gly
1175 1180 1185

Ala Thr Met Thr Asp Asp Ile Gly Met Gly Val Thr Tyr Leu Ala
1190 1195 1200

Leu Leu Ala Ala Phe Lys Val Arg Pro Thr Phe Ala Ala Gly Leu
1205 1210 1215

Leu Leu Arg Lys Leu Thr Ser Lys Glu Leu Met Met Thr Thr Ile
1220 1225 1230

Gly Ile Val Leu Leu Ser Gln Ser Thr Leu Pro Glu Thr Ile Leu

1235 1240 1245

Glu Leu Thr Asp Ala Leu Ala Leu Gly Met Met Val Leu Lys Met
1250 1255 1260

Val Arg Asn Met Glu Lys Tyr Gln Leu Ala Val Thr Ile Met Ala
1265 1270 1275

Ile Leu Cys Val Pro Asn Ala Val Ile Leu Gln Asn Ala Trp Lys
1280 1285 1290

Val Ser Cys Thr Ile Leu Ala Val Val Ser Val Ser Pro Leu Phe
1295 1300 1305

Leu Thr Ser Ser Gln Gln Lys Thr Asp Trp Ile Pro Leu Ala Leu
1310 1315 1320

Thr Ile Lys Gly Leu Asn Pro Thr Ala Ile Phe Leu Thr Thr Leu
1325 1330 1335

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

Met Ala Val Gly Met Val Ser Ile Leu Ala Ser Ser Leu Leu Lys
1355 1360 1365

Asn Asp Ile Pro Met Thr Gly Pro Leu Val Ala Gly Gly Leu Leu
1370 1375 1380

Thr Val Cys Tyr Val Leu Thr Gly Arg Ser Ala Asp Leu Glu Leu
1385 1390 1395

Glu Arg Ala Ala Asp Val Lys Trp Glu Asp Gln Ala Glu Ile Ser
1400 1405 1410

Gly Ser Ser Pro Ile Leu Ser Ile Thr Ile Ser Glu Asp Gly Ser
1415 1420 1425

Met Ser Ile Lys Asn Glu Glu Glu Asp Gln Thr Leu Thr Ile Leu
1430 1435 1440

Ile Arg Thr Gly Leu Leu Val Ile Ser Gly Leu Phe Pro Val Ser
1445 1450 1455

Ile Pro Ile Thr Ala Ala Ala Trp Tyr Leu Trp Glu Val Lys Lys
1460 1465 1470

Gln Arg Ala Gly Val Leu Trp Asp Val Pro Ser Pro Pro Pro Met
1475 1480 1485

Gly Lys Ala Glu Leu Glu Asp Gly Ala Tyr Arg Ile Lys Gln Lys

1490 1495 1500

Gly Ile Leu Gly Tyr Ser Gln Ile Gly Ala Gly Val Tyr Lys Glu
1505 1510 1515

Gly Thr Phe His Thr Met Trp His Val Thr Arg Gly Ala Val Leu
1520 1525 1530

Met His Lys Gly Lys Arg Ile Glu Pro Ser Trp Ala Asp Val Lys
1535 1540 1545

Lys Asp Leu Ile Ser Tyr Gly Gly Gly Trp Lys Leu Glu Gly Glu
1550 1555 1560

Trp Lys Glu Gly Glu Glu Val Gln Val Leu Ala Leu Glu Pro Gly
1565 1570 1575

Lys Asn Pro Arg Ala Val Gln Thr Lys Pro Gly Leu Phe Lys Thr
1580 1585 1590

Asn Ala Gly Thr Ile Gly Ala Val Ser Leu Asp Phe Ser Pro Gly
1595 1600 1605

Thr Ser Gly Ser Pro Ile Ile Asp Lys Lys Gly Lys Val Val Gly
1610 1615 1620

Leu Tyr Gly Asn Gly Val Val Thr Arg Ser Gly Ala Tyr Val Ser
1625 1630 1635

Ala Ile Ala Gln Thr Glu Lys Ser Ile Glu Asp Asn Pro Glu Ile
1640 1645 1650

Glu Asp Asp Ile Phe Arg Lys Arg Arg Leu Thr Ile Met Asp Leu
1655 1660 1665

His Pro Gly Ala Gly Lys Thr Lys Arg Tyr Leu Pro Ala Ile Val
1670 1675 1680

Arg Glu Ala Ile Lys Arg Gly Leu Arg Thr Leu Ile Leu Ala Pro
1685 1690 1695

Thr Arg Val Val Ala Ala Glu Met Glu Glu Ala Leu Arg Gly Leu
1700 1705 1710

Pro Ile Arg Tyr Gln Thr Pro Ala Ile Arg Ala Val His Thr Gly
1715 1720 1725

Arg Glu Ile Val Asp Leu Met Cys His Ala Thr Phe Thr Met Arg
1730 1735 1740

Leu Leu Ser Pro Val Arg Val Pro Asn Tyr Asn Leu Ile Ile Met

1745 1750 1755

Asp Glu Ala His Phe Thr Asp Pro Ala Ser Ile Ala Ala Arg Gly
1760 1765 1770

Tyr Ile Ser Thr Arg Val Glu Met Gly Glu Ala Ala Gly Ile Phe
1775 1780 1785

Met Thr Ala Thr Pro Pro Gly Ser Arg Asp Pro Phe Pro Gln Ser
1790 1795 1800

Asn Ala Pro Ile Ile Asp Glu Glu Arg Glu Ile Pro Glu Arg Ser
1805 1810 1815

Trp Asn Ser Gly His Glu Trp Val Thr Asp Phe Lys Gly Lys Thr
1820 1825 1830

Val Trp Phe Val Pro Ser Ile Lys Ala Gly Asn Asp Ile Ala Ala
1835 1840 1845

Cys Leu Arg Lys Asn Gly Lys Lys Val Ile Gln Leu Ser Arg Lys
1850 1855 1860

Thr Phe Asp Ser Glu Tyr Val Lys Thr Arg Thr Asn Asp Trp Asp
1865 1870 1875

Phe Val Val Thr Thr Asp Ile Ser Glu Met Gly Ala Asn Phe Lys
1880 1885 1890

Ala Glu Arg Val Ile Asp Pro Arg Arg Cys Met Lys Pro Val Ile
1895 1900 1905

Leu Thr Asp Gly Glu Glu Arg Val Ile Leu Ala Gly Pro Met Pro
1910 1915 1920

Val Thr His Ser Ser Ala Ala Gln Arg Arg Gly Arg Ile Gly Arg
1925 1930 1935

Asn Pro Lys Asn Glu Asn Asp Gln Tyr Ile Tyr Met Gly Glu Pro
1940 1945 1950

Leu Glu Asn Asp Glu Asp Cys Ala His Trp Lys Glu Ala Lys Met
1955 1960 1965

Leu Leu Asp Asn Ile Asn Thr Pro Glu Gly Ile Ile Pro Ser Met
1970 1975 1980

Phe Glu Pro Glu Arg Glu Lys Val Asp Ala Ile Asp Gly Glu Tyr
1985 1990 1995

Arg Leu Arg Gly Glu Ala Arg Lys Thr Phe Val Asp Leu Met Arg

2000 2005 2010

Arg Gly Asp Leu Pro Val Trp Leu Ala Tyr Arg Val Ala Ala Glu
2015 2020 2025

Gly Ile Asn Tyr Ala Asp Arg Arg Trp Cys Phe Asp Gly Val Lys
2030 2035 2040

Asn Asn Gln Ile Leu Glu Glu Asn Val Glu Val Glu Ile Trp Thr
2045 2050 2055

Lys Glu Gly Glu Arg Lys Lys Leu Lys Pro Arg Trp Leu Asp Ala
2060 2065 2070

Arg Ile Tyr Ser Asp Pro Leu Ala Leu Lys Glu Phe Lys Glu Phe
2075 2080 2085

Ala Ala Gly Arg Lys Ser Leu Thr Leu Asn Leu Ile Thr Glu Met
2090 2095 2100

Gly Arg Leu Pro Thr Phe Met Thr Gln Lys Ala Arg Asp Ala Leu
2105 2110 2115

Asp Asn Leu Ala Val Leu His Thr Ala Glu Ala Gly Gly Arg Ala
2120 2125 2130

Tyr Asn His Ala Leu Ser Glu Leu Pro Glu Thr Leu Glu Thr Leu
2135 2140 2145

Leu Leu Leu Thr Leu Leu Ala Thr Val Thr Gly Gly Ile Phe Leu
2150 2155 2160

Phe Leu Met Ser Ala Arg Gly Ile Gly Lys Met Thr Leu Gly Met
2165 2170 2175

Cys Cys Ile Ile Thr Ala Ser Ile Leu Leu Trp Tyr Ala Gln Ile
2180 2185 2190

Gln Pro His Trp Ile Ala Ala Ser Ile Ile Leu Glu Phe Phe Leu
2195 2200 2205

Ile Val Leu Leu Ile Pro Glu Pro Glu Lys Gln Arg Thr Pro Gln
2210 2215 2220

Asp Asn Gln Leu Thr Tyr Val Val Ile Ala Ile Leu Thr Val Val
2225 2230 2235

Ala Ala Thr Met Ala Asn Glu Met Gly Phe Leu Glu Lys Thr Lys
2240 2245 2250

Lys Asp Leu Gly Leu Gly Ser Ile Ala Thr Gln Gln Pro Glu Ser

2255 2260 2265

Asn Ile Leu Asp Ile Asp Leu Arg Pro Ala Ser Ala Trp Thr Leu
2270 2275 2280

Tyr Ala Val Ala Thr Thr Phe Val Thr Pro Met Leu Arg His Ser
2285 2290 2295

Ile Glu Asn Ser Ser Val Asn Val Ser Leu Thr Ala Ile Ala Asn
2300 2305 2310

Gln Ala Thr Val Leu Met Gly Leu Gly Lys Gly Trp Pro Leu Ser
2315 2320 2325

Lys Met Asp Ile Gly Val Pro Leu Leu Ala Ile Gly Cys Tyr Ser
2330 2335 2340

Gln Val Asn Pro Ile Thr Leu Thr Ala Ala Leu Phe Leu Leu Val
2345 2350 2355

Ala His Tyr Ala Ile Ile Gly Pro Gly Leu Gln Ala Lys Ala Thr
2360 2365 2370

Arg Glu Ala Gln Lys Arg Ala Ala Ala Gly Ile Met Lys Asn Pro
2375 2380 2385

Thr Val Asp Gly Ile Thr Val Ile Asp Leu Asp Pro Ile Pro Tyr
2390 2395 2400

Asp Pro Lys Phe Glu Lys Gln Leu Gly Gln Val Met Leu Leu Val
2405 2410 2415

Leu Cys Val Thr Gln Val Leu Met Met Arg Thr Thr Trp Ala Leu
2420 2425 2430

Cys Glu Ala Leu Thr Leu Ala Thr Gly Pro Ile Ser Thr Leu Trp
2435 2440 2445

Glu Gly Asn Pro Gly Arg Phe Trp Asn Thr Thr Ile Ala Val Ser
2450 2455 2460

Met Ala Asn Ile Phe Arg Gly Ser Tyr Leu Ala Gly Ala Gly Leu
2465 2470 2475

Leu Phe Ser Ile Met Lys Asn Thr Thr Asn Thr Arg Arg Gly Thr
2480 2485 2490

Gly Asn Ile Gly Glu Thr Leu Gly Glu Lys Trp Lys Ser Arg Leu
2495 2500 2505

Asn Ala Leu Gly Lys Ser Glu Phe Gln Ile Tyr Lys Lys Ser Gly

2510 2515 2520

Ile Gln Glu Val Asp Arg Thr Leu Ala Lys Glu Gly Ile Lys Arg
2525 2530 2535

Gly Glu Thr Asp His His Ala Val Ser Arg Gly Ser Ala Lys Leu
2540 2545 2550

Arg Trp Phe Val Glu Arg Asn Met Val Thr Pro Glu Gly Lys Val
2555 2560 2565

Val Asp Leu Gly Cys Gly Arg Gly Gly Trp Ser Tyr Tyr Cys Gly
2570 2575 2580

Gly Leu Lys Asn Val Arg Glu Val Lys Gly Leu Thr Lys Gly Gly
2585 2590 2595

Pro Gly His Glu Glu Pro Ile Pro Met Ser Thr Tyr Gly Trp Asn
2600 2605 2610

Leu Val Arg Leu Gln Ser Gly Val Asp Val Phe Phe Ile Pro Pro
2615 2620 2625

Glu Lys Cys Asp Thr Leu Leu Cys Asp Ile Gly Glu Ser Ser Pro
2630 2635 2640

Asn Pro Thr Val Glu Ala Gly Arg Thr Leu Arg Val Leu Asn Leu
2645 2650 2655

Val Glu Asn Trp Leu Asn Asn Asn Thr Gln Phe Cys Ile Lys Val
2660 2665 2670

Leu Asn Pro Tyr Met Pro Ser Val Ile Glu Lys Met Glu Ala Leu
2675 2680 2685

Gln Arg Lys Tyr Gly Gly Ala Leu Val Arg Asn Pro Leu Ser Arg
2690 2695 2700

Asn Ser Thr His Glu Met Tyr Trp Val Ser Asn Ala Ser Gly Asn
2705 2710 2715

Ile Val Ser Ser Val Asn Met Ile Ser Arg Met Leu Ile Asn Arg
2720 2725 2730

Phe Thr Met Arg Tyr Lys Lys Ala Thr Tyr Glu Pro Asp Val Asp
2735 2740 2745

Leu Gly Ser Gly Thr Arg Asn Ile Gly Ile Glu Ser Glu Ile Pro
2750 2755 2760

Asn Leu Asp Ile Ile Gly Lys Arg Ile Glu Lys Ile Lys Gln Glu

2765 2770 2775

His Glu Thr Ser Trp His Tyr Asp Gln Asp His Pro Tyr Lys Thr
2780 2785 2790

Trp Ala Tyr His Gly Ser Tyr Glu Thr Lys Gln Thr Gly Ser Ala
2795 2800 2805

Ser Ser Met Val Asn Gly Val Val Arg Leu Leu Thr Lys Pro Trp
2810 2815 2820

Asp Val Val Pro Met Val Thr Gln Met Ala Met Thr Asp Thr Thr
2825 2830 2835

Pro Phe Gly Gln Gln Arg Val Phe Lys Glu Lys Val Asp Thr Arg
2840 2845 2850

Thr Gln Glu Pro Lys Glu Gly Thr Lys Lys Leu Met Lys Ile Thr
2855 2860 2865

Ala Glu Trp Leu Trp Lys Glu Leu Gly Lys Lys Lys Thr Pro Arg
2870 2875 2880

Met Cys Thr Arg Glu Glu Phe Thr Arg Lys Val Arg Ser Asn Ala
2885 2890 2895

Ala Leu Gly Ala Ile Phe Thr Asp Glu Asn Lys Trp Lys Ser Ala
2900 2905 2910

Arg Glu Ala Val Glu Asp Ser Arg Phe Trp Glu Leu Val Asp Lys
2915 2920 2925

Glu Arg Asn Leu His Leu Glu Gly Lys Cys Glu Thr Cys Val Tyr
2930 2935 2940

Asn Met Met Gly Lys Arg Glu Lys Lys Leu Gly Glu Phe Gly Lys
2945 2950 2955

Ala Lys Gly Ser Arg Ala Ile Trp Tyr Met Trp Leu Gly Ala Arg
2960 2965 2970

Phe Leu Glu Phe Glu Ala Leu Gly Phe Leu Asn Glu Asp His Trp
2975 2980 2985

Phe Ser Arg Glu Asn Ser Leu Ser Gly Val Glu Gly Glu Gly Leu
2990 2995 3000

His Lys Leu Gly Tyr Ile Leu Arg Asp Val Ser Lys Lys Glu Gly
3005 3010 3015

Gly Ala Met Tyr Ala Asp Asp Thr Ala Gly Trp Asp Thr Arg Ile

```
    3020                3025                3030

Thr Leu  Glu Asp Leu Lys Asn  Glu Glu Met Val Thr  Asn His Met
    3035                3040                3045

Glu Gly  Glu His Lys Lys Leu  Ala Glu Ala Ile Phe  Lys Leu Thr
    3050                3055                3060

Tyr Gln  Asn Lys Val Val Arg  Val Gln Arg Pro Thr  Pro Arg Gly
    3065                3070                3075

Thr Val  Met Asp Ile Ile Ser  Arg Arg Asp Gln Arg  Gly Ser Gly
    3080                3085                3090

Gln Val  Gly Thr Tyr Gly Leu  Asn Thr Phe Thr Asn  Met Glu Ala
    3095                3100                3105

Gln Leu  Ile Arg Gln Met Glu  Gly Glu Gly Val Phe  Lys Ser Ile
    3110                3115                3120

Gln His  Leu Thr Ile Thr Glu  Glu Ile Ala Val Gln  Asn Trp Leu
    3125                3130                3135

Ala Arg  Val Gly Arg Glu Arg  Leu Ser Arg Met Ala  Ile Ser Gly
    3140                3145                3150

Asp Asp  Cys Val Val Lys Pro  Leu Asp Asp Arg Phe  Ala Ser Ala
    3155                3160                3165

Leu Thr  Ala Leu Asn Asp Met  Gly Lys Ile Arg Lys  Asp Ile Gln
    3170                3175                3180

Gln Trp  Glu Pro Ser Arg Gly  Trp Asn Asp Trp Thr  Gln Val Pro
    3185                3190                3195

Phe Cys  Ser His His Phe His  Glu Leu Ile Met Lys  Asp Gly Arg
    3200                3205                3210

Val Leu  Val Val Pro Cys Arg  Asn Gln Asp Glu Leu  Ile Gly Arg
    3215                3220                3225

Ala Arg  Ile Ser Gln Gly Ala  Gly Trp Ser Leu Arg  Glu Thr Ala
    3230                3235                3240

Cys Leu  Gly Lys Ser Tyr Ala  Gln Met Trp Ser Leu  Met Tyr Phe
    3245                3250                3255

His Arg  Arg Asp Leu Arg Leu  Ala Ala Asn Ala Ile  Cys Ser Ala
    3260                3265                3270

Val Pro  Ser His Trp Val Pro  Thr Ser Arg Thr Thr  Trp Ser Ile
```

3275 3280 3285

His Ala Lys His Glu Trp Met Thr Thr Glu Asp Met Leu Thr Val
3290 3295 3300

Trp Asn Arg Val Trp Ile Gln Glu Asn Pro Trp Met Glu Asp Lys
3305 3310 3315

Thr Pro Val Glu Ser Trp Glu Glu Ile Pro Tyr Leu Gly Lys Arg
3320 3325 3330

Glu Asp Gln Trp Cys Gly Ser Leu Ile Gly Leu Thr Ser Arg Ala
3335 3340 3345

Thr Trp Ala Lys Asn Ile Gln Ala Ala Ile Asn Gln Val Arg Ser
3350 3355 3360

Leu Ile Gly Asn Glu Glu Tyr Thr Asp Tyr Met Pro Ser Met Lys
3365 3370 3375

Arg Phe Arg Arg Glu Glu Glu Glu Ala Gly Val Leu Trp
3380 3385 3390

<210> 4
<211> 10723
<212> DNA
<213> Artificial Sequence

<220>
<223> Dengue virus serotype 2, BVS

<400> 4
agttgttagt ctacgtggac cgacaaagac agattctttg agggagctaa gctcaatgta 60

gttctaacag ttttttaatt agagagcaga tctctgatga ataaccaacg gaaaaaggcg 120

aaaaacacgc ctttcaatat gctgaaacgc gagagaaacc gcgtgtcgac tgtgcaacag 180

ctgacaaaga gattctcact tggaatgctg cagggacgag gaccattaaa actgttcatg 240

gccctggtgg cgttccttcg tttcctaaca atcccaccaa cagcagggat attgaagaga 300

tggggaacaa ttaaaaaatc aaaagctatt aatgttttga gagggttcag gaaagagatt 360

ggaaggatgc tgaacatctt gaataggaga cgcagatctg caggcatgat cattatgctg 420

attccaacag tgatggcgtt ccatttaacc acacgtaacg gagaaccaca catgatcgtc 480

agcagacaag agaaagggaa aagtcttctg tttaaaacag aggttggcgt gaacatgtgt 540

accctcatgg ccatggacct tggtgaattg tgtgaagaca caatcacgta cgagtgtccc 600

cttctcaggc agaatgagcc agaagacata gactgttggt gcaactctac gtccacgtgg 660

gtaacttatg ggacgtgtac caccatggga gaacatagaa gagaaaaaag atcagtggca 720

ctcgttccac atgtgggaat gggactggag acacgaactg aaacatggat gtcatcagaa 780

ggggcctgga aacatgtcca gagaattgaa acttggatct tgagacatcc aggcttcacc 840

atgatggcag caatcctggc atacaccata ggaacgacac atttccaaag agccctgatc 900

ttcatcttac tgacagctgt cactccttca atgacaatgc gttgcatagg aatgtcaaat 960

agagactttg tggaaggggt ttcaggagga agctgggttg acatagtctt agaacatgga 1020

agctgtgtga cgacgatggc aaaaaacaaa ccaacattgg attttgaact gataaaaaca 1080

gaagccaaac agcctgccac cctaaggaag tactgtatag aggcaaagct aaccaacaca 1140

acaacagaat ctcgctgccc aacacaaggg gaacccagcc taaatgaaga gcaggacaaa 1200

aggttcgtct gcaaacactc catggtagac agaggatggg gaaatggatg tggactattt 1260

ggaaagggag gcattgtgac ctgtgctatg ttcagatgca aaaagaacat ggaaggaaaa 1320

gttgtgcaac cagaaaactt ggaatacacc attgtgataa cacctcactc aggggaagag 1380

catgcagtcg gaaatgacac aggaaaacat ggcaaggaaa tcaaaataac accacagagt 1440

tccatcacag aagcagaatt gacaggttat ggcactgtca caatggagtg ctctccaaga 1500

acgggcctcg acttcaatga gatggtgttg ctgcagatgg aaaataaagc ttggctggtg 1560

cacaggcaat ggttcctaga cctgccgtta ccatggttgc ccggagcgga cacacaaggg 1620

tcaaattgga tacagaaaga gacattggtc actttcaaaa atccccatgc gaagaaacag 1680

gatgttgttg ttttaggatc ccaagaaggg gccatgcaca cagcacttac aggggccaca 1740

gaaatccaaa tgtcatcagg aaacttactc ttcacaggac atctcaagtg caggctgaga 1800

atggacaagc tacagctcaa aggaatgtca tactctatgt gcacaggaaa gtttaaagtt 1860

gtgaaggaaa tagcagaaac acaacatgga acaatagtta tcagagtgca atatgaaggg 1920

gacggctctc catgcaagat ccctttgag ataatggatt tggaaaaaag acatgtctta 1980

ggtcgcctga ttacagtcaa cccaattgtg acagaaaaag atagcccagt caacatagaa 2040

gcagaacctc catttggaga cagctacatc atcataggag tagagccggg acaactgaag 2100

ctcaactggt ttaagaaagg aagttctatc ggccaaatgt ttgagacaac aatgaggggg 2160

gcgaagagaa tggccatttt aggtgacaca gcctgggatt ttggatcctt gggaggagtg 2220

tttacatcta taggaaaggc tctccaccaa gtctttggag caatctatgg agctgccttc 2280

agtggggttt catggactat gaaaatcctc ataggagtca ttatcacatg gataggaatg 2340

aattcacgca gcacctcact gtctgtgaca ctagtattgg tgggaattgt gacactgtat 2400

ttgggagtca tggtgcaggc cgatagtggt tgcgttgtga gctggaaaaa caaagaactg 2460

aaatgtggca gtgggatttt catcacagac aacgtgcaca catggacaga acaatacaag 2520

ttccaaccag aatccccttc aaaactagct tcagctatcc agaaagccca tgaagaggac 2580

atttgtggaa tccgctcagt aacaagactg gagaatctga tgtggaaaca aataacacca 2640

gaattgaatc acattctatc agaaaatgag gtgaagttaa ctattatgac aggagacatc 2700

aaaggaatca tgcaggcagg aaaacgatct ctgcggcctc agcccactga gctgaagtat 2760

tcatggaaaa catggggcaa agcaaaaatg ctctctacag agtctcataa ccagaccttt 2820

ctcattgatg gccccgaaac agcagaatgc cccaacacaa atagagcttg gaattcgttg 2880

gaagttgaag actatggctt tggagtattc accaccaata tatggctaaa attgaaagaa 2940

aaacaggatg tattctgcga ctcaaaactc atgtcagcgg ccataaaaga caacagagcc 3000

gtccatgccg atatgggtta ttggatagaa agtgcactca atgacacatg gaagatagag 3060

aaagcctctt tcattgaagt taaaaactgc cactggccaa aatcacacac cctctggagc 3120

aatggagtgc tagaaagtga gatgataatt ccaaagaatc tcgctggacc agtgtctcaa 3180

cacaactata gaccaggcta ccatacacaa ataacaggac catggcatct aggtaagctt 3240

gagatggact ttgatttctg tgatggaaca acagtggtag tgactgagga ctgcggaaat 3300

agaggaccct ctttgagaac aaccactgcc tctggaaaac tcataacaga atggtgctgc 3360

cgatcttgca cattaccacc gctaagatac agaggtgagg atgggtgctg gtacgggatg 3420

gaaatcagac cattgaagga gaaagaagag aatttggtca actccttggt cacagctgga 3480

catgggcagg tcgacaactt ttcactagga gtcttgggaa tggcattgtt cctggaggaa 3540

atgcttagga cccgagtagg aacgaaacat gcaatactac tagttgcagt ttcttttgtg 3600

acattgatca cagggaacat gtcctttaga gacctgggaa gagtgatggt tatggtaggc 3660

gccactatga cggatgacat aggtatgggc gtgacttatc ttgccctact agcagccttc 3720

aaagtcagac caacttttgc agctggacta ctcttgagaa agctgacctc caaggaattg 3780

atgatgacta ctataggaat tgtactcctc tcccagagca ccataccaga gaccattctt 3840

gagttgactg atgcgttagc cttaggcatg atggtcctca aaatggtgag aaatatggaa 3900

aagtatcaat tggcagtgac tatcatggct atcttgtgcg tcccaaacgc agtgatatta 3960

caaaacgcat ggaaagtgag ttgcacaata ttggcagtgg tgtccgtttc cccactgttc 4020

ttaacatcct cacagcaaaa aacagattgg ataccattag cattgacgat caaaggtctc 4080

aatccaacag ctatttttct aacaaccctc tcaagaacca gcaagaaaag gagctggcca 4140

ttaaatgagg ctatcatggc agtcgggatg gtgagcattt tagccagttc tctcctaaaa 4200

aatgatattc ccatgacagg accattagtg gctggagggc tcctcactgt gtgctacgtg 4260

ctcactggac gatcggccga tttggaactg gagagagcag ccgatgtcaa atgggaagac 4320

caggcagaga tatcaggaag cagtccaatc ctgtcaataa caatatcaga agatggtagc 4380

atgtcgataa aaaatgaaga ggaagaacaa acactgacca tactcattag aacaggattg 4440

ctggtgatct caggactttt tcctgtatca ataccaatca cggcagcagc atggtacctg 4500

tgggaagtga agaaacaacg ggccggagta ttgtgggatg ttccttcacc cccacccatg 4560

ggaaaggctg aactggaaga tggagcctat agaattaagc aaaaagggat tcttggatat 4620

tcccagatcg gagccggagt ttacaaagaa ggaacattcc atacaatgtg gcatgtcaca 4680

cgtggcgctg ttctaatgca taaaggaaag aggattgaac catcatgggc ggacgtcaag 4740

aaagacctaa tatcatatgg aggaggctgg aagttagaag gagaatggaa ggaaggagaa 4800

gaagtccagg tattggcact ggagcctgga aaaaatccaa gagccgtcca aacgaaacct 4860

ggtcttttca aaaccaacgc cggaacaata ggtgctgtat ctctggactt ttctcctgga 4920

acgtcaggat ctccaattat cgacaaaaaa ggaaaagttg tgggtcttta tggtaatggt 4980

gttgttacaa ggagtggagc atatgtgagt gctatagccc agactgaaaa aagcattgaa 5040

gacaacccag agatcgaaga tgacattttc cgaaagagaa gactgaccat catggacctc 5100

cacccaggag cgggaaagac gaagagatac cttccggcca tagtcagaga agctataaaa 5160

cggggtttga gaacattaat cttggccccc actagagttg tggcagctga aatggaggaa 5220

gcccttagag gacttccaat aagataccag accccagcca tcagagctgt gcacaccggg 5280

cgggagattg tggacctaat gtgtcatgcc acatttacca tgaggctgct atcaccagtt 5340

agagtgccaa actacaacct gattatcatg gacgaagccc atttcacaga cccagcaagt 5400

atagcagcta gaggatacat ctcaactcga gtggagatgg gtgaggcagc tgggattttt 5460

atgacagcca ctcccccggg aagcagagac ccatttcctc agagcaatgc accaatcata 5520

gatgaagaaa gagaaatccc tgaacgctcg tggaattccg gacatgaatg ggtcacggat 5580

tttaaaggga agactgtttg gttcgttcca agtataaaag caggaaatga tatagcagct 5640

tgcctgagga aaaatggaaa gaaagtgata caactcagta ggaagacctt tgattctgag 5700

tatgtcaaga ctagaaccaa tgattgggac ttcgtggtta caactgacat ttcagaaatg 5760

ggtgccaatt tcaaggctga gagggttata gaccccagac gctgcatgaa accagtcata 5820

ctaacagatg gtgaagagcg ggtgattctg gcaggaccta tgccagtgac ccactctagt 5880

gcagcacaaa gaagagggag aataggaaga aatccaaaaa atgagaatga ccagtacata 5940

tacatggggg aacctctgga aaatgatgaa gactgtgcac actggaaaga agctaaaatg 6000

ctcctagata acatcaacac gccagaagga atcattccta gcatgttcga accagagcgt 6060

gaaaaggtgg atgccattga tggcgaatac cgcttgagag gagaagcaag gaaaaccttt 6120

gtagacttaa tgagaagagg agacctacca gtctggttgg cctacagagt ggcagctgaa 6180

ggcatcaact acgcagacag aaggtggtgt tttgatggag tcaagaacaa ccaaatccta 6240

gaagaaaacg tggaagttga aatctggaca aaagaagggg aaaggaagaa attgaaaccc 6300

agatggttgg atgctaggat ctattctgac ccactggcgc taaaagaatt taaggaattt 6360

gcagccggaa gaaagtctct gaccctgaac ctaatcacag aaatgggtag gctcccaacc 6420

ttcatgactc agaaggcaag agacgcactg gacaacttag cagtgctgca cacggctgag 6480

ccaggtggaa gggcgtacaa ccatgctctc agtgaactgc cggagaccct ggagacattg 6540

cttttactga cacttctggc tacagtcacg ggagggatct ttttattctt gatgagcgca 6600

aggggcatag ggaagatgac cctgggaatg tgctgcataa tcacggctag catcctccta 6660

tggtacgcac aaatacagcc acactggata gcagcttcaa taatactgga gtttttttctc 6720

atagttttgc ttattccaga acctgaaaaa cagagaacac cccaagacaa ccaactgacc 6780

tacgttgtca tagccatcct cacagtggtg gccgcaacca tggcaaacga gatgggtttc 6840

ctagaaaaaa cgaagaaaga tctcggattg ggaagcattg caacccagca acccgagagc 6900

aacatcctgg acatagatct acgtcctgca tcagcatgga cgctgtatgc cgtggccaca 6960

acatttgtta caccaatgtt gagacatagc attgaaaatt cctcagtgaa tgtgtcccta 7020
acagctatag ccaaccaagc cacagtgtta atgggtctcg ggaaaggatg gccattgtca 7080
aagatggaca tcggagttcc ccttctcgcc attggatgct actcacaagt caaccccata 7140
actctcacag cagctttttt cttattggta gcacattatg ccatcatagg gccaggactc 7200
caagcaaaag caaccagaga agctcagaaa agagcagcgg cgggcatcat gaaaaaccca 7260
actgtcgatg gaataacagt gattgaccta gatccaatac cttatgatcc aaagtttgaa 7320
aagcagttgg gacaagtaat gctcctagtc ctctgcgtga ctcaagtatt gatgatgagg 7380
actacatggg ctctgtgtga ggctttaacc ttagctaccg ggcccatctc cacattgtgg 7440
gaaggaaatc cagggaggtt ttggaacact accattgcgg tgtcaatggc taacattttt 7500
agagggagtt acttggccgg agctggactt ctcttttcta ttatgaagaa cacaaccaac 7560
acaagaaggg gaactggcaa cataggagag acgcttggag agaaatggaa aagccgattg 7620
aacgcattgg gaaaaagtga attccagatc tacaagaaaa gtggaatcca ggaagtggat 7680
agaaccttag caaaagaagg cattaaaaga ggagaaacgg accatcacgc tgtgtcgcga 7740
ggctcagcaa aactgagatg gttcgttgag agaaacatgg tcacaccaga agggaaagta 7800
gtggacctcg gttgtggcag aggaggctgg tcatactatt gtggaggact aaagaatgta 7860
agagaagtca aaggcctaac aaaaggagga ccaggacacg aagaacccat ccccatgtca 7920
acatatgggt ggaatctagt gcgtcttcaa agtggagttg acgttttctt catcccgcca 7980
gaaaagtgtg acacattatt gtgtgacata ggggagtcat caccaaatcc cacagtggaa 8040
gcaggacgaa cactcagagt ccttaactta gtagaaaatt ggttgaacaa caacactcaa 8100
ttttgcataa aggttctcaa cccatatatg ccctcagtca tagaaaaaat ggaagcacta 8160
caaaggaaat atggaggagc cttagtgagg aatccactct cacgaaactc cacacatgag 8220
atgtactggg tatccaatgc ttccgggaac atagtgtcat cagtgaacat gatttcaagg 8280
atgttgatca acagatttac aatgagatac aagaaagcca cttacgagcc ggatgttgac 8340
ctcggaagcg gaacccgtaa catcgggatt gaaagtgaga taccaaacct agatataatt 8400
gggaaaagaa tagaaaaaat aaagcaagag catgaaacat catggcacta tgaccaagac 8460
cacccataca aaacgtgggc ataccatggt agctatgaaa caaaacagac tggatcagca 8520
tcatccatgg tcaacggagt ggtcaggctg ctgacaaaac cttgggacgt cgtccccatg 8580
gtgacacaga tggcaatgac agacacgact ccatttggac aacagcgcgt ttttaaagag 8640
aaagtggaca cgagaaccca agaaccgaaa gaaggcacga agaaactaat gaaaataaca 8700
gcagagtggc tttggaaaga attagggaag aaaaagacac ccaggatgtg caccagagaa 8760
gaattcacaa gaaaggtgag aagcaatgca gccttggggg ccgtattcac tgatgagaac 8820
aagtggaagt cggcacgtga ggctgttgaa gatagtaggt tttgggagct ggttgacaag 8880
gaaaggaatc tccatcttga aggaaagtgt gaaacatgtg tgtacaacat gatgggaaaa 8940
agagagaaga agctagggga attcggcaag gcaaaaggca gcagagccat atggtacatg 9000

tggcttggag cacgcttctt agagtttgaa gccctaggat tcttaaatga agatcactgg 9060

ttctccagag agaactccct gagtggagtg gaaggagaag ggctgcacaa gctaggttac 9120

attctaagag acgtgagcaa gaaagaggga ggagcaatgt atgccgatga caccgcagga 9180

tgggatacaa gaatcacact agaagaccta aaaaatgaag aaatggtaac aaaccacatg 9240

gaaggagaac acaagaaact agccgaggcc attttcaaac taacgtacca aaacaaggtg 9300

gtgcgtgtgc aaagaccaac accaagaggc acagtaatgg acatcatatc gagaagagac 9360

caaagaggta gtggacaagt tggcacctat ggactcaata ctttcaccaa tatggaagcc 9420

caactaatca gacagatgga gggagaagga gtctttaaaa gcattcagca cctaacaatc 9480

acagaagaaa tcgctgtgca aaactggtta gcaagagtgg ggcgcgaaag gttatcaaga 9540

atggccatca gtggagatga ttgtgttgtg aaacctttag atgacaggtt cgcaagcgct 9600

ttaacagctc taaatgacat gggaaagatt aggaaagaca tacaacaatg ggaaccttca 9660

agaggatgga atgattggac acaagtgccc ttctgttcac accatttcca tgagttaatc 9720

atgaaagacg gtcgcgtact cgttgttcca tgtagaaacc aagatgaact gattggcaga 9780

gcccgaatct cccaaggagc agggtggtct ttgcgggaga cggcctgttt ggggaagtct 9840

tacgcccaaa tgtggagctt gatgtacttc cacagacgcg acctcaggct ggcggcaaat 9900

gctatttgct cggcagtacc atcacattgg gttccaacaa gtcgaacaac ctggtccata 9960

catgctaaac atgaatggat gacaacggaa gacatgctga cagtctggaa cagggtgtgg 10020

attcaagaaa acccatggat ggaagacaaa actccagtgg aatcatggga ggaaatccca 10080

tacttgggga aaagagaaga ccaatggtgc ggctcattga ttgggttaac aagcagggcc 10140

acctgggcaa agaacatcca agcagcaata aatcaagtta gatcccttat aggcaatgaa 10200

gaatacacag attacatgcc atccatgaaa agattcagaa gagaagagga agaagcagga 10260

gttctgtggt agaaagcaaa actaacatga aacaaggcta gaagtcaggt cggattaagc 10320

catagtacgg aaaaaactat gctacctgtg agccccgtcc aaggacgtta aaagaagtca 10380

ggccatcata aatgccatag cttgagtaaa ctatgcagcc tgtagctcca cctgagaagg 10440

tgtaaaaaat ccgggaggcc acaaaccatg gaagctgtac gcatggcgta gtggactagc 10500

ggttagagga gacccctccc ttacaaatcg cagcaacaat gggggcccaa ggcgagatga 10560

agctgtagtc tcgctggaag gactagaggt tagaggagac ccccccgaaa caaaaaacag 10620

catattgacg ctgggaaaga ccagagatcc tgctgtctcc tcagcatcat tccaggcaca 10680

gaacgccaga aaatggaatg gtgctgttga atcaacaggt tct 10723

<210> 5
<211> 3391
<212> PRT
<213> Artificial Sequence

<220>
<223> Dengue virus serotype 2, BVS

<400> 5

```
Met Asn Asn Gln Arg Lys Lys Ala Lys Asn Thr Pro Phe Asn Met Leu
1               5                  10                  15

Lys Arg Glu Arg Asn Arg Val Ser Thr Val Gln Gln Leu Thr Lys Arg
            20                  25                  30

Phe Ser Leu Gly Met Leu Gln Gly Arg Gly Pro Leu Lys Leu Phe Met
        35                  40                  45

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

Ile Leu Lys Arg Trp Gly Thr Ile Lys Lys Ser Lys Ala Ile Asn Val
65                  70                  75                  80

Leu Arg Gly Phe Arg Lys Glu Ile Gly Arg Met Leu Asn Ile Leu Asn
                85                  90                  95

Arg Arg Arg Arg Ser Ala Gly Met Ile Ile Met Leu Ile Pro Thr Val
            100                 105                 110

Met Ala Phe His Leu Thr Thr Arg Asn Gly Glu Pro His Met Ile Val
        115                 120                 125

Ser Arg Gln Glu Lys Gly Lys Ser Leu Leu Phe Lys Thr Glu Val Gly
    130                 135                 140

Val Asn Met Cys Thr Leu Met Ala Met Asp Leu Gly Glu Leu Cys Glu
145                 150                 155                 160

Asp Thr Ile Thr Tyr Glu Cys Pro Leu Leu Arg Gln Asn Glu Pro Glu
                165                 170                 175

Asp Ile Asp Cys Trp Cys Asn Ser Thr Ser Thr Trp Val Thr Tyr Gly
            180                 185                 190

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

Leu Val Pro His Val Gly Met Gly Leu Glu Thr Arg Thr Glu Thr Trp
    210                 215                 220

Met Ser Ser Glu Gly Ala Trp Lys His Val Gln Arg Ile Glu Thr Trp
225                 230                 235                 240

Ile Leu Arg His Pro Gly Phe Thr Met Met Ala Ala Ile Leu Ala Tyr
                245                 250                 255

Thr Ile Gly Thr Thr His Phe Gln Arg Ala Leu Ile Phe Ile Leu Leu
            260                 265                 270
```

```
Thr Ala Val Thr Pro Ser Met Thr Met Arg Cys Ile Gly Met Ser Asn
        275                 280                 285

Arg Asp Phe Val Glu Gly Val Ser Gly Gly Ser Trp Val Asp Ile Val
    290                 295                 300

Leu Glu His Gly Ser Cys Val Thr Thr Met Ala Lys Asn Lys Pro Thr
305                 310                 315                 320

Leu Asp Phe Glu Leu Ile Lys Thr Glu Ala Lys Gln Pro Ala Thr Leu
                325                 330                 335

Arg Lys Tyr Cys Ile Glu Ala Lys Leu Thr Asn Thr Thr Thr Glu Ser
            340                 345                 350

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

Arg Phe Val Cys Lys His Ser Met Val Asp Arg Gly Trp Gly Asn Gly
    370                 375                 380

Cys Gly Leu Phe Gly Lys Gly Gly Ile Val Thr Cys Ala Met Phe Arg
385                 390                 395                 400

Cys Lys Lys Asn Met Glu Gly Lys Val Val Gln Pro Glu Asn Leu Glu
                405                 410                 415

Tyr Thr Ile Val Ile Thr Pro His Ser Gly Glu Glu His Ala Val Gly
            420                 425                 430

Asn Asp Thr Gly Lys His Gly Lys Glu Ile Lys Ile Thr Pro Gln Ser
        435                 440                 445

Ser Ile Thr Glu Ala Glu Leu Thr Gly Tyr Gly Thr Val Thr Met Glu
    450                 455                 460

Cys Ser Pro Arg Thr Gly Leu Asp Phe Asn Glu Met Val Leu Leu Gln
465                 470                 475                 480

Met Glu Asn Lys Ala Trp Leu Val His Arg Gln Trp Phe Leu Asp Leu
                485                 490                 495

Pro Leu Pro Trp Leu Pro Gly Ala Asp Thr Gln Gly Ser Asn Trp Ile
            500                 505                 510

Gln Lys Glu Thr Leu Val Thr Phe Lys Asn Pro His Ala Lys Lys Gln
        515                 520                 525

Asp Val Val Val Leu Gly Ser Gln Glu Gly Ala Met His Thr Ala Leu
    530                 535                 540
```

```
Thr Gly Ala Thr Glu Ile Gln Met Ser Ser Gly Asn Leu Leu Phe Thr
545                 550                 555                 560

Gly His Leu Lys Cys Arg Leu Arg Met Asp Lys Leu Gln Leu Lys Gly
                565                 570                 575

Met Ser Tyr Ser Met Cys Thr Gly Lys Phe Lys Val Val Lys Glu Ile
            580                 585                 590

Ala Glu Thr Gln His Gly Thr Ile Val Ile Arg Val Gln Tyr Glu Gly
        595                 600                 605

Asp Gly Ser Pro Cys Lys Ile Pro Phe Glu Ile Met Asp Leu Glu Lys
    610                 615                 620

Arg His Val Leu Gly Arg Leu Ile Thr Val Asn Pro Ile Val Thr Glu
625                 630                 635                 640

Lys Asp Ser Pro Val Asn Ile Glu Ala Glu Pro Pro Phe Gly Asp Ser
                645                 650                 655

Tyr Ile Ile Ile Gly Val Glu Pro Gly Gln Leu Lys Leu Asn Trp Phe
            660                 665                 670

Lys Lys Gly Ser Ser Ile Gly Gln Met Phe Glu Thr Thr Met Arg Gly
        675                 680                 685

Ala Lys Arg Met Ala Ile Leu Gly Asp Thr Ala Trp Asp Phe Gly Ser
    690                 695                 700

Leu Gly Gly Val Phe Thr Ser Ile Gly Lys Ala Leu His Gln Val Phe
705                 710                 715                 720

Gly Ala Ile Tyr Gly Ala Ala Phe Ser Gly Val Ser Trp Thr Met Lys
                725                 730                 735

Ile Leu Ile Gly Val Ile Ile Thr Trp Ile Gly Met Asn Ser Arg Ser
            740                 745                 750

Thr Ser Leu Ser Val Thr Leu Val Leu Val Gly Ile Val Thr Leu Tyr
        755                 760                 765

Leu Gly Val Met Val Gln Ala Asp Ser Gly Cys Val Val Ser Trp Lys
    770                 775                 780

Asn Lys Glu Leu Lys Cys Gly Ser Gly Ile Phe Ile Thr Asp Asn Val
785                 790                 795                 800

His Thr Trp Thr Glu Gln Tyr Lys Phe Gln Pro Glu Ser Pro Ser Lys
                805                 810                 815
```

```
Leu Ala Ser Ala Ile Gln Lys Ala His Glu Glu Asp Ile Cys Gly Ile
            820                 825                 830

Arg Ser Val Thr Arg Leu Glu Asn Leu Met Trp Lys Gln Ile Thr Pro
        835                 840                 845

Glu Leu Asn His Ile Leu Ser Glu Asn Glu Val Lys Leu Thr Ile Met
    850                 855                 860

Thr Gly Asp Ile Lys Gly Ile Met Gln Ala Gly Lys Arg Ser Leu Arg
865                 870                 875                 880

Pro Gln Pro Thr Glu Leu Lys Tyr Ser Trp Lys Thr Trp Gly Lys Ala
                885                 890                 895

Lys Met Leu Ser Thr Glu Ser His Asn Gln Thr Phe Leu Ile Asp Gly
            900                 905                 910

Pro Glu Thr Ala Glu Cys Pro Asn Thr Asn Arg Ala Trp Asn Ser Leu
        915                 920                 925

Glu Val Glu Asp Tyr Gly Phe Gly Val Phe Thr Thr Asn Ile Trp Leu
    930                 935                 940

Lys Leu Lys Glu Lys Gln Asp Val Phe Cys Asp Ser Lys Leu Met Ser
945                 950                 955                 960

Ala Ala Ile Lys Asp Asn Arg Ala Val His Ala Asp Met Gly Tyr Trp
                965                 970                 975

Ile Glu Ser Ala Leu Asn Asp Thr Trp Lys Ile Glu Lys Ala Ser Phe
            980                 985                 990

Ile Glu Val Lys Asn Cys His Trp  Pro Lys Ser His Thr  Leu Trp Ser
        995                 1000                1005

Asn Gly  Val Leu Glu Ser Glu  Met Ile Ile Pro Lys  Asn Leu Ala
    1010                1015                1020

Gly Pro  Val Ser Gln His Asn  Tyr Arg Pro Gly Tyr  His Thr Gln
    1025                1030                1035

Ile Thr  Gly Pro Trp His Leu  Gly Lys Leu Glu Met  Asp Phe Asp
    1040                1045                1050

Phe Cys  Asp Gly Thr Thr Val  Val Val Thr Glu Asp  Cys Gly Asn
    1055                1060                1065

Arg Gly  Pro Ser Leu Arg Thr  Thr Thr Ala Ser Gly  Lys Leu Ile
    1070                1075                1080
```

```
Thr Glu  Trp Cys Cys Arg Ser  Cys Thr Leu Pro Pro  Leu Arg Tyr
    1085                 1090                 1095

Arg Gly  Glu Asp Gly Cys Trp  Tyr Gly Met Glu Ile  Arg Pro Leu
    1100                 1105                 1110

Lys Glu  Lys Glu Glu Asn Leu  Val Asn Ser Leu Val  Thr Ala Gly
    1115                 1120                 1125

His Gly  Gln Val Asp Asn Phe  Ser Leu Gly Val Leu  Gly Met Ala
    1130                 1135                 1140

Leu Phe  Leu Glu Glu Met Leu  Arg Thr Arg Val Gly  Thr Lys His
    1145                 1150                 1155

Ala Ile  Leu Leu Val Ala Val  Ser Phe Val Thr Leu  Ile Thr Gly
    1160                 1165                 1170

Asn Met  Ser Phe Arg Asp Leu  Gly Arg Val Met Val  Met Val Gly
    1175                 1180                 1185

Ala Thr  Met Thr Asp Asp Ile  Gly Met Gly Val Thr  Tyr Leu Ala
    1190                 1195                 1200

Leu Leu  Ala Ala Phe Lys Val  Arg Pro Thr Phe Ala  Ala Gly Leu
    1205                 1210                 1215

Leu Leu  Arg Lys Leu Thr Ser  Lys Glu Leu Met Met  Thr Thr Ile
    1220                 1225                 1230

Gly Ile  Val Leu Leu Ser Gln  Ser Thr Ile Pro Glu  Thr Ile Leu
    1235                 1240                 1245

Glu Leu  Thr Asp Ala Leu Ala  Leu Gly Met Met Val  Leu Lys Met
    1250                 1255                 1260

Val Arg  Asn Met Glu Lys Tyr  Gln Leu Ala Val Thr  Ile Met Ala
    1265                 1270                 1275

Ile Leu  Cys Val Pro Asn Ala  Val Ile Leu Gln Asn  Ala Trp Lys
    1280                 1285                 1290

Val Ser  Cys Thr Ile Leu Ala  Val Val Ser Val Ser  Pro Leu Phe
    1295                 1300                 1305

Leu Thr  Ser Ser Gln Gln Lys  Thr Asp Trp Ile Pro  Leu Ala Leu
    1310                 1315                 1320

Thr Ile  Lys Gly Leu Asn Pro  Thr Ala Ile Phe Leu  Thr Thr Leu
    1325                 1330                 1335
```

```
Ser Arg  Thr Ser Lys Lys Arg  Ser Trp Pro Leu Asn  Glu Ala Ile
    1340                 1345                 1350

Met Ala  Val Gly Met Val Ser  Ile Leu Ala Ser Ser  Leu Leu Lys
    1355                 1360                 1365

Asn Asp  Ile Pro Met Thr Gly  Pro Leu Val Ala Gly  Gly Leu Leu
    1370                 1375                 1380

Thr Val  Cys Tyr Val Leu Thr  Gly Arg Ser Ala Asp  Leu Glu Leu
    1385                 1390                 1395

Glu Arg  Ala Ala Asp Val Lys  Trp Glu Asp Gln Ala  Glu Ile Ser
    1400                 1405                 1410

Gly Ser  Ser Pro Ile Leu Ser  Ile Thr Ile Ser Glu  Asp Gly Ser
    1415                 1420                 1425

Met Ser  Ile Lys Asn Glu Glu  Glu Glu Gln Thr Leu  Thr Ile Leu
    1430                 1435                 1440

Ile Arg  Thr Gly Leu Leu Val  Ile Ser Gly Leu Phe  Pro Val Ser
    1445                 1450                 1455

Ile Pro  Ile Thr Ala Ala Ala  Trp Tyr Leu Trp Glu  Val Lys Lys
    1460                 1465                 1470

Gln Arg  Ala Gly Val Leu Trp  Asp Val Pro Ser Pro  Pro Pro Met
    1475                 1480                 1485

Gly Lys  Ala Glu Leu Glu Asp  Gly Ala Tyr Arg Ile  Lys Gln Lys
    1490                 1495                 1500

Gly Ile  Leu Gly Tyr Ser Gln  Ile Gly Ala Gly Val  Tyr Lys Glu
    1505                 1510                 1515

Gly Thr  Phe His Thr Met Trp  His Val Thr Arg Gly  Ala Val Leu
    1520                 1525                 1530

Met His  Lys Gly Lys Arg Ile  Glu Pro Ser Trp Ala  Asp Val Lys
    1535                 1540                 1545

Lys Asp  Leu Ile Ser Tyr Gly  Gly Gly Trp Lys Leu  Glu Gly Glu
    1550                 1555                 1560

Trp Lys  Glu Gly Glu Glu Val  Gln Val Leu Ala Leu  Glu Pro Gly
    1565                 1570                 1575

Lys Asn  Pro Arg Ala Val Gln  Thr Lys Pro Gly Leu  Phe Lys Thr
    1580                 1585                 1590
```

```
Asn Ala  Gly Thr Ile Gly Ala  Val Ser Leu Asp Phe  Ser Pro Gly
    1595                 1600                 1605

Thr Ser  Gly Ser Pro Ile Ile  Asp Lys Lys Gly Lys  Val Val Gly
    1610                 1615                 1620

Leu Tyr  Gly Asn Gly Val Val  Thr Arg Ser Gly Ala  Tyr Val Ser
    1625                 1630                 1635

Ala Ile  Ala Gln Thr Glu Lys  Ser Ile Glu Asp Asn  Pro Glu Ile
    1640                 1645                 1650

Glu Asp  Asp Ile Phe Arg Lys  Arg Arg Leu Thr Ile  Met Asp Leu
    1655                 1660                 1665

His Pro  Gly Ala Gly Lys Thr  Lys Arg Tyr Leu Pro  Ala Ile Val
    1670                 1675                 1680

Arg Glu  Ala Ile Lys Arg Gly  Leu Arg Thr Leu Ile  Leu Ala Pro
    1685                 1690                 1695

Thr Arg  Val Val Ala Ala Glu  Met Glu Glu Ala Leu  Arg Gly Leu
    1700                 1705                 1710

Pro Ile  Arg Tyr Gln Thr Pro  Ala Ile Arg Ala Val  His Thr Gly
    1715                 1720                 1725

Arg Glu  Ile Val Asp Leu Met  Cys His Ala Thr Phe  Thr Met Arg
    1730                 1735                 1740

Leu Leu  Ser Pro Val Arg Val  Pro Asn Tyr Asn Leu  Ile Ile Met
    1745                 1750                 1755

Asp Glu  Ala His Phe Thr Asp  Pro Ala Ser Ile Ala  Ala Arg Gly
    1760                 1765                 1770

Tyr Ile  Ser Thr Arg Val Glu  Met Gly Glu Ala Ala  Gly Ile Phe
    1775                 1780                 1785

Met Thr  Ala Thr Pro Pro Gly  Ser Arg Asp Pro Phe  Pro Gln Ser
    1790                 1795                 1800

Asn Ala  Pro Ile Ile Asp Glu  Glu Arg Glu Ile Pro  Glu Arg Ser
    1805                 1810                 1815

Trp Asn  Ser Gly His Glu Trp  Val Thr Asp Phe Lys  Gly Lys Thr
    1820                 1825                 1830

Val Trp  Phe Val Pro Ser Ile  Lys Ala Gly Asn Asp  Ile Ala Ala
    1835                 1840                 1845
```

```
Cys Leu  Arg Lys Asn Gly Lys  Lys Val Ile Gln Leu  Ser Arg Lys
    1850                 1855                 1860

Thr Phe  Asp Ser Glu Tyr Val  Lys Thr Arg Thr Asn  Asp Trp Asp
    1865                 1870                 1875

Phe Val  Val Thr Thr Asp Ile  Ser Glu Met Gly Ala  Asn Phe Lys
    1880                 1885                 1890

Ala Glu  Arg Val Ile Asp Pro  Arg Arg Cys Met Lys  Pro Val Ile
    1895                 1900                 1905

Leu Thr  Asp Gly Glu Glu Arg  Val Ile Leu Ala Gly  Pro Met Pro
    1910                 1915                 1920

Val Thr  His Ser Ser Ala Ala  Gln Arg Arg Gly Arg  Ile Gly Arg
    1925                 1930                 1935

Asn Pro  Lys Asn Glu Asn Asp  Gln Tyr Ile Tyr Met  Gly Glu Pro
    1940                 1945                 1950

Leu Glu  Asn Asp Glu Asp Cys  Ala His Trp Lys Glu  Ala Lys Met
    1955                 1960                 1965

Leu Leu  Asp Asn Ile Asn Thr  Pro Glu Gly Ile Ile  Pro Ser Met
    1970                 1975                 1980

Phe Glu  Pro Glu Arg Glu Lys  Val Asp Ala Ile Asp  Gly Glu Tyr
    1985                 1990                 1995

Arg Leu  Arg Gly Glu Ala Arg  Lys Thr Phe Val Asp  Leu Met Arg
    2000                 2005                 2010

Arg Gly  Asp Leu Pro Val Trp  Leu Ala Tyr Arg Val  Ala Ala Glu
    2015                 2020                 2025

Gly Ile  Asn Tyr Ala Asp Arg  Arg Trp Cys Phe Asp  Gly Val Lys
    2030                 2035                 2040

Asn Asn  Gln Ile Leu Glu Glu  Asn Val Glu Val Glu  Ile Trp Thr
    2045                 2050                 2055

Lys Glu  Gly Glu Arg Lys Lys  Leu Lys Pro Arg Trp  Leu Asp Ala
    2060                 2065                 2070

Arg Ile  Tyr Ser Asp Pro Leu  Ala Leu Lys Glu Phe  Lys Glu Phe
    2075                 2080                 2085

Ala Ala  Gly Arg Lys Ser Leu  Thr Leu Asn Leu Ile  Thr Glu Met
    2090                 2095                 2100
```

```
Gly Arg  Leu Pro Thr Phe Met  Thr Gln Lys Ala Arg  Asp Ala Leu
    2105                 2110                 2115

Asp Asn  Leu Ala Val Leu His  Thr Ala Glu Pro Gly  Gly Arg Ala
    2120                 2125                 2130

Tyr Asn  His Ala Leu Ser Glu  Leu Pro Glu Thr Leu  Glu Thr Leu
    2135                 2140                 2145

Leu Leu  Leu Thr Leu Leu Ala  Thr Val Thr Gly Gly  Ile Phe Leu
    2150                 2155                 2160

Phe Leu  Met Ser Ala Arg Gly  Ile Gly Lys Met Thr  Leu Gly Met
    2165                 2170                 2175

Cys Cys  Ile Ile Thr Ala Ser  Ile Leu Leu Trp Tyr  Ala Gln Ile
    2180                 2185                 2190

Gln Pro  His Trp Ile Ala Ala  Ser Ile Ile Leu Glu  Phe Phe Leu
    2195                 2200                 2205

Ile Val  Leu Leu Ile Pro Glu  Pro Glu Lys Gln Arg  Thr Pro Gln
    2210                 2215                 2220

Asp Asn  Gln Leu Thr Tyr Val  Val Ile Ala Ile Leu  Thr Val Val
    2225                 2230                 2235

Ala Ala  Thr Met Ala Asn Glu  Met Gly Phe Leu Glu  Lys Thr Lys
    2240                 2245                 2250

Lys Asp  Leu Gly Leu Gly Ser  Ile Ala Thr Gln Gln  Pro Glu Ser
    2255                 2260                 2265

Asn Ile  Leu Asp Ile Asp Leu  Arg Pro Ala Ser Ala  Trp Thr Leu
    2270                 2275                 2280

Tyr Ala  Val Ala Thr Thr Phe  Val Thr Pro Met Leu  Arg His Ser
    2285                 2290                 2295

Ile Glu  Asn Ser Ser Val Asn  Val Ser Leu Thr Ala  Ile Ala Asn
    2300                 2305                 2310

Gln Ala  Thr Val Leu Met Gly  Leu Gly Lys Gly Trp  Pro Leu Ser
    2315                 2320                 2325

Lys Met  Asp Ile Gly Val Pro  Leu Leu Ala Ile Gly  Cys Tyr Ser
    2330                 2335                 2340

Gln Val  Asn Pro Ile Thr Leu  Thr Ala Ala Phe Phe  Leu Leu Val
    2345                 2350                 2355
```

```
Ala His  Tyr Ala Ile Ile Gly  Pro Gly Leu Gln Ala  Lys Ala Thr
    2360                 2365                 2370

Arg Glu  Ala Gln Lys Arg Ala  Ala Ala Gly Ile Met  Lys Asn Pro
    2375                 2380                 2385

Thr Val  Asp Gly Ile Thr Val  Ile Asp Leu Asp Pro  Ile Pro Tyr
    2390                 2395                 2400

Asp Pro  Lys Phe Glu Lys Gln  Leu Gly Gln Val Met  Leu Leu Val
    2405                 2410                 2415

Leu Cys  Val Thr Gln Val Leu  Met Met Arg Thr Thr  Trp Ala Leu
    2420                 2425                 2430

Cys Glu  Ala Leu Thr Leu Ala  Thr Gly Pro Ile Ser  Thr Leu Trp
    2435                 2440                 2445

Glu Gly  Asn Pro Gly Arg Phe  Trp Asn Thr Thr Ile  Ala Val Ser
    2450                 2455                 2460

Met Ala  Asn Ile Phe Arg Gly  Ser Tyr Leu Ala Gly  Ala Gly Leu
    2465                 2470                 2475

Leu Phe  Ser Ile Met Lys Asn  Thr Thr Asn Thr Arg  Arg Gly Thr
    2480                 2485                 2490

Gly Asn  Ile Gly Glu Thr Leu  Gly Glu Lys Trp Lys  Ser Arg Leu
    2495                 2500                 2505

Asn Ala  Leu Gly Lys Ser Glu  Phe Gln Ile Tyr Lys  Lys Ser Gly
    2510                 2515                 2520

Ile Gln  Glu Val Asp Arg Thr  Leu Ala Lys Glu Gly  Ile Lys Arg
    2525                 2530                 2535

Gly Glu  Thr Asp His His Ala  Val Ser Arg Gly Ser  Ala Lys Leu
    2540                 2545                 2550

Arg Trp  Phe Val Glu Arg Asn  Met Val Thr Pro Glu  Gly Lys Val
    2555                 2560                 2565

Val Asp  Leu Gly Cys Gly Arg  Gly Gly Trp Ser Tyr  Tyr Cys Gly
    2570                 2575                 2580

Gly Leu  Lys Asn Val Arg Glu  Val Lys Gly Leu Thr  Lys Gly Gly
    2585                 2590                 2595

Pro Gly  His Glu Glu Pro Ile  Pro Met Ser Thr Tyr  Gly Trp Asn
    2600                 2605                 2610
```

```
Leu Val  Arg Leu Gln Ser Gly  Val Asp Val Phe Phe  Ile Pro Pro
    2615                 2620                 2625

Glu Lys  Cys Asp Thr Leu Leu  Cys Asp Ile Gly Glu  Ser Ser Pro
    2630                 2635                 2640

Asn Pro  Thr Val Glu Ala Gly  Arg Thr Leu Arg Val  Leu Asn Leu
    2645                 2650                 2655

Val Glu  Asn Trp Leu Asn Asn  Asn Thr Gln Phe Cys  Ile Lys Val
    2660                 2665                 2670

Leu Asn  Pro Tyr Met Pro Ser  Val Ile Glu Lys Met  Glu Ala Leu
    2675                 2680                 2685

Gln Arg  Lys Tyr Gly Gly Ala  Leu Val Arg Asn Pro  Leu Ser Arg
    2690                 2695                 2700

Asn Ser  Thr His Glu Met Tyr  Trp Val Ser Asn Ala  Ser Gly Asn
    2705                 2710                 2715

Ile Val  Ser Ser Val Asn Met  Ile Ser Arg Met Leu  Ile Asn Arg
    2720                 2725                 2730

Phe Thr  Met Arg Tyr Lys Lys  Ala Thr Tyr Glu Pro  Asp Val Asp
    2735                 2740                 2745

Leu Gly  Ser Gly Thr Arg Asn  Ile Gly Ile Glu Ser  Glu Ile Pro
    2750                 2755                 2760

Asn Leu  Asp Ile Ile Gly Lys  Arg Ile Glu Lys Ile  Lys Gln Glu
    2765                 2770                 2775

His Glu  Thr Ser Trp His Tyr  Asp Gln Asp His Pro  Tyr Lys Thr
    2780                 2785                 2790

Trp Ala  Tyr His Gly Ser Tyr  Glu Thr Lys Gln Thr  Gly Ser Ala
    2795                 2800                 2805

Ser Ser  Met Val Asn Gly Val  Val Arg Leu Leu Thr  Lys Pro Trp
    2810                 2815                 2820

Asp Val  Val Pro Met Val Thr  Gln Met Ala Met Thr  Asp Thr Thr
    2825                 2830                 2835

Pro Phe  Gly Gln Gln Arg Val  Phe Lys Glu Lys Val  Asp Thr Arg
    2840                 2845                 2850

Thr Gln  Glu Pro Lys Glu Gly  Thr Lys Lys Leu Met  Lys Ile Thr
    2855                 2860                 2865
```

```
Ala Glu  Trp Leu Trp Lys Glu  Leu Gly Lys Lys Lys  Thr Pro Arg
    2870                 2875                 2880

Met Cys  Thr Arg Glu Glu Phe  Thr Arg Lys Val Arg  Ser Asn Ala
    2885                 2890                 2895

Ala Leu  Gly Ala Val Phe Thr  Asp Glu Asn Lys Trp  Lys Ser Ala
    2900                 2905                 2910

Arg Glu  Ala Val Glu Asp Ser  Arg Phe Trp Glu Leu  Val Asp Lys
    2915                 2920                 2925

Glu Arg  Asn Leu His Leu Glu  Gly Lys Cys Glu Thr  Cys Val Tyr
    2930                 2935                 2940

Asn Met  Met Gly Lys Arg Glu  Lys Lys Leu Gly Glu  Phe Gly Lys
    2945                 2950                 2955

Ala Lys  Gly Ser Arg Ala Ile  Trp Tyr Met Trp Leu  Gly Ala Arg
    2960                 2965                 2970

Phe Leu  Glu Phe Glu Ala Leu  Gly Phe Leu Asn Glu  Asp His Trp
    2975                 2980                 2985

Phe Ser  Arg Glu Asn Ser Leu  Ser Gly Val Glu Gly  Glu Gly Leu
    2990                 2995                 3000

His Lys  Leu Gly Tyr Ile Leu  Arg Asp Val Ser Lys  Lys Glu Gly
    3005                 3010                 3015

Gly Ala  Met Tyr Ala Asp Asp  Thr Ala Gly Trp Asp  Thr Arg Ile
    3020                 3025                 3030

Thr Leu  Glu Asp Leu Lys Asn  Glu Glu Met Val Thr  Asn His Met
    3035                 3040                 3045

Glu Gly  Glu His Lys Lys Leu  Ala Glu Ala Ile Phe  Lys Leu Thr
    3050                 3055                 3060

Tyr Gln  Asn Lys Val Val Arg  Val Gln Arg Pro Thr  Pro Arg Gly
    3065                 3070                 3075

Thr Val  Met Asp Ile Ile Ser  Arg Arg Asp Gln Arg  Gly Ser Gly
    3080                 3085                 3090

Gln Val  Gly Thr Tyr Gly Leu  Asn Thr Phe Thr Asn  Met Glu Ala
    3095                 3100                 3105

Gln Leu  Ile Arg Gln Met Glu  Gly Glu Gly Val Phe  Lys Ser Ile
    3110                 3115                 3120
```

```
Gln His  Leu Thr Ile Thr Glu  Glu Ile Ala Val Gln  Asn Trp Leu
    3125                3130                3135

Ala Arg  Val Gly Arg Glu Arg  Leu Ser Arg Met Ala  Ile Ser Gly
    3140                3145                3150

Asp Asp  Cys Val Val Lys Pro  Leu Asp Asp Arg Phe  Ala Ser Ala
    3155                3160                3165

Leu Thr  Ala Leu Asn Asp Met  Gly Lys Ile Arg Lys  Asp Ile Gln
    3170                3175                3180

Gln Trp  Glu Pro Ser Arg Gly  Trp Asn Asp Trp Thr  Gln Val Pro
    3185                3190                3195

Phe Cys  Ser His His Phe His  Glu Leu Ile Met Lys  Asp Gly Arg
    3200                3205                3210

Val Leu  Val Val Pro Cys Arg  Asn Gln Asp Glu Leu  Ile Gly Arg
    3215                3220                3225

Ala Arg  Ile Ser Gln Gly Ala  Gly Trp Ser Leu Arg  Glu Thr Ala
    3230                3235                3240

Cys Leu  Gly Lys Ser Tyr Ala  Gln Met Trp Ser Leu  Met Tyr Phe
    3245                3250                3255

His Arg  Arg Asp Leu Arg Leu  Ala Ala Asn Ala Ile  Cys Ser Ala
    3260                3265                3270

Val Pro  Ser His Trp Val Pro  Thr Ser Arg Thr Thr  Trp Ser Ile
    3275                3280                3285

His Ala  Lys His Glu Trp Met  Thr Thr Glu Asp Met  Leu Thr Val
    3290                3295                3300

Trp Asn  Arg Val Trp Ile Gln  Glu Asn Pro Trp Met  Glu Asp Lys
    3305                3310                3315

Thr Pro  Val Glu Ser Trp Glu  Glu Ile Pro Tyr Leu  Gly Lys Arg
    3320                3325                3330

Glu Asp  Gln Trp Cys Gly Ser  Leu Ile Gly Leu Thr  Ser Arg Ala
    3335                3340                3345

Thr Trp  Ala Lys Asn Ile Gln  Ala Ala Ile Asn Gln  Val Arg Ser
    3350                3355                3360

Leu Ile  Gly Asn Glu Glu Tyr  Thr Asp Tyr Met Pro  Ser Met Lys
    3365                3370                3375
```

Arg Phe Arg Arg Glu Glu Glu Glu Ala Gly Val Leu Trp
3380 3385 3390

<210> 6
<211> 3391
<212> PRT
<213> Artificial Sequence

<220>
<223> Dengue virus serotype 2, MVS

<400> 6

Met Asn Asn Gln Arg Lys Lys Ala Lys Asn Thr Pro Phe Asn Met Leu
1 5 10 15

Lys Arg Glu Arg Asn Arg Val Ser Thr Val Gln Gln Leu Thr Lys Arg
20 25 30

Phe Ser Leu Gly Met Leu Gln Gly Arg Gly Pro Leu Lys Leu Phe Met
35 40 45

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

Ile Leu Lys Arg Trp Gly Thr Ile Lys Lys Ser Lys Ala Ile Asn Val
65 70 75 80

Leu Arg Gly Phe Arg Lys Glu Ile Gly Arg Met Leu Asn Ile Leu Asn
85 90 95

Arg Arg Arg Arg Ser Ala Gly Met Ile Ile Met Leu Ile Pro Thr Val
100 105 110

Met Ala Phe His Leu Thr Thr Arg Asn Gly Glu Pro His Met Ile Val
115 120 125

Ser Arg Gln Glu Lys Gly Lys Ser Leu Leu Phe Lys Thr Glu Val Gly
130 135 140

Val Asn Met Cys Thr Leu Met Ala Met Asp Leu Gly Glu Leu Cys Glu
145 150 155 160

Asp Thr Ile Thr Tyr Glu Cys Pro Leu Leu Arg Gln Asn Glu Pro Glu
165 170 175

Asp Ile Asp Cys Trp Cys Asn Ser Thr Ser Thr Trp Val Thr Tyr Gly
180 185 190

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

Leu Val Pro His Val Gly Met Gly Leu Glu Thr Arg Thr Glu Thr Trp

210 215 220

Met Ser Ser Glu Gly Ala Trp Lys His Val Gln Arg Ile Glu Thr Trp
225 230 235 240

Ile Leu Arg His Pro Gly Phe Thr Met Met Ala Ala Ile Leu Ala Tyr
245 250 255

Thr Ile Gly Thr Thr His Phe Gln Arg Ala Leu Ile Phe Ile Leu Leu
260 265 270

Thr Ala Val Thr Pro Ser Met Thr Met Arg Cys Ile Gly Met Ser Asn
275 280 285

Arg Asp Phe Val Glu Gly Val Ser Gly Gly Ser Trp Val Asp Ile Val
290 295 300

Leu Glu His Gly Ser Cys Val Thr Thr Met Ala Lys Asn Lys Pro Thr
305 310 315 320

Leu Asp Phe Glu Leu Ile Lys Thr Glu Ala Lys Gln Pro Ala Thr Leu
325 330 335

Arg Lys Tyr Cys Ile Glu Ala Lys Leu Thr Asn Thr Thr Thr Glu Ser
340 345 350

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

Arg Phe Val Cys Lys His Ser Met Val Asp Arg Gly Trp Gly Asn Gly
370 375 380

Cys Gly Leu Phe Gly Lys Gly Gly Ile Val Thr Cys Ala Met Phe Arg
385 390 395 400

Cys Lys Lys Asn Met Glu Gly Lys Val Val Gln Pro Glu Asn Leu Glu
405 410 415

Tyr Thr Ile Val Ile Thr Pro His Ser Gly Glu Glu His Ala Val Gly
420 425 430

Asn Asp Thr Gly Lys His Gly Lys Glu Ile Lys Ile Thr Pro Gln Ser
435 440 445

Ser Ile Thr Glu Ala Glu Leu Thr Gly Tyr Gly Thr Val Thr Met Glu
450 455 460

Cys Ser Pro Arg Thr Gly Leu Asp Phe Asn Glu Met Val Leu Leu Gln
465 470 475 480

Met Glu Asn Lys Ala Trp Leu Val His Arg Gln Trp Phe Leu Asp Leu

485 490 495

Pro Leu Pro Trp Leu Pro Gly Ala Asp Thr Gln Gly Ser Asn Trp Ile
500 505 510

Gln Lys Glu Thr Leu Val Thr Phe Lys Asn Pro His Ala Lys Lys Gln
515 520 525

Asp Val Val Val Leu Gly Ser Gln Glu Gly Ala Met His Thr Ala Leu
530 535 540

Thr Gly Ala Thr Glu Ile Gln Met Ser Ser Gly Asn Leu Leu Phe Thr
545 550 555 560

Gly His Leu Lys Cys Arg Leu Arg Met Asp Lys Leu Gln Leu Lys Gly
565 570 575

Met Ser Tyr Ser Met Cys Thr Gly Lys Phe Lys Val Val Lys Glu Ile
580 585 590

Ala Glu Thr Gln His Gly Thr Ile Val Ile Arg Val Gln Tyr Glu Gly
595 600 605

Asp Gly Ser Pro Cys Lys Ile Pro Phe Glu Ile Met Asp Leu Glu Lys
610 615 620

Arg His Val Leu Gly Arg Leu Ile Thr Val Asn Pro Ile Val Thr Glu
625 630 635 640

Lys Asp Ser Pro Val Asn Ile Glu Ala Glu Pro Pro Phe Gly Asp Ser
645 650 655

Tyr Ile Ile Ile Gly Val Glu Pro Gly Gln Leu Lys Leu Asn Trp Phe
660 665 670

Lys Lys Gly Ser Ser Ile Gly Gln Met Phe Glu Thr Thr Met Arg Gly
675 680 685

Ala Lys Arg Met Ala Ile Leu Gly Asp Thr Ala Trp Asp Phe Gly Ser
690 695 700

Leu Gly Gly Val Phe Thr Ser Ile Gly Lys Ala Leu His Gln Val Phe
705 710 715 720

Gly Ala Ile Tyr Gly Ala Ala Phe Ser Gly Val Ser Trp Thr Met Lys
725 730 735

Ile Leu Ile Gly Val Ile Ile Thr Trp Ile Gly Met Asn Ser Arg Ser
740 745 750

Thr Ser Leu Ser Val Thr Leu Val Leu Val Gly Ile Val Thr Leu Tyr

```
        755                 760                 765

Leu Gly Val Met Val Gln Ala Asp Ser Gly Cys Val Val Ser Trp Lys
    770                 775                 780

Asn Lys Glu Leu Lys Cys Gly Ser Gly Ile Phe Ile Thr Asp Asn Val
785                 790                 795                 800

His Thr Trp Thr Glu Gln Tyr Lys Phe Gln Pro Glu Ser Pro Ser Lys
                805                 810                 815

Leu Ala Ser Ala Ile Gln Lys Ala His Glu Glu Asp Ile Cys Gly Ile
            820                 825                 830

Arg Ser Val Thr Arg Leu Glu Asn Leu Met Trp Lys Gln Ile Thr Pro
        835                 840                 845

Glu Leu Asn His Ile Leu Ser Glu Asn Glu Val Lys Leu Thr Ile Met
    850                 855                 860

Thr Gly Asp Ile Lys Gly Ile Met Gln Ala Gly Lys Arg Ser Leu Arg
865                 870                 875                 880

Pro Gln Pro Thr Glu Leu Lys Tyr Ser Trp Lys Thr Trp Gly Lys Ala
                885                 890                 895

Lys Met Leu Ser Thr Glu Ser His Asn Gln Thr Phe Leu Ile Asp Gly
            900                 905                 910

Pro Glu Thr Ala Glu Cys Pro Asn Thr Asn Arg Ala Trp Asn Ser Leu
        915                 920                 925

Glu Val Glu Asp Tyr Gly Phe Gly Val Phe Thr Thr Asn Ile Trp Leu
    930                 935                 940

Lys Leu Lys Glu Lys Gln Asp Val Phe Cys Asp Ser Lys Leu Met Ser
945                 950                 955                 960

Ala Ala Ile Lys Asp Asn Arg Ala Val His Ala Asp Met Gly Tyr Trp
                965                 970                 975

Ile Glu Ser Ala Leu Asn Asp Thr Trp Lys Ile Glu Lys Ala Ser Phe
            980                 985                 990

Ile Glu Val Lys Asn Cys His Trp  Pro Lys Ser His Thr  Leu Trp Ser
        995                 1000                1005

Asn Gly  Val Leu Glu Ser Glu  Met Ile Ile Pro Lys  Asn Leu Ala
    1010                1015                1020

Gly Pro  Val Ser Gln His Asn  Tyr Arg Pro Gly Tyr  His Thr Gln
```

1025 1030 1035

Ile Thr Gly Pro Trp His Leu Gly Lys Leu Glu Met Asp Phe Asp
1040 1045 1050

Phe Cys Asp Gly Thr Thr Val Val Val Thr Glu Asp Cys Gly Asn
1055 1060 1065

Arg Gly Pro Ser Leu Arg Thr Thr Thr Ala Ser Gly Lys Leu Ile
1070 1075 1080

Thr Glu Trp Cys Cys Arg Ser Cys Thr Leu Pro Pro Leu Arg Tyr
1085 1090 1095

Arg Gly Glu Asp Gly Cys Trp Tyr Gly Met Glu Ile Arg Pro Leu
1100 1105 1110

Lys Glu Lys Glu Glu Asn Leu Val Asn Ser Leu Val Thr Ala Gly
1115 1120 1125

His Gly Gln Val Asp Asn Phe Ser Leu Gly Val Leu Gly Met Ala
1130 1135 1140

Leu Phe Leu Glu Glu Met Leu Arg Thr Arg Val Gly Thr Lys His
1145 1150 1155

Ala Ile Leu Leu Val Ala Val Ser Phe Val Thr Leu Ile Thr Gly
1160 1165 1170

Asn Met Ser Phe Arg Asp Leu Gly Arg Val Met Val Met Val Gly
1175 1180 1185

Ala Thr Met Thr Asp Asp Ile Gly Met Gly Val Thr Tyr Leu Ala
1190 1195 1200

Leu Leu Ala Ala Phe Lys Val Arg Pro Thr Phe Ala Ala Gly Leu
1205 1210 1215

Leu Leu Arg Lys Leu Thr Ser Lys Glu Leu Met Met Thr Thr Ile
1220 1225 1230

Gly Ile Val Leu Leu Ser Gln Ser Thr Ile Pro Glu Thr Ile Leu
1235 1240 1245

Glu Leu Thr Asp Ala Leu Ala Leu Gly Met Met Val Leu Lys Met
1250 1255 1260

Val Arg Asn Met Glu Lys Tyr Gln Leu Ala Val Thr Ile Met Ala
1265 1270 1275

Ile Leu Cys Val Pro Asn Ala Val Ile Leu Gln Asn Ala Trp Lys

1280 1285 1290

Val Ser Cys Thr Ile Leu Ala Val Val Ser Val Ser Pro Leu Phe
1295 1300 1305

Leu Thr Ser Ser Gln Gln Lys Thr Asp Trp Ile Pro Leu Ala Leu
1310 1315 1320

Thr Ile Lys Gly Leu Asn Pro Thr Ala Ile Phe Leu Thr Thr Leu
1325 1330 1335

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

Met Ala Val Gly Met Val Ser Ile Leu Ala Ser Ser Leu Leu Lys
1355 1360 1365

Asn Asp Ile Pro Met Thr Gly Pro Leu Val Ala Gly Gly Leu Leu
1370 1375 1380

Thr Val Cys Tyr Val Leu Thr Gly Arg Ser Ala Asp Leu Glu Leu
1385 1390 1395

Glu Arg Ala Ala Asp Val Lys Trp Glu Asp Gln Ala Glu Ile Ser
1400 1405 1410

Gly Ser Ser Pro Ile Leu Ser Ile Thr Ile Ser Glu Asp Gly Ser
1415 1420 1425

Met Ser Ile Lys Asn Glu Glu Glu Glu Gln Thr Leu Thr Ile Leu
1430 1435 1440

Ile Arg Thr Gly Leu Leu Val Ile Ser Gly Leu Phe Pro Val Ser
1445 1450 1455

Ile Pro Ile Thr Ala Ala Ala Trp Tyr Leu Trp Glu Val Lys Lys
1460 1465 1470

Gln Arg Ala Gly Val Leu Trp Asp Val Pro Ser Pro Pro Pro Met
1475 1480 1485

Gly Lys Ala Glu Leu Glu Asp Gly Ala Tyr Arg Ile Lys Gln Lys
1490 1495 1500

Gly Ile Leu Gly Tyr Ser Gln Ile Gly Ala Gly Val Tyr Lys Glu
1505 1510 1515

Gly Thr Phe His Thr Met Trp His Val Thr Arg Gly Ala Val Leu
1520 1525 1530

Met His Lys Gly Lys Arg Ile Glu Pro Ser Trp Ala Asp Val Lys

1535 1540 1545

Lys Asp Leu Ile Ser Tyr Gly Gly Gly Trp Lys Leu Glu Gly Glu
1550 1555 1560

Trp Lys Glu Gly Glu Glu Val Gln Val Leu Ala Leu Glu Pro Gly
1565 1570 1575

Lys Asn Pro Arg Ala Val Gln Thr Lys Pro Gly Leu Phe Lys Thr
1580 1585 1590

Asn Ala Gly Thr Ile Gly Ala Val Ser Leu Asp Phe Ser Pro Gly
1595 1600 1605

Thr Ser Gly Ser Pro Ile Ile Asp Lys Lys Gly Lys Val Val Gly
1610 1615 1620

Leu Tyr Gly Asn Gly Val Val Thr Arg Ser Gly Ala Tyr Val Ser
1625 1630 1635

Ala Ile Ala Gln Thr Glu Lys Ser Ile Glu Asp Asn Pro Glu Ile
1640 1645 1650

Glu Asp Asp Ile Phe Arg Lys Arg Arg Leu Thr Ile Met Asp Leu
1655 1660 1665

His Pro Gly Ala Gly Lys Thr Lys Arg Tyr Leu Pro Ala Ile Val
1670 1675 1680

Arg Glu Ala Ile Lys Arg Gly Leu Arg Thr Leu Ile Leu Ala Pro
1685 1690 1695

Thr Arg Val Val Ala Ala Glu Met Glu Glu Ala Leu Arg Gly Leu
1700 1705 1710

Pro Ile Arg Tyr Gln Thr Pro Ala Ile Arg Ala Val His Thr Gly
1715 1720 1725

Arg Glu Ile Val Asp Leu Met Cys His Ala Thr Phe Thr Met Arg
1730 1735 1740

Leu Leu Ser Pro Val Arg Val Pro Asn Tyr Asn Leu Ile Ile Met
1745 1750 1755

Asp Glu Ala His Phe Thr Asp Pro Ala Ser Ile Ala Ala Arg Gly
1760 1765 1770

Tyr Ile Ser Thr Arg Val Glu Met Gly Glu Ala Ala Gly Ile Phe
1775 1780 1785

Met Thr Ala Thr Pro Pro Gly Ser Arg Asp Pro Phe Pro Gln Ser

1790 1795 1800

Asn Ala Pro Ile Ile Asp Glu Glu Arg Glu Ile Pro Glu Arg Ser
1805 1810 1815

Trp Asn Ser Gly His Glu Trp Val Thr Asp Phe Lys Gly Lys Thr
1820 1825 1830

Val Trp Phe Val Pro Ser Ile Lys Ala Gly Asn Asp Ile Ala Ala
1835 1840 1845

Cys Leu Arg Lys Asn Gly Lys Lys Val Ile Gln Leu Ser Arg Lys
1850 1855 1860

Thr Phe Asp Ser Glu Tyr Val Lys Thr Arg Thr Asn Asp Trp Asp
1865 1870 1875

Phe Val Val Thr Thr Asp Ile Ser Glu Met Gly Ala Asn Phe Lys
1880 1885 1890

Ala Glu Arg Val Ile Asp Pro Arg Arg Cys Met Lys Pro Val Ile
1895 1900 1905

Leu Thr Asp Gly Glu Glu Arg Val Ile Leu Ala Gly Pro Met Pro
1910 1915 1920

Val Thr His Ser Ser Ala Ala Gln Arg Arg Gly Arg Ile Gly Arg
1925 1930 1935

Asn Pro Lys Asn Glu Asn Asp Gln Tyr Ile Tyr Met Gly Glu Pro
1940 1945 1950

Leu Glu Asn Asp Glu Asp Cys Ala His Trp Lys Glu Ala Lys Met
1955 1960 1965

Leu Leu Asp Asn Ile Asn Thr Pro Glu Gly Ile Ile Pro Ser Met
1970 1975 1980

Phe Glu Pro Glu Arg Glu Lys Val Asp Ala Ile Asp Gly Glu Tyr
1985 1990 1995

Arg Leu Arg Gly Glu Ala Arg Lys Thr Phe Val Asp Leu Met Arg
2000 2005 2010

Arg Gly Asp Leu Pro Val Trp Leu Ala Tyr Arg Val Ala Ala Glu
2015 2020 2025

Gly Ile Asn Tyr Ala Asp Arg Arg Trp Cys Phe Asp Gly Val Lys
2030 2035 2040

Asn Asn Gln Ile Leu Glu Glu Asn Val Glu Val Glu Ile Trp Thr

2045 2050 2055

Lys Glu Gly Glu Arg Lys Lys Leu Lys Pro Arg Trp Leu Asp Ala
2060 2065 2070

Arg Ile Tyr Ser Asp Pro Leu Ala Leu Lys Glu Phe Lys Glu Phe
2075 2080 2085

Ala Ala Gly Arg Lys Ser Leu Thr Leu Asn Leu Ile Thr Glu Met
2090 2095 2100

Gly Arg Leu Pro Thr Phe Met Thr Gln Lys Ala Arg Asp Ala Leu
2105 2110 2115

Asp Asn Leu Ala Val Leu His Thr Ala Glu Ala Gly Gly Arg Ala
2120 2125 2130

Tyr Asn His Ala Leu Ser Glu Leu Pro Glu Thr Leu Glu Thr Leu
2135 2140 2145

Leu Leu Leu Thr Leu Leu Ala Thr Val Thr Gly Gly Ile Phe Leu
2150 2155 2160

Phe Leu Met Ser Ala Arg Gly Ile Gly Lys Met Thr Leu Gly Met
2165 2170 2175

Cys Cys Ile Ile Thr Ala Ser Ile Leu Leu Trp Tyr Ala Gln Ile
2180 2185 2190

Gln Pro His Trp Ile Ala Ala Ser Ile Ile Leu Glu Phe Phe Leu
2195 2200 2205

Ile Val Leu Leu Ile Pro Glu Pro Glu Lys Gln Arg Thr Pro Gln
2210 2215 2220

Asp Asn Gln Leu Thr Tyr Val Val Ile Ala Ile Leu Thr Val Val
2225 2230 2235

Ala Ala Thr Met Ala Asn Glu Met Gly Phe Leu Glu Lys Thr Lys
2240 2245 2250

Lys Asp Leu Gly Leu Gly Ser Ile Ala Thr Gln Gln Pro Glu Ser
2255 2260 2265

Asn Ile Leu Asp Ile Asp Leu Arg Pro Ala Ser Ala Trp Thr Leu
2270 2275 2280

Tyr Ala Val Ala Thr Thr Phe Val Thr Pro Met Leu Arg His Ser
2285 2290 2295

Ile Glu Asn Ser Ser Val Asn Val Ser Leu Thr Ala Ile Ala Asn

2300 2305 2310

Gln Ala Thr Val Leu Met Gly Leu Gly Lys Gly Trp Pro Leu Ser
2315 2320 2325

Lys Met Asp Ile Gly Val Pro Leu Leu Ala Ile Gly Cys Tyr Ser
2330 2335 2340

Gln Val Asn Pro Ile Thr Leu Thr Ala Ala Leu Phe Leu Leu Val
2345 2350 2355

Ala His Tyr Ala Ile Ile Gly Pro Gly Leu Gln Ala Lys Ala Thr
2360 2365 2370

Arg Glu Ala Gln Lys Arg Ala Ala Ala Gly Ile Met Lys Asn Pro
2375 2380 2385

Thr Val Asp Gly Ile Thr Val Ile Asp Leu Asp Pro Ile Pro Tyr
2390 2395 2400

Asp Pro Lys Phe Glu Lys Gln Leu Gly Gln Val Met Leu Leu Val
2405 2410 2415

Leu Cys Val Thr Gln Val Leu Met Met Arg Thr Thr Trp Ala Leu
2420 2425 2430

Cys Glu Ala Leu Thr Leu Ala Thr Gly Pro Ile Ser Thr Leu Trp
2435 2440 2445

Glu Gly Asn Pro Gly Arg Phe Trp Asn Thr Thr Ile Ala Val Ser
2450 2455 2460

Met Ala Asn Ile Phe Arg Gly Ser Tyr Leu Ala Gly Ala Gly Leu
2465 2470 2475

Leu Phe Ser Ile Met Lys Asn Thr Thr Asn Thr Arg Arg Gly Thr
2480 2485 2490

Gly Asn Ile Gly Glu Thr Leu Gly Glu Lys Trp Lys Ser Arg Leu
2495 2500 2505

Asn Ala Leu Gly Lys Ser Glu Phe Gln Ile Tyr Lys Lys Ser Gly
2510 2515 2520

Ile Gln Glu Val Asp Arg Thr Leu Ala Lys Glu Gly Ile Lys Arg
2525 2530 2535

Gly Glu Thr Asp His His Ala Val Ser Arg Gly Ser Ala Lys Leu
2540 2545 2550

Arg Trp Phe Val Glu Arg Asn Met Val Thr Pro Glu Gly Lys Val

2555 2560 2565

Val Asp Leu Gly Cys Gly Arg Gly Gly Trp Ser Tyr Tyr Cys Gly
2570 2575 2580

Gly Leu Lys Asn Val Arg Glu Val Lys Gly Leu Thr Lys Gly Gly
2585 2590 2595

Pro Gly His Glu Glu Pro Ile Pro Met Ser Thr Tyr Gly Trp Asn
2600 2605 2610

Leu Val Arg Leu Gln Ser Gly Val Asp Val Phe Phe Ile Pro Pro
2615 2620 2625

Glu Lys Cys Asp Thr Leu Leu Cys Asp Ile Gly Glu Ser Ser Pro
2630 2635 2640

Asn Pro Thr Val Glu Ala Gly Arg Thr Leu Arg Val Leu Asn Leu
2645 2650 2655

Val Glu Asn Trp Leu Asn Asn Asn Thr Gln Phe Cys Ile Lys Val
2660 2665 2670

Leu Asn Pro Tyr Met Pro Ser Val Ile Glu Lys Met Glu Ala Leu
2675 2680 2685

Gln Arg Lys Tyr Gly Gly Ala Leu Val Arg Asn Pro Leu Ser Arg
2690 2695 2700

Asn Ser Thr His Glu Met Tyr Trp Val Ser Asn Ala Ser Gly Asn
2705 2710 2715

Ile Val Ser Ser Val Asn Met Ile Ser Arg Met Leu Ile Asn Arg
2720 2725 2730

Phe Thr Met Arg Tyr Lys Lys Ala Thr Tyr Glu Pro Asp Val Asp
2735 2740 2745

Leu Gly Ser Gly Thr Arg Asn Ile Gly Ile Glu Ser Glu Ile Pro
2750 2755 2760

Asn Leu Asp Ile Ile Gly Lys Arg Ile Glu Lys Ile Lys Gln Glu
2765 2770 2775

His Glu Thr Ser Trp His Tyr Asp Gln Asp His Pro Tyr Lys Thr
2780 2785 2790

Trp Ala Tyr His Gly Ser Tyr Glu Thr Lys Gln Thr Gly Ser Ala
2795 2800 2805

Ser Ser Met Val Asn Gly Val Val Arg Leu Leu Thr Lys Pro Trp

2810 2815 2820

Asp Val Val Pro Met Val Thr Gln Met Ala Met Thr Asp Thr Thr
2825 2830 2835

Pro Phe Gly Gln Gln Arg Val Phe Lys Glu Lys Val Asp Thr Arg
2840 2845 2850

Thr Gln Glu Pro Lys Glu Gly Thr Lys Lys Leu Met Lys Ile Thr
2855 2860 2865

Ala Glu Trp Leu Trp Lys Glu Leu Gly Lys Lys Lys Thr Pro Arg
2870 2875 2880

Met Cys Thr Arg Glu Glu Phe Thr Arg Lys Val Arg Ser Asn Ala
2885 2890 2895

Ala Leu Gly Ala Val Phe Thr Asp Glu Asn Lys Trp Lys Ser Ala
2900 2905 2910

Arg Glu Ala Val Glu Asp Ser Arg Phe Trp Glu Leu Val Asp Lys
2915 2920 2925

Glu Arg Asn Leu His Leu Glu Gly Lys Cys Glu Thr Cys Val Tyr
2930 2935 2940

Asn Met Met Gly Lys Arg Glu Lys Lys Leu Gly Glu Phe Gly Lys
2945 2950 2955

Ala Lys Gly Ser Arg Ala Ile Trp Tyr Met Trp Leu Gly Ala Arg
2960 2965 2970

Phe Leu Glu Phe Glu Ala Leu Gly Phe Leu Asn Glu Asp His Trp
2975 2980 2985

Phe Ser Arg Glu Asn Ser Leu Ser Gly Val Glu Gly Glu Gly Leu
2990 2995 3000

His Lys Leu Gly Tyr Ile Leu Arg Asp Val Ser Lys Lys Glu Gly
3005 3010 3015

Gly Ala Met Tyr Ala Asp Asp Thr Ala Gly Trp Asp Thr Arg Ile
3020 3025 3030

Thr Leu Glu Asp Leu Lys Asn Glu Glu Met Val Thr Asn His Met
3035 3040 3045

Glu Gly Glu His Lys Lys Leu Ala Glu Ala Ile Phe Lys Leu Thr
3050 3055 3060

Tyr Gln Asn Lys Val Val Arg Val Gln Arg Pro Thr Pro Arg Gly

3065 3070 3075

Thr Val Met Asp Ile Ile Ser Arg Arg Asp Gln Arg Gly Ser Gly
3080 3085 3090

Gln Val Gly Thr Tyr Gly Leu Asn Thr Phe Thr Asn Met Glu Ala
3095 3100 3105

Gln Leu Ile Arg Gln Met Glu Gly Glu Gly Val Phe Lys Ser Ile
3110 3115 3120

Gln His Leu Thr Ile Thr Glu Glu Ile Ala Val Gln Asn Trp Leu
3125 3130 3135

Ala Arg Val Gly Arg Glu Arg Leu Ser Arg Met Ala Ile Ser Gly
3140 3145 3150

Asp Asp Cys Val Val Lys Pro Leu Asp Asp Arg Phe Ala Ser Ala
3155 3160 3165

Leu Thr Ala Leu Asn Asp Met Gly Lys Ile Arg Lys Asp Ile Gln
3170 3175 3180

Gln Trp Glu Pro Ser Arg Gly Trp Asn Asp Trp Thr Gln Val Pro
3185 3190 3195

Phe Cys Ser His His Phe His Glu Leu Ile Met Lys Asp Gly Arg
3200 3205 3210

Val Leu Val Val Pro Cys Arg Asn Gln Asp Glu Leu Ile Gly Arg
3215 3220 3225

Ala Arg Ile Ser Gln Gly Ala Gly Trp Ser Leu Arg Glu Thr Ala
3230 3235 3240

Cys Leu Gly Lys Ser Tyr Ala Gln Met Trp Ser Leu Met Tyr Phe
3245 3250 3255

His Arg Arg Asp Leu Arg Leu Ala Ala Asn Ala Ile Cys Ser Ala
3260 3265 3270

Val Pro Ser His Trp Val Pro Thr Ser Arg Thr Thr Trp Ser Ile
3275 3280 3285

His Ala Lys His Glu Trp Met Thr Thr Glu Asp Met Leu Thr Val
3290 3295 3300

Trp Asn Arg Val Trp Ile Gln Glu Asn Pro Trp Met Glu Asp Lys
3305 3310 3315

Thr Pro Val Glu Ser Trp Glu Glu Ile Pro Tyr Leu Gly Lys Arg

3320 3325 3330

Glu Asp Gln Trp Cys Gly Ser Leu Ile Gly Leu Thr Ser Arg Ala
3335 3340 3345

Thr Trp Ala Lys Asn Ile Gln Ala Ala Ile Asn Gln Val Arg Ser
3350 3355 3360

Leu Ile Gly Asn Glu Glu Tyr Thr Asp Tyr Met Pro Ser Met Lys
3365 3370 3375

Arg Phe Arg Arg Glu Glu Glu Glu Ala Gly Val Leu Trp
3380 3385 3390

<210> 7
<211> 10717
<212> DNA
<213> Artificial Sequence

<220>
<223> Dengue virus serotype 3, BVS

<400> 7
agttgttagt ctacgtggac cgacaaagac agattctttg agggagctaa gctcaatgta 60

gttctaacag ttttttaatt agagagcaga tctctgatga ataaccaacg gaaaaaggcg 120

aaaaacacgc ctttcaatat gctgaaacgc gagagaaacc gcgtgtcgac tgtgcaacag 180

ctgacaaaga gattctcact tggaatgctg cagggacgag gaccattaaa actgttcatg 240

gccctggtgg cgttccttcg tttcctaaca atcccaccaa cagcagggat attgaagaga 300

tggggaacaa ttaaaaaatc aaaagctatt aatgttttga gagggttcag gaaagagatt 360

ggaaggatgc tgaacatctt gaataggaga cgcagatctg caggcatgat cattatgctg 420

attccaacag tgatggcgtt ccatttaacc acgcgtgatg gagagccgcg catgattgtg 480

gggaagaatg aaagaggaaa atccctactt ttcaagacag cctctggaat caacatgtgc 540

acactcatag ctatggatct gggagagatg tgtgatgaca cggtcactta caaatgcccc 600

cacattaccg aagtggagcc tgaagacatt gactgctggt gcaaccttac atcgacatgg 660

gtgacttatg gaacatgcaa tcaagctgga gagcatagac gcgataagag atcagtggcg 720

ttagctcccc atgttggcat gggactggac acacgcactc aaacctggat gtcggctgaa 780

ggagcttgga gacaagtcga gaaggtagag acatgggccc ttaggcaccc agggtttacc 840

atactagccc tatttcttgc ccattacata ggcacttcct tgacccagaa agtggttatt 900

tttatactat taatgctggt taccccatcc atgacaatga gatgtgtagg agtaggaaac 960

agagattttg tggaaggcct atcgggagct acgtgggttg acgtggtgct cgagcacggt 1020

gggtgtgtga ctaccatggc taagaacaag cccacgctgg acatagagct tcagaagacc 1080

gaggccaccc aactggcgac cctaaggaag ctatgcattg agggaaaaat taccaacata 1140

acaaccgact caagatgtcc cacccaaggg gaagcgattt tacctgagga gcaggaccag 1200

aactacgtgt gtaagcatac atacgtggac agaggctggg gaaacggttg tggtttgttt 1260
ggcaagggaa gcttggtgac atgcgcgaaa tttcaatgtt tagaatcaat agagggaaaa 1320
gtggtgcaac atgagaacct caaatacacc gtcatcatca cagtgcacac aggagaccaa 1380
caccaggtgg gaaatgaaac gcagggagtc acggctgaga taacacccca ggcatcaacc 1440
gctgaagcca ttttacctga atatggaacc ctcgggctag aatgctcacc acggacaggt 1500
ttggatttca atgaaatgat ctcattgaca atgaagaaca aagcatggat ggtacataga 1560
caatggttct ttgacttacc cctaccatgg acatcaggag cttcagcaga aacaccaact 1620
tggaacagga aagagcttct tgtgacattt aaaaatgcac atgcaaaaaa gcaagaagta 1680
gttgttcttg gatcacaaga gggagcaatg catacagcac tgacaggagc tacagagatc 1740
caaacctcag gaggcacaag tatctttgcg gggcacttaa aatgtagact caagatggac 1800
aaattggaac tcaaggggat gagctatgca atgtgcttga gtagctttgt gttgaagaaa 1860
gaagtctccg aaacgcagca tgggacaata ctcattaagg ttgagtacaa aggggaagat 1920
gcaccctgca agattccttt ctccacggag gatggacaag gaaaagctct caatggcaga 1980
ctgatcacag ccaatccagt ggtgaccaag aaggaggagc ctgtcaacat tgaggctgaa 2040
cctccttttg gagaaagtaa catagtaatt ggaattggag acaaagccct gaaaatcaac 2100
tggtacaaga agggaagctc gattgggaag atgttcgagg ccactgccag aggtgcaagg 2160
cgcatggcca tcttgggaga cacagcctgg gactttggat cagtgggtgg tgttttgaat 2220
tcattaggga aaatggtcca ccaaatattt gggagtgctt acacagccct atttggtgga 2280
gtctcctgga tgatgaaaat tggaataggt gtcctcttaa cctggatagg gttgaactca 2340
aaaaatactt ctatgtcatt ttcatgcatc gcggccggca ttgtgacact gtatttggga 2400
gtcatggtgc aggccgatag tggttgcgtt gtgagctgga aaaacaaaga actgaaatgt 2460
ggcagtggga ttttcatcac agacaacgtg cacacatgga cagaacaata caagttccaa 2520
ccagaatccc cttcaaaact agcttcagct atccagaaag cccatgaaga ggacatttgt 2580
ggaatccgct cagtaacaag actggagaat ctgatgtgga aacaaataac accagaattg 2640
aatcacattc tatcagaaaa tgaggtgaag ttaactatta tgacaggaga catcaaagga 2700
atcatgcagg caggaaaacg atctctgcgg cctcagccca ctgagctgaa gtattcatgg 2760
aaaacatggg gcaaagcaaa aatgctctct acagagtctc ataaccagac ctttctcatt 2820
gatggccccg aaacagcaga atgccccaac acaaatagag cttggaattc gttggaagtt 2880
gaagactatg gctttggagt attcaccacc aatatatggc taaaattgaa agaaaaacag 2940
gatgtattct gcgactcaaa actcatgtca gcggccataa aagacaacag agccgtccat 3000
gccgatatgg gttattggat agaaagtgca ctcaatgaca catggaagat agagaaagcc 3060
tctttcattg aagttaaaaa ctgccactgg ccaaaatcac acaccctctg gagcaatgga 3120
gtgctagaaa gtgagatgat aattccaaag aatctcgctg gaccagtgtc tcaacacaac 3180
tatagaccag gctaccatac acaaataaca ggaccatggc atctaggtaa gcttgagatg 3240

gactttgatt tctgtgatgg aacaacagtg gtagtgactg aggactgcgg aaatagagga 3300

ccctctttga gaacaaccac tgcctctgga aaactcataa cagaatggtg ctgccgatct 3360

tgcacattac caccgctaag atacagaggt gaggatgggt gctggtacgg gatggaaatc 3420

agaccattga aggagaaaga agagaatttg gtcaactcct tggtcacagc tggacatggg 3480

caggtcgaca acttttcact aggagtcttg ggaatggcat tgttcctgga ggaaatgctt 3540

aggacccgag taggaacgaa acatgcaata ctactagttg cagtttcttt tgtgacattg 3600

atcacaggga acatgtcctt tagagacctg ggaagagtga tggttatggt aggcgccact 3660

atgacggatg acataggtat gggcgtgact tatcttgccc tactagcagc cttcaaagtc 3720

agaccaactt ttgcagctgg actactcttg agaaagctga cctccaagga attgatgatg 3780

actactatag gaattgtact cctctcccag agcaccatac cagagaccat tcttgagttg 3840

actgatgcgt tagccttagg catgatggtc ctcaaaatgg tgagaaatat ggaaaagtat 3900

caattggcag tgactatcat ggctatcttg tgcgtcccaa acgcagtgat attacaaaac 3960

gcatggaaag tgagttgcac aatattggca gtggtgtccg tttccccact gttcttaaca 4020

tcctcacagc aaaaaacaga ttggatacca ttagcattga cgatcaaagg tctcaatcca 4080

acagctattt ttctaacaac cctctcaaga accagcaaga aaaggagctg gccattaaat 4140

gaggctatca tggcagtcgg gatggtgagc attttagcca gttctctcct aaaaaatgat 4200

attcccatga caggaccatt agtggctgga gggctcctca ctgtgtgcta cgtgctcact 4260

ggacgatcgg ccgatttgga actggagaga gcagccgatg tcaaatggga agaccaggca 4320

gagatatcag gaagcagtcc aatcctgtca ataacaatat cagaagatgg tagcatgtcg 4380

ataaaaaatg aagaggaaga acaaacactg accatactca ttagaacagg attgctggtg 4440

atctcaggac tttttcctgt atcaatacca atcacggcag cagcatggta cctgtgggaa 4500

gtgaagaaac aacgggccgg agtattgtgg gatgttcctt cacccccacc catgggaaag 4560

gctgaactgg aagatggagc ctatagaatt aagcaaaaag ggattcttgg atattcccag 4620

atcggagccg gagtttacaa agaaggaaca ttccatacaa tgtggcatgt cacacgtggc 4680

gctgttctaa tgcataaagg aaagaggatt gaaccatcat gggcggacgt caagaaagac 4740

ctaatatcat atggaggagg ctggaagtta gaaggagaat ggaaggaagg agaagaagtc 4800

caggtattgg cactggagcc tggaaaaaat ccaagagccg tccaaacgaa acctggtctt 4860

ttcaaaacca acgccggaac aataggtgct gtatctctgg acttttctcc tggaacgtca 4920

ggatctccaa ttatcgacaa aaaaggaaaa gttgtgggtc tttatggtaa tggtgttgtt 4980

acaaggagtg gagcatatgt gagtgctata gcccagactg aaaaaagcat tgaagacaac 5040

ccagagatcg aagatgacat tttccgaaag agaagactga ccatcatgga cctccaccca 5100

ggagcgggaa agacgaagag ataccttccg gccatagtca gagaagctat aaaacggggt 5160

ttgagaacat taatcttggc ccccactaga gttgtggcag ctgaaatgga ggaagccctt 5220

agaggacttc caataagata ccagacccca gccatcagag ctgtgcacac cgggcgggag 5280

attgtggacc taatgtgtca tgccacattt accatgaggc tgctatcacc agttagagtg 5340

ccaaactaca acctgattat catggacgaa gcccatttca cagacccagc aagtatagca 5400

gctagaggat acatctcaac tcgagtggag atgggtgagg cagctgggat ttttatgaca 5460

gccactcccc cgggaagcag agacccattt cctcagagca atgcaccaat catagatgaa 5520

gaaagagaaa tccctgaacg ctcgtggaat tccggacatg aatgggtcac ggattttaaa 5580

gggaagactg tttggttcgt tccaagtata aaagcaggaa atgatatagc agcttgcctg 5640

aggaaaaatg gaaagaaagt gatacaactc agtaggaaga cctttgattc tgagtatgtc 5700

aagactagaa ccaatgattg ggacttcgtg gttacaactg acatttcaga aatgggtgcc 5760

aatttcaagg ctgagagggt tatagacccc agacgctgca tgaaaccagt catactaaca 5820

gatggtgaag agcgggtgat tctggcagga cctatgccag tgacccactc tagtgcagca 5880

caaagaagag ggagaatagg aagaaatcca aaaaatgaga atgaccagta catatacatg 5940

ggggaacctc tggaaaatga tgaagactgt gcacactgga aagaagctaa aatgctccta 6000

gataacatca acacgccaga aggaatcatt cctagcatgt tcgaaccaga gcgtgaaaag 6060

gtggatgcca ttgatggcga ataccgcttg agaggagaag caaggaaaac ctttgtagac 6120

ttaatgagaa gaggagacct accagtctgg ttggcctaca gagtggcagc tgaaggcatc 6180

aactacgcag acagaaggtg gtgttttgat ggagtcaaga acaaccaaat cctagaagaa 6240

aacgtggaag ttgaaatctg gacaaaagaa ggggaaagga agaaattgaa acccagatgg 6300

ttggatgcta ggatctattc tgacccactg gcgctaaaag aatttaagga atttgcagcc 6360

ggaagaaagt ctctgaccct gaacctaatc acagaaatgg gtaggctccc aaccttcatg 6420

actcagaagg caagaaacgc actggacaac ttagcagtgc tgcacacggc tgaggcaggt 6480

ggaagggcgt acaaccatgc tctcagtgaa ctgccggaga ccctggagac attgctttta 6540

ctgacacttc tggctacagt cacgggaggg atctttttat tcttgatgag cgcaaggggc 6600

atagggaaga tgaccctggg aatgtgctgc ataatcacgg ctagcatcct cctatggtac 6660

gcacaaatac agccacactg gatagcagct tcaataatac tggagttttt tctcatagtt 6720

ttgcttattc cagaacctga aaaacagaga acaccccaag acaaccaact gacctacgtt 6780

gtcatagcca tcctcacagt ggtggccgca accatggcaa acgagatggg tttcctagaa 6840

aaaacgaaga aagatctcgg attgggaagc attgcaaccc agcaacccga gagcaacatc 6900

ctggacatag atctacgtcc tgcatcagca tggacgctgt atgccgtggc cacaacattt 6960

gttacaccaa tgttgagaca tagcattgaa aattcctcag tgaatgtgtc cctaacagct 7020

atagccaacc aagccacagt gttaatgggt ctcgggaaag gatggccatt gtcaaagatg 7080

gacatcggag ttccccttct cgccattgga tgctactcac aagtcaaccc cataactctc 7140

acagcagctc ttttcttatt ggtagcacat tatgccatca tagggccagg actccaagca 7200

aaagcaacca gagaagctca gaaaagagca gcggcgggca tcatgaaaaa cccaactgtc 7260

gatggaataa cagtgattga cctagatcca ataccttatg atccaaagtt tgaaaagcag 7320

ttgggacaag taatgctcct agtcctctgc gtgactcaag tattgatgat gaggactaca 7380

tgggctctgt gtgaggcttt aaccttagct accgggccca tctccacatt gtgggaagga 7440

aatccaggga ggttttggaa cactaccatt gcggtgtcaa tggctaacat ttttagaggg 7500

agttacttgg ccggagctgg acttctcttt tctattatga agaacacaac caacacaaga 7560

aggggaactg gcaacatagg agagacgctt ggagagaaat ggaaaagccg attgaacgcg 7620

ttgggaaaaa gtgaattcca gatctacaag aaaagtggaa tccaggaagt ggatagaacc 7680

ttagcaaaag aaggcattaa aagaggagaa acggaccatc acgctgtgtc gcgaggctca 7740

gcaaaactga gatggttcgt tgagagaaac atggtcacac cagaagggaa agtagtggac 7800

ctcggttgtg gcagaggagg ctggtcatac tattgtggag gactaaagaa tgtaagagaa 7860

gtcaaaggcc taacaaaagg aggaccagga cacgaagaac ccatccccat gtcaacatat 7920

gggtggaatc tagtgcgtct tcaaagtgga gttgacgttt tcttcatccc gccagaaaag 7980

tgtgacacat tattgtgtga cataggggag tcatcaccaa atcccacagt ggaagcagga 8040

cgaacactca gagtccttaa cttagtagaa aattggttga acaacaacac tcaattttgc 8100

ataaaggttc tcaacccata tatgccctca gtcatagaaa aaatggaagc actacaaagg 8160

aaatatggag gagccttagt gaggaatcca ctctcacgaa actccacaca tgagatgtac 8220

tgggtatcca atgcttccgg gaacatagtg tcatcagtga acatgatttc aaggatgttg 8280

atcaacagat ttacaatgag atacaagaaa gccacttacg agccggatgt tgacctcgga 8340

agcggaaccc gtaacatcgg gattgaaagt gagataccaa acctagatat aattgggaaa 8400

agaatagaaa aaataaagca agagcatgaa acatcatggc actatgacca agaccaccca 8460

tacaaaacgt gggcatacca tggtagctat gaaacaaaac agactggatc agcatcatcc 8520

atggtcaacg gagtggtcag gctgctgaca aaaccttggg acgtcgtccc catggtgaca 8580

cagatggcaa tgacagacac gactccattt ggacaacagc gcgtttttaa agagaaagtg 8640

gacacgagaa cccaagaacc gaaagaaggc acgaagaaac taatgaaaat aacagcagag 8700

tggctttgga aagaattagg gaagaaaaag acacccagga tgtgcaccag agaagaattc 8760

acaagaaagg tgagaagcaa tgcagccttg ggggccatat tcactgatga gaacaagtgg 8820

aagtcggcac gtgaggctgt tgaagatagt aggttttggg agctggttga caaggaaagg 8880

aatctccatc ttgaaggaaa gtgtgaaaca tgtgtgtaca acatgatggg aaaaagagag 8940

aagaagctag gggaattcgg caaggcaaaa ggcagcagag ccatatggta catgtggctt 9000

ggagcacgct tcttagagtt tgaagcccta ggattcttaa atgaagatca ctggttctcc 9060

agagagaact ccctgagtgg agtggaagga gaagggctgc acaagctagg ttacattcta 9120

agagacgtga gcaagaaaga gggaggagca atgtatgccg atgacaccgc aggatgggat 9180

acaagaatca cactagaaga cctaaaaaat gaagaaatgg taacaaacca catggaagga 9240

gaacacaaga aactagccga ggccattttc aaactaacgt accaaaacaa ggtggtgcgt 9300

gtgcaaagac caacaccaag aggcacagta atggacatca tatcgagaag agaccaaaga 9360

ggtagtggac aagttggcac ctatggactc aatactttca ccaatatgga agcccaacta 9420

atcagacaga tggagggaga aggagtcttt aaaagcattc agcacctaac aatcacagaa 9480

gaaatcgctg tgcaaaactg gttagcaaga gtggggcgcg aaaggttatc aagaatggcc 9540

atcagtggag atgattgtgt tgtgaaacct ttagatgaca ggttcgcaag cgctttaaca 9600

gctctaaatg acatgggaaa gattaggaaa gacatacaac aatgggaacc ttcaagagga 9660

tggaatgatt ggacacaagt gcccttctgt tcacaccatt tccatgagtt aatcatgaaa 9720

gacggtcgcg tactcgttgt tccatgtaga aaccaagatg aactgattgg cagagcccga 9780

atctcccaag gagcagggtg gtctttgcgg gagacggcct gtttggggaa gtcttacgcc 9840

caaatgtgga gcttgatgta cttccacaga cgcgacctca ggctggcggc aaatgctatt 9900

tgctcggcag taccatcaca ttgggttcca acaagtcgaa caacctggtc catacatgct 9960

aaacatgaat ggatgacaac ggaagacatg ctgacagtct ggaacagggt gtggattcaa 10020

gaaaacccat ggatggaaga caaaactcca gtggaatcat gggaggaaat cccatacttg 10080

gggaaaagag aagaccaatg gtgcggctca ttgattgggt taacaagcag ggccacctgg 10140

gcaaagaaca tccaagcagc aataaatcaa gttagatccc ttataggcaa tgaagaatac 10200

acagattaca tgccatccat gaaaagattc agaagagaag aggaagaagc aggagttctg 10260

tggtagaaag caaaactaac atgaaacaag gctagaagtc aggtcggatt aagccatagt 10320

acggaaaaaa ctatgctacc tgtgagcccc gtccaaggac gttaaaagaa gtcaggccat 10380

cataaatgcc atagcttgag taaactatgc agcctgtagc tccacctgag aaggtgtaaa 10440

aaatccggga ggccacaaac catggaagct gtacgcatgg cgtagtggac tagcggttag 10500

aggagacccc tcccttacaa atcgcagcaa caatgggggc ccaaggcgag atgaagctgt 10560

agtctcgctg gaaggactag aggttagagg agacccccc gaaacaaaaa acagcatatt 10620

gacgctggga aagaccagag atcctgctgt ctcctcagca tcattccagg cacagaacgc 10680

cagaaaatgg aatggtgctg ttgaatcaac aggttct 10717

<210> 8
<211> 3389
<212> PRT
<213> Artificial Sequence

<220>
<223> Dengue virus serotype 3, BVS

<400> 8

Met Asn Asn Gln Arg Lys Lys Ala Lys Asn Thr Pro Phe Asn Met Leu
1 5 10 15

Lys Arg Glu Arg Asn Arg Val Ser Thr Val Gln Gln Leu Thr Lys Arg
20 25 30

Phe Ser Leu Gly Met Leu Gln Gly Arg Gly Pro Leu Lys Leu Phe Met
35 40 45

```
Ala Leu Val Ala Phe Leu Arg Phe Leu Thr Ile Pro Pro Thr Ala Gly
    50                  55                  60

Ile Leu Lys Arg Trp Gly Thr Ile Lys Lys Ser Lys Ala Ile Asn Val
65                  70                  75                  80

Leu Arg Gly Phe Arg Lys Glu Ile Gly Arg Met Leu Asn Ile Leu Asn
                85                  90                  95

Arg Arg Arg Arg Ser Ala Gly Met Ile Ile Met Leu Ile Pro Thr Val
            100                 105                 110

Met Ala Phe His Leu Thr Thr Arg Asp Gly Glu Pro Arg Met Ile Val
        115                 120                 125

Gly Lys Asn Glu Arg Gly Lys Ser Leu Leu Phe Lys Thr Ala Ser Gly
    130                 135                 140

Ile Asn Met Cys Thr Leu Ile Ala Met Asp Leu Gly Glu Met Cys Asp
145                 150                 155                 160

Asp Thr Val Thr Tyr Lys Cys Pro His Ile Thr Glu Val Glu Pro Glu
                165                 170                 175

Asp Ile Asp Cys Trp Cys Asn Leu Thr Ser Thr Trp Val Thr Tyr Gly
            180                 185                 190

Thr Cys Asn Gln Ala Gly Glu His Arg Arg Asp Lys Arg Ser Val Ala
        195                 200                 205

Leu Ala Pro His Val Gly Met Gly Leu Asp Thr Arg Thr Gln Thr Trp
    210                 215                 220

Met Ser Ala Glu Gly Ala Trp Arg Gln Val Glu Lys Val Glu Thr Trp
225                 230                 235                 240

Ala Leu Arg His Pro Gly Phe Thr Ile Leu Ala Leu Phe Leu Ala His
                245                 250                 255

Tyr Ile Gly Thr Ser Leu Thr Gln Lys Val Val Ile Phe Ile Leu Leu
            260                 265                 270

Met Leu Val Thr Pro Ser Met Thr Met Arg Cys Val Gly Val Gly Asn
        275                 280                 285

Arg Asp Phe Val Glu Gly Leu Ser Gly Ala Thr Trp Val Asp Val Val
    290                 295                 300

Leu Glu His Gly Gly Cys Val Thr Thr Met Ala Lys Asn Lys Pro Thr
305                 310                 315                 320
```

```
Leu Asp Ile Glu Leu Gln Lys Thr Glu Ala Thr Gln Leu Ala Thr Leu
                325                 330                 335

Arg Lys Leu Cys Ile Glu Gly Lys Ile Thr Asn Ile Thr Thr Asp Ser
            340                 345                 350

Arg Cys Pro Thr Gln Gly Glu Ala Ile Leu Pro Glu Glu Gln Asp Gln
        355                 360                 365

Asn Tyr Val Cys Lys His Thr Tyr Val Asp Arg Gly Trp Gly Asn Gly
    370                 375                 380

Cys Gly Leu Phe Gly Lys Gly Ser Leu Val Thr Cys Ala Lys Phe Gln
385                 390                 395                 400

Cys Leu Glu Ser Ile Glu Gly Lys Val Val Gln His Glu Asn Leu Lys
                405                 410                 415

Tyr Thr Val Ile Ile Thr Val His Thr Gly Asp Gln His Gln Val Gly
            420                 425                 430

Asn Glu Thr Gln Gly Val Thr Ala Glu Ile Thr Pro Gln Ala Ser Thr
        435                 440                 445

Ala Glu Ala Ile Leu Pro Glu Tyr Gly Thr Leu Gly Leu Glu Cys Ser
    450                 455                 460

Pro Arg Thr Gly Leu Asp Phe Asn Glu Met Ile Ser Leu Thr Met Lys
465                 470                 475                 480

Asn Lys Ala Trp Met Val His Arg Gln Trp Phe Phe Asp Leu Pro Leu
                485                 490                 495

Pro Trp Thr Ser Gly Ala Ser Ala Glu Thr Pro Thr Trp Asn Arg Lys
            500                 505                 510

Glu Leu Leu Val Thr Phe Lys Asn Ala His Ala Lys Lys Gln Glu Val
        515                 520                 525

Val Val Leu Gly Ser Gln Glu Gly Ala Met His Thr Ala Leu Thr Gly
    530                 535                 540

Ala Thr Glu Ile Gln Thr Ser Gly Gly Thr Ser Ile Phe Ala Gly His
545                 550                 555                 560

Leu Lys Cys Arg Leu Lys Met Asp Lys Leu Glu Leu Lys Gly Met Ser
                565                 570                 575

Tyr Ala Met Cys Leu Ser Ser Phe Val Leu Lys Lys Glu Val Ser Glu
            580                 585                 590
```

Thr Gln His Gly Thr Ile Leu Ile Lys Val Glu Tyr Lys Gly Glu Asp
595 600 605

Ala Pro Cys Lys Ile Pro Phe Ser Thr Glu Asp Gly Gln Gly Lys Ala
610 615 620

Leu Asn Gly Arg Leu Ile Thr Ala Asn Pro Val Val Thr Lys Lys Glu
625 630 635 640

Glu Pro Val Asn Ile Glu Ala Glu Pro Pro Phe Gly Glu Ser Asn Ile
645 650 655

Val Ile Gly Ile Gly Asp Lys Ala Leu Lys Ile Asn Trp Tyr Lys Lys
660 665 670

Gly Ser Ser Ile Gly Lys Met Phe Glu Ala Thr Ala Arg Gly Ala Arg
675 680 685

Arg Met Ala Ile Leu Gly Asp Thr Ala Trp Asp Phe Gly Ser Val Gly
690 695 700

Gly Val Leu Asn Ser Leu Gly Lys Met Val His Gln Ile Phe Gly Ser
705 710 715 720

Ala Tyr Thr Ala Leu Phe Gly Gly Val Ser Trp Met Met Lys Ile Gly
725 730 735

Ile Gly Val Leu Leu Thr Trp Ile Gly Leu Asn Ser Lys Asn Thr Ser
740 745 750

Met Ser Phe Ser Cys Ile Ala Ala Gly Ile Val Thr Leu Tyr Leu Gly
755 760 765

Val Met Val Gln Ala Asp Ser Gly Cys Val Val Ser Trp Lys Asn Lys
770 775 780

Glu Leu Lys Cys Gly Ser Gly Ile Phe Ile Thr Asp Asn Val His Thr
785 790 795 800

Trp Thr Glu Gln Tyr Lys Phe Gln Pro Glu Ser Pro Ser Lys Leu Ala
805 810 815

Ser Ala Ile Gln Lys Ala His Glu Glu Asp Ile Cys Gly Ile Arg Ser
820 825 830

Val Thr Arg Leu Glu Asn Leu Met Trp Lys Gln Ile Thr Pro Glu Leu
835 840 845

Asn His Ile Leu Ser Glu Asn Glu Val Lys Leu Thr Ile Met Thr Gly
850 855 860

```
Asp Ile Lys Gly Ile Met Gln Ala Gly Lys Arg Ser Leu Arg Pro Gln
865                 870                 875                 880

Pro Thr Glu Leu Lys Tyr Ser Trp Lys Thr Trp Gly Lys Ala Lys Met
                885                 890                 895

Leu Ser Thr Glu Ser His Asn Gln Thr Phe Leu Ile Asp Gly Pro Glu
            900                 905                 910

Thr Ala Glu Cys Pro Asn Thr Asn Arg Ala Trp Asn Ser Leu Glu Val
        915                 920                 925

Glu Asp Tyr Gly Phe Gly Val Phe Thr Thr Asn Ile Trp Leu Lys Leu
    930                 935                 940

Lys Glu Lys Gln Asp Val Phe Cys Asp Ser Lys Leu Met Ser Ala Ala
945                 950                 955                 960

Ile Lys Asp Asn Arg Ala Val His Ala Asp Met Gly Tyr Trp Ile Glu
                965                 970                 975

Ser Ala Leu Asn Asp Thr Trp Lys Ile Glu Lys Ala Ser Phe Ile Glu
            980                 985                 990

Val Lys Asn Cys His Trp Pro Lys  Ser His Thr Leu Trp  Ser Asn Gly
        995                 1000                1005

Val Leu  Glu Ser Glu Met Ile  Ile Pro Lys Asn Leu  Ala Gly Pro
    1010                1015                1020

Val Ser  Gln His Asn Tyr Arg  Pro Gly Tyr His Thr  Gln Ile Thr
    1025                1030                1035

Gly Pro  Trp His Leu Gly Lys  Leu Glu Met Asp Phe  Asp Phe Cys
    1040                1045                1050

Asp Gly  Thr Thr Val Val Val  Thr Glu Asp Cys Gly  Asn Arg Gly
    1055                1060                1065

Pro Ser  Leu Arg Thr Thr Thr  Ala Ser Gly Lys Leu  Ile Thr Glu
    1070                1075                1080

Trp Cys  Cys Arg Ser Cys Thr  Leu Pro Pro Leu Arg  Tyr Arg Gly
    1085                1090                1095

Glu Asp  Gly Cys Trp Tyr Gly  Met Glu Ile Arg Pro  Leu Lys Glu
    1100                1105                1110

Lys Glu  Glu Asn Leu Val Asn  Ser Leu Val Thr Ala  Gly His Gly
    1115                1120                1125
```

```
Gln Val  Asp Asn Phe Ser Leu  Gly Val Leu Gly Met  Ala Leu Phe
    1130                1135                1140

Leu Glu  Glu Met Leu Arg Thr  Arg Val Gly Thr Lys  His Ala Ile
    1145                1150                1155

Leu Leu  Val Ala Val Ser Phe  Val Thr Leu Ile Thr  Gly Asn Met
    1160                1165                1170

Ser Phe  Arg Asp Leu Gly Arg  Val Met Val Met Val  Gly Ala Thr
    1175                1180                1185

Met Thr  Asp Asp Ile Gly Met  Gly Val Thr Tyr Leu  Ala Leu Leu
    1190                1195                1200

Ala Ala  Phe Lys Val Arg Pro  Thr Phe Ala Ala Gly  Leu Leu Leu
    1205                1210                1215

Arg Lys  Leu Thr Ser Lys Glu  Leu Met Met Thr Thr  Ile Gly Ile
    1220                1225                1230

Val Leu  Leu Ser Gln Ser Thr  Ile Pro Glu Thr Ile  Leu Glu Leu
    1235                1240                1245

Thr Asp  Ala Leu Ala Leu Gly  Met Met Val Leu Lys  Met Val Arg
    1250                1255                1260

Asn Met  Glu Lys Tyr Gln Leu  Ala Val Thr Ile Met  Ala Ile Leu
    1265                1270                1275

Cys Val  Pro Asn Ala Val Ile  Leu Gln Asn Ala Trp  Lys Val Ser
    1280                1285                1290

Cys Thr  Ile Leu Ala Val Val  Ser Val Ser Pro Leu  Phe Leu Thr
    1295                1300                1305

Ser Ser  Gln Gln Lys Thr Asp  Trp Ile Pro Leu Ala  Leu Thr Ile
    1310                1315                1320

Lys Gly  Leu Asn Pro Thr Ala  Ile Phe Leu Thr Thr  Leu Ser Arg
    1325                1330                1335

Thr Ser  Lys Lys Arg Ser Trp  Pro Leu Asn Glu Ala  Ile Met Ala
    1340                1345                1350

Val Gly  Met Val Ser Ile Leu  Ala Ser Ser Leu Leu  Lys Asn Asp
    1355                1360                1365

Ile Pro  Met Thr Gly Pro Leu  Val Ala Gly Gly Leu  Leu Thr Val
    1370                1375                1380
```

```
Cys Tyr  Val Leu Thr Gly Arg  Ser Ala Asp Leu Glu  Leu Glu Arg
    1385                 1390                 1395

Ala Ala  Asp Val Lys Trp Glu  Asp Gln Ala Glu Ile  Ser Gly Ser
    1400                 1405                 1410

Ser Pro  Ile Leu Ser Ile Thr  Ile Ser Glu Asp Gly  Ser Met Ser
    1415                 1420                 1425

Ile Lys  Asn Glu Glu Glu Glu  Gln Thr Leu Thr Ile  Leu Ile Arg
    1430                 1435                 1440

Thr Gly  Leu Leu Val Ile Ser  Gly Leu Phe Pro Val  Ser Ile Pro
    1445                 1450                 1455

Ile Thr  Ala Ala Ala Trp Tyr  Leu Trp Glu Val Lys  Lys Gln Arg
    1460                 1465                 1470

Ala Gly  Val Leu Trp Asp Val  Pro Ser Pro Pro Pro  Met Gly Lys
    1475                 1480                 1485

Ala Glu  Leu Glu Asp Gly Ala  Tyr Arg Ile Lys Gln  Lys Gly Ile
    1490                 1495                 1500

Leu Gly  Tyr Ser Gln Ile Gly  Ala Gly Val Tyr Lys  Glu Gly Thr
    1505                 1510                 1515

Phe His  Thr Met Trp His Val  Thr Arg Gly Ala Val  Leu Met His
    1520                 1525                 1530

Lys Gly  Lys Arg Ile Glu Pro  Ser Trp Ala Asp Val  Lys Lys Asp
    1535                 1540                 1545

Leu Ile  Ser Tyr Gly Gly Gly  Trp Lys Leu Glu Gly  Glu Trp Lys
    1550                 1555                 1560

Glu Gly  Glu Glu Val Gln Val  Leu Ala Leu Glu Pro  Gly Lys Asn
    1565                 1570                 1575

Pro Arg  Ala Val Gln Thr Lys  Pro Gly Leu Phe Lys  Thr Asn Ala
    1580                 1585                 1590

Gly Thr  Ile Gly Ala Val Ser  Leu Asp Phe Ser Pro  Gly Thr Ser
    1595                 1600                 1605

Gly Ser  Pro Ile Ile Asp Lys  Lys Gly Lys Val Val  Gly Leu Tyr
    1610                 1615                 1620

Gly Asn  Gly Val Val Thr Arg  Ser Gly Ala Tyr Val  Ser Ala Ile
    1625                 1630                 1635
```

```
Ala Gln  Thr Glu Lys Ser Ile  Glu Asp Asn Pro Glu  Ile Glu Asp
    1640                1645                1650

Asp Ile  Phe Arg Lys Arg Arg  Leu Thr Ile Met Asp  Leu His Pro
    1655                1660                1665

Gly Ala  Gly Lys Thr Lys Arg  Tyr Leu Pro Ala Ile  Val Arg Glu
    1670                1675                1680

Ala Ile  Lys Arg Gly Leu Arg  Thr Leu Ile Leu Ala  Pro Thr Arg
    1685                1690                1695

Val Val  Ala Ala Glu Met Glu  Glu Ala Leu Arg Gly  Leu Pro Ile
    1700                1705                1710

Arg Tyr  Gln Thr Pro Ala Ile  Arg Ala Val His Thr  Gly Arg Glu
    1715                1720                1725

Ile Val  Asp Leu Met Cys His  Ala Thr Phe Thr Met  Arg Leu Leu
    1730                1735                1740

Ser Pro  Val Arg Val Pro Asn  Tyr Asn Leu Ile Ile  Met Asp Glu
    1745                1750                1755

Ala His  Phe Thr Asp Pro Ala  Ser Ile Ala Ala Arg  Gly Tyr Ile
    1760                1765                1770

Ser Thr  Arg Val Glu Met Gly  Glu Ala Ala Gly Ile  Phe Met Thr
    1775                1780                1785

Ala Thr  Pro Pro Gly Ser Arg  Asp Pro Phe Pro Gln  Ser Asn Ala
    1790                1795                1800

Pro Ile  Ile Asp Glu Glu Arg  Glu Ile Pro Glu Arg  Ser Trp Asn
    1805                1810                1815

Ser Gly  His Glu Trp Val Thr  Asp Phe Lys Gly Lys  Thr Val Trp
    1820                1825                1830

Phe Val  Pro Ser Ile Lys Ala  Gly Asn Asp Ile Ala  Ala Cys Leu
    1835                1840                1845

Arg Lys  Asn Gly Lys Lys Val  Ile Gln Leu Ser Arg  Lys Thr Phe
    1850                1855                1860

Asp Ser  Glu Tyr Val Lys Thr  Arg Thr Asn Asp Trp  Asp Phe Val
    1865                1870                1875

Val Thr  Thr Asp Ile Ser Glu  Met Gly Ala Asn Phe  Lys Ala Glu
    1880                1885                1890
```

```
Arg Val  Ile Asp Pro Arg Arg  Cys Met Lys Pro Val  Ile Leu Thr
    1895                1900                1905

Asp Gly  Glu Glu Arg Val Ile  Leu Ala Gly Pro Met  Pro Val Thr
    1910                1915                1920

His Ser  Ser Ala Ala Gln Arg  Arg Gly Arg Ile Gly  Arg Asn Pro
    1925                1930                1935

Lys Asn  Glu Asn Asp Gln Tyr  Ile Tyr Met Gly Glu  Pro Leu Glu
    1940                1945                1950

Asn Asp  Glu Asp Cys Ala His  Trp Lys Glu Ala Lys  Met Leu Leu
    1955                1960                1965

Asp Asn  Ile Asn Thr Pro Glu  Gly Ile Ile Pro Ser  Met Phe Glu
    1970                1975                1980

Pro Glu  Arg Glu Lys Val Asp  Ala Ile Asp Gly Glu  Tyr Arg Leu
    1985                1990                1995

Arg Gly  Glu Ala Arg Lys Thr  Phe Val Asp Leu Met  Arg Arg Gly
    2000                2005                2010

Asp Leu  Pro Val Trp Leu Ala  Tyr Arg Val Ala Ala  Glu Gly Ile
    2015                2020                2025

Asn Tyr  Ala Asp Arg Arg Trp  Cys Phe Asp Gly Val  Lys Asn Asn
    2030                2035                2040

Gln Ile  Leu Glu Glu Asn Val  Glu Val Glu Ile Trp  Thr Lys Glu
    2045                2050                2055

Gly Glu  Arg Lys Lys Leu Lys  Pro Arg Trp Leu Asp  Ala Arg Ile
    2060                2065                2070

Tyr Ser  Asp Pro Leu Ala Leu  Lys Glu Phe Lys Glu  Phe Ala Ala
    2075                2080                2085

Gly Arg  Lys Ser Leu Thr Leu  Asn Leu Ile Thr Glu  Met Gly Arg
    2090                2095                2100

Leu Pro  Thr Phe Met Thr Gln  Lys Ala Arg Asn Ala  Leu Asp Asn
    2105                2110                2115

Leu Ala  Val Leu His Thr Ala  Glu Ala Gly Gly Arg  Ala Tyr Asn
    2120                2125                2130

His Ala  Leu Ser Glu Leu Pro  Glu Thr Leu Glu Thr  Leu Leu Leu
    2135                2140                2145
```

```
Leu Thr  Leu Leu Ala Thr Val  Thr Gly Gly Ile Phe  Leu Phe Leu
    2150                2155                2160

Met Ser  Ala Arg Gly Ile Gly  Lys Met Thr Leu Gly  Met Cys Cys
    2165                2170                2175

Ile Ile  Thr Ala Ser Ile Leu  Leu Trp Tyr Ala Gln  Ile Gln Pro
    2180                2185                2190

His Trp  Ile Ala Ala Ser Ile  Ile Leu Glu Phe Phe  Leu Ile Val
    2195                2200                2205

Leu Leu  Ile Pro Glu Pro Glu  Lys Gln Arg Thr Pro  Gln Asp Asn
    2210                2215                2220

Gln Leu  Thr Tyr Val Val Ile  Ala Ile Leu Thr Val  Val Ala Ala
    2225                2230                2235

Thr Met  Ala Asn Glu Met Gly  Phe Leu Glu Lys Thr  Lys Lys Asp
    2240                2245                2250

Leu Gly  Leu Gly Ser Ile Ala  Thr Gln Gln Pro Glu  Ser Asn Ile
    2255                2260                2265

Leu Asp  Ile Asp Leu Arg Pro  Ala Ser Ala Trp Thr  Leu Tyr Ala
    2270                2275                2280

Val Ala  Thr Thr Phe Val Thr  Pro Met Leu Arg His  Ser Ile Glu
    2285                2290                2295

Asn Ser  Ser Val Asn Val Ser  Leu Thr Ala Ile Ala  Asn Gln Ala
    2300                2305                2310

Thr Val  Leu Met Gly Leu Gly  Lys Gly Trp Pro Leu  Ser Lys Met
    2315                2320                2325

Asp Ile  Gly Val Pro Leu Leu  Ala Ile Gly Cys Tyr  Ser Gln Val
    2330                2335                2340

Asn Pro  Ile Thr Leu Thr Ala  Ala Leu Phe Leu Leu  Val Ala His
    2345                2350                2355

Tyr Ala  Ile Ile Gly Pro Gly  Leu Gln Ala Lys Ala  Thr Arg Glu
    2360                2365                2370

Ala Gln  Lys Arg Ala Ala Ala  Gly Ile Met Lys Asn  Pro Thr Val
    2375                2380                2385

Asp Gly  Ile Thr Val Ile Asp  Leu Asp Pro Ile Pro  Tyr Asp Pro
    2390                2395                2400
```

```
Lys Phe  Glu Lys Gln Leu Gly  Gln Val Met Leu Leu  Val Leu Cys
    2405                 2410                 2415

Val Thr  Gln Val Leu Met Met  Arg Thr Thr Trp Ala  Leu Cys Glu
    2420                 2425                 2430

Ala Leu  Thr Leu Ala Thr Gly  Pro Ile Ser Thr Leu  Trp Glu Gly
    2435                 2440                 2445

Asn Pro  Gly Arg Phe Trp Asn  Thr Thr Ile Ala Val  Ser Met Ala
    2450                 2455                 2460

Asn Ile  Phe Arg Gly Ser Tyr  Leu Ala Gly Ala Gly  Leu Leu Phe
    2465                 2470                 2475

Ser Ile  Met Lys Asn Thr Thr  Asn Thr Arg Arg Gly  Thr Gly Asn
    2480                 2485                 2490

Ile Gly  Glu Thr Leu Gly Glu  Lys Trp Lys Ser Arg  Leu Asn Ala
    2495                 2500                 2505

Leu Gly  Lys Ser Glu Phe Gln  Ile Tyr Lys Lys Ser  Gly Ile Gln
    2510                 2515                 2520

Glu Val  Asp Arg Thr Leu Ala  Lys Glu Gly Ile Lys  Arg Gly Glu
    2525                 2530                 2535

Thr Asp  His His Ala Val Ser  Arg Gly Ser Ala Lys  Leu Arg Trp
    2540                 2545                 2550

Phe Val  Glu Arg Asn Met Val  Thr Pro Glu Gly Lys  Val Val Asp
    2555                 2560                 2565

Leu Gly  Cys Gly Arg Gly Gly  Trp Ser Tyr Tyr Cys  Gly Gly Leu
    2570                 2575                 2580

Lys Asn  Val Arg Glu Val Lys  Gly Leu Thr Lys Gly  Gly Pro Gly
    2585                 2590                 2595

His Glu  Glu Pro Ile Pro Met  Ser Thr Tyr Gly Trp  Asn Leu Val
    2600                 2605                 2610

Arg Leu  Gln Ser Gly Val Asp  Val Phe Phe Ile Pro  Pro Glu Lys
    2615                 2620                 2625

Cys Asp  Thr Leu Leu Cys Asp  Ile Gly Glu Ser Ser  Pro Asn Pro
    2630                 2635                 2640

Thr Val  Glu Ala Gly Arg Thr  Leu Arg Val Leu Asn  Leu Val Glu
    2645                 2650                 2655
```

```
Asn Trp  Leu Asn Asn Asn Thr  Gln Phe Cys Ile Lys  Val Leu Asn
    2660                 2665                 2670

Pro Tyr  Met Pro Ser Val Ile  Glu Lys Met Glu Ala  Leu Gln Arg
    2675                 2680                 2685

Lys Tyr  Gly Gly Ala Leu Val  Arg Asn Pro Leu Ser  Arg Asn Ser
    2690                 2695                 2700

Thr His  Glu Met Tyr Trp Val  Ser Asn Ala Ser Gly  Asn Ile Val
    2705                 2710                 2715

Ser Ser  Val Asn Met Ile Ser  Arg Met Leu Ile Asn  Arg Phe Thr
    2720                 2725                 2730

Met Arg  Tyr Lys Lys Ala Thr  Tyr Glu Pro Asp Val  Asp Leu Gly
    2735                 2740                 2745

Ser Gly  Thr Arg Asn Ile Gly  Ile Glu Ser Glu Ile  Pro Asn Leu
    2750                 2755                 2760

Asp Ile  Ile Gly Lys Arg Ile  Glu Lys Ile Lys Gln  Glu His Glu
    2765                 2770                 2775

Thr Ser  Trp His Tyr Asp Gln  Asp His Pro Tyr Lys  Thr Trp Ala
    2780                 2785                 2790

Tyr His  Gly Ser Tyr Glu Thr  Lys Gln Thr Gly Ser  Ala Ser Ser
    2795                 2800                 2805

Met Val  Asn Gly Val Val Arg  Leu Leu Thr Lys Pro  Trp Asp Val
    2810                 2815                 2820

Val Pro  Met Val Thr Gln Met  Ala Met Thr Asp Thr  Thr Pro Phe
    2825                 2830                 2835

Gly Gln  Gln Arg Val Phe Lys  Glu Lys Val Asp Thr  Arg Thr Gln
    2840                 2845                 2850

Glu Pro  Lys Glu Gly Thr Lys  Lys Leu Met Lys Ile  Thr Ala Glu
    2855                 2860                 2865

Trp Leu  Trp Lys Glu Leu Gly  Lys Lys Lys Thr Pro  Arg Met Cys
    2870                 2875                 2880

Thr Arg  Glu Glu Phe Thr Arg  Lys Val Arg Ser Asn  Ala Ala Leu
    2885                 2890                 2895

Gly Ala  Ile Phe Thr Asp Glu  Asn Lys Trp Lys Ser  Ala Arg Glu
    2900                 2905                 2910
```

```
Ala Val  Glu Asp Ser Arg Phe  Trp Glu Leu Val Asp  Lys Glu Arg
    2915                 2920                 2925

Asn Leu  His Leu Glu Gly Lys  Cys Glu Thr Cys Val  Tyr Asn Met
    2930                 2935                 2940

Met Gly  Lys Arg Glu Lys Lys  Leu Gly Glu Phe Gly  Lys Ala Lys
    2945                 2950                 2955

Gly Ser  Arg Ala Ile Trp Tyr  Met Trp Leu Gly Ala  Arg Phe Leu
    2960                 2965                 2970

Glu Phe  Glu Ala Leu Gly Phe  Leu Asn Glu Asp His  Trp Phe Ser
    2975                 2980                 2985

Arg Glu  Asn Ser Leu Ser Gly  Val Glu Gly Glu Gly  Leu His Lys
    2990                 2995                 3000

Leu Gly  Tyr Ile Leu Arg Asp  Val Ser Lys Lys Glu  Gly Gly Ala
    3005                 3010                 3015

Met Tyr  Ala Asp Asp Thr Ala  Gly Trp Asp Thr Arg  Ile Thr Leu
    3020                 3025                 3030

Glu Asp  Leu Lys Asn Glu Glu  Met Val Thr Asn His  Met Glu Gly
    3035                 3040                 3045

Glu His  Lys Lys Leu Ala Glu  Ala Ile Phe Lys Leu  Thr Tyr Gln
    3050                 3055                 3060

Asn Lys  Val Val Arg Val Gln  Arg Pro Thr Pro Arg  Gly Thr Val
    3065                 3070                 3075

Met Asp  Ile Ile Ser Arg Arg  Asp Gln Arg Gly Ser  Gly Gln Val
    3080                 3085                 3090

Gly Thr  Tyr Gly Leu Asn Thr  Phe Thr Asn Met Glu  Ala Gln Leu
    3095                 3100                 3105

Ile Arg  Gln Met Glu Gly Glu  Gly Val Phe Lys Ser  Ile Gln His
    3110                 3115                 3120

Leu Thr  Ile Thr Glu Glu Ile  Ala Val Gln Asn Trp  Leu Ala Arg
    3125                 3130                 3135

Val Gly  Arg Glu Arg Leu Ser  Arg Met Ala Ile Ser  Gly Asp Asp
    3140                 3145                 3150

Cys Val  Val Lys Pro Leu Asp  Asp Arg Phe Ala Ser  Ala Leu Thr
    3155                 3160                 3165
```

```
Ala Leu  Asn Asp Met Gly Lys  Ile Arg Lys Asp Ile  Gln Gln Trp
    3170                 3175                 3180

Glu Pro  Ser Arg Gly Trp Asn  Asp Trp Thr Gln Val  Pro Phe Cys
    3185                 3190                 3195

Ser His  His Phe His Glu Leu  Ile Met Lys Asp Gly  Arg Val Leu
    3200                 3205                 3210

Val Val  Pro Cys Arg Asn Gln  Asp Glu Leu Ile Gly  Arg Ala Arg
    3215                 3220                 3225

Ile Ser  Gln Gly Ala Gly Trp  Ser Leu Arg Glu Thr  Ala Cys Leu
    3230                 3235                 3240

Gly Lys  Ser Tyr Ala Gln Met  Trp Ser Leu Met Tyr  Phe His Arg
    3245                 3250                 3255

Arg Asp  Leu Arg Leu Ala Ala  Asn Ala Ile Cys Ser  Ala Val Pro
    3260                 3265                 3270

Ser His  Trp Val Pro Thr Ser  Arg Thr Thr Trp Ser  Ile His Ala
    3275                 3280                 3285

Lys His  Glu Trp Met Thr Thr  Glu Asp Met Leu Thr  Val Trp Asn
    3290                 3295                 3300

Arg Val  Trp Ile Gln Glu Asn  Pro Trp Met Glu Asp  Lys Thr Pro
    3305                 3310                 3315

Val Glu  Ser Trp Glu Glu Ile  Pro Tyr Leu Gly Lys  Arg Glu Asp
    3320                 3325                 3330

Gln Trp  Cys Gly Ser Leu Ile  Gly Leu Thr Ser Arg  Ala Thr Trp
    3335                 3340                 3345

Ala Lys  Asn Ile Gln Ala Ala  Ile Asn Gln Val Arg  Ser Leu Ile
    3350                 3355                 3360

Gly Asn  Glu Glu Tyr Thr Asp  Tyr Met Pro Ser Met  Lys Arg Phe
    3365                 3370                 3375

Arg Arg  Glu Glu Glu Glu Ala  Gly Val Leu Trp
    3380                 3385
```

<210> 9
<211> 3389
<212> PRT
<213> Artificial Sequence

<220>
<223> Dengue virus serotype 3, MVS

<400> 9

```
Met Asn Asn Gln Arg Lys Lys Ala Lys Asn Thr Pro Phe Asn Met Leu
1               5                  10                  15

Lys Arg Glu Arg Asn Arg Val Ser Thr Val Gln Gln Leu Thr Lys Arg
            20                  25                  30

Phe Ser Leu Gly Met Leu Gln Gly Arg Gly Pro Leu Lys Leu Phe Met
        35                  40                  45

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

Ile Leu Lys Arg Trp Gly Thr Ile Lys Lys Ser Lys Ala Ile Asn Val
65                  70                  75                  80

Leu Arg Gly Phe Arg Lys Glu Ile Gly Arg Met Leu Asn Ile Leu Asn
                85                  90                  95

Arg Arg Arg Arg Ser Ala Gly Met Ile Ile Met Leu Ile Pro Thr Val
            100                 105                 110

Met Ala Phe His Leu Thr Thr Arg Asp Gly Glu Pro Arg Met Ile Val
        115                 120                 125

Gly Lys Asn Glu Arg Gly Lys Ser Leu Leu Phe Lys Thr Ala Ser Gly
    130                 135                 140

Ile Asn Met Cys Thr Leu Ile Ala Met Asp Leu Gly Glu Met Cys Asp
145                 150                 155                 160

Asp Thr Val Thr Tyr Lys Cys Pro His Ile Thr Glu Val Glu Pro Glu
                165                 170                 175

Asp Ile Asp Cys Trp Cys Asn Leu Thr Ser Thr Trp Val Thr Tyr Gly
            180                 185                 190

Thr Cys Asn Gln Ala Gly Glu His Arg Arg Asp Lys Arg Ser Val Ala
        195                 200                 205

Leu Ala Pro His Val Gly Met Gly Leu Asp Thr Arg Thr Gln Thr Trp
    210                 215                 220

Met Ser Ala Glu Gly Ala Trp Arg Gln Val Glu Lys Val Glu Thr Trp
225                 230                 235                 240

Ala Leu Arg His Pro Gly Phe Thr Ile Leu Ala Leu Phe Leu Ala His
                245                 250                 255

Tyr Ile Gly Thr Ser Leu Thr Gln Lys Val Val Ile Phe Ile Leu Leu
```

260 265 270

Met Leu Val Thr Pro Ser Met Thr Met Arg Cys Val Gly Val Gly Asn
275 280 285

Arg Asp Phe Val Glu Gly Leu Ser Gly Ala Thr Trp Val Asp Val Val
290 295 300

Leu Glu His Gly Gly Cys Val Thr Thr Met Ala Lys Asn Lys Pro Thr
305 310 315 320

Leu Asp Ile Glu Leu Gln Lys Thr Glu Ala Thr Gln Leu Ala Thr Leu
325 330 335

Arg Lys Leu Cys Ile Glu Gly Lys Ile Thr Asn Ile Thr Thr Asp Ser
340 345 350

Arg Cys Pro Thr Gln Gly Glu Ala Ile Leu Pro Glu Glu Gln Asp Gln
355 360 365

Asn Tyr Val Cys Lys His Thr Tyr Val Asp Arg Gly Trp Gly Asn Gly
370 375 380

Cys Gly Leu Phe Gly Lys Gly Ser Leu Val Thr Cys Ala Lys Phe Gln
385 390 395 400

Cys Leu Glu Ser Ile Glu Gly Lys Val Val Gln His Glu Asn Leu Lys
405 410 415

Tyr Thr Val Ile Ile Thr Val His Thr Gly Asp Gln His Gln Val Gly
420 425 430

Asn Glu Thr Gln Gly Val Thr Ala Glu Ile Thr Pro Gln Ala Ser Thr
435 440 445

Ala Glu Ala Ile Leu Pro Glu Tyr Gly Thr Leu Gly Leu Glu Cys Ser
450 455 460

Pro Arg Thr Gly Leu Asp Phe Asn Glu Met Ile Ser Leu Thr Met Lys
465 470 475 480

Asn Lys Ala Trp Met Val His Arg Gln Trp Phe Phe Asp Leu Pro Leu
485 490 495

Pro Trp Thr Ser Gly Ala Ser Ala Glu Thr Pro Thr Trp Asn Arg Lys
500 505 510

Glu Leu Leu Val Thr Phe Lys Asn Ala His Ala Lys Lys Gln Glu Val
515 520 525

Val Val Leu Gly Ser Gln Glu Gly Ala Met His Thr Ala Leu Thr Gly

530 535 540

Ala Thr Glu Ile Gln Thr Ser Gly Gly Thr Ser Ile Phe Ala Gly His
545 550 555 560

Leu Lys Cys Arg Leu Lys Met Asp Lys Leu Glu Leu Lys Gly Met Ser
565 570 575

Tyr Ala Met Cys Leu Ser Ser Phe Val Leu Lys Lys Glu Val Ser Glu
580 585 590

Thr Gln His Gly Thr Ile Leu Ile Lys Val Glu Tyr Lys Gly Glu Asp
595 600 605

Ala Pro Cys Lys Ile Pro Phe Ser Thr Glu Asp Gly Gln Gly Lys Ala
610 615 620

Leu Asn Gly Arg Leu Ile Thr Ala Asn Pro Val Val Thr Lys Lys Glu
625 630 635 640

Glu Pro Val Asn Ile Glu Ala Glu Pro Pro Phe Gly Glu Ser Asn Ile
645 650 655

Val Ile Gly Ile Gly Asp Lys Ala Leu Lys Ile Asn Trp Tyr Lys Lys
660 665 670

Gly Ser Ser Ile Gly Lys Met Phe Glu Ala Thr Ala Arg Gly Ala Arg
675 680 685

Arg Met Ala Ile Leu Gly Asp Thr Ala Trp Asp Phe Gly Ser Val Gly
690 695 700

Gly Val Leu Asn Ser Leu Gly Lys Met Val His Gln Ile Phe Gly Ser
705 710 715 720

Ala Tyr Thr Ala Leu Phe Gly Gly Val Ser Trp Met Met Lys Ile Gly
725 730 735

Ile Gly Val Leu Leu Thr Trp Ile Gly Leu Asn Ser Lys Asn Thr Ser
740 745 750

Met Ser Phe Ser Cys Ile Ala Ala Gly Ile Val Thr Leu Tyr Leu Gly
755 760 765

Val Met Val Gln Ala Asp Ser Gly Cys Val Val Ser Trp Lys Asn Lys
770 775 780

Glu Leu Lys Cys Gly Ser Gly Ile Phe Ile Thr Asp Asn Val His Thr
785 790 795 800

Trp Thr Glu Gln Tyr Lys Phe Gln Pro Glu Ser Pro Ser Lys Leu Ala

805 810 815

Ser Ala Ile Gln Lys Ala His Glu Glu Asp Ile Cys Gly Ile Arg Ser
820 825 830

Val Thr Arg Leu Glu Asn Leu Met Trp Lys Gln Ile Thr Pro Glu Leu
835 840 845

Asn His Ile Leu Ser Glu Asn Glu Val Lys Leu Thr Ile Met Thr Gly
850 855 860

Asp Ile Lys Gly Ile Met Gln Ala Gly Lys Arg Ser Leu Arg Pro Gln
865 870 875 880

Pro Thr Glu Leu Lys Tyr Ser Trp Lys Thr Trp Gly Lys Ala Lys Met
885 890 895

Leu Ser Thr Glu Ser His Asn Gln Thr Phe Leu Ile Asp Gly Pro Glu
900 905 910

Thr Ala Glu Cys Pro Asn Thr Asn Arg Ala Trp Asn Ser Leu Glu Val
915 920 925

Glu Asp Tyr Gly Phe Gly Val Phe Thr Thr Asn Ile Trp Leu Lys Leu
930 935 940

Lys Glu Lys Gln Asp Val Phe Cys Asp Ser Lys Leu Met Ser Ala Ala
945 950 955 960

Ile Lys Asp Asn Arg Ala Val His Ala Asp Met Gly Tyr Trp Ile Glu
965 970 975

Ser Ala Leu Asn Asp Thr Trp Lys Ile Glu Lys Ala Ser Phe Ile Glu
980 985 990

Val Lys Asn Cys His Trp Pro Lys Ser His Thr Leu Trp Ser Asn Gly
995 1000 1005

Val Leu Glu Ser Glu Met Ile Ile Pro Lys Asn Leu Ala Gly Pro
1010 1015 1020

Val Ser Gln His Asn Tyr Arg Pro Gly Tyr His Thr Gln Ile Thr
1025 1030 1035

Gly Pro Trp His Leu Gly Lys Leu Glu Met Asp Phe Asp Phe Cys
1040 1045 1050

Asp Gly Thr Thr Val Val Val Thr Glu Asp Cys Gly Asn Arg Gly
1055 1060 1065

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

1070 1075 1080

Trp Cys Cys Arg Ser Cys Thr Leu Pro Pro Leu Arg Tyr Arg Gly
1085 1090 1095

Glu Asp Gly Cys Trp Tyr Gly Met Glu Ile Arg Pro Leu Lys Glu
1100 1105 1110

Lys Glu Glu Asn Leu Val Asn Ser Leu Val Thr Ala Gly His Gly
1115 1120 1125

Gln Val Asp Asn Phe Ser Leu Gly Val Leu Gly Met Ala Leu Phe
1130 1135 1140

Leu Glu Glu Met Leu Arg Thr Arg Val Gly Thr Lys His Ala Ile
1145 1150 1155

Leu Leu Val Ala Val Ser Phe Val Thr Leu Ile Thr Gly Asn Met
1160 1165 1170

Ser Phe Arg Asp Leu Gly Arg Val Met Val Met Val Gly Ala Thr
1175 1180 1185

Met Thr Asp Asp Ile Gly Met Gly Val Thr Tyr Leu Ala Leu Leu
1190 1195 1200

Ala Ala Phe Lys Val Arg Pro Thr Phe Ala Ala Gly Leu Leu Leu
1205 1210 1215

Arg Lys Leu Thr Ser Lys Glu Leu Met Met Thr Thr Ile Gly Ile
1220 1225 1230

Val Leu Leu Ser Gln Ser Thr Ile Pro Glu Thr Ile Leu Glu Leu
1235 1240 1245

Thr Asp Ala Leu Ala Leu Gly Met Met Val Leu Lys Met Val Arg
1250 1255 1260

Asn Met Glu Lys Tyr Gln Leu Ala Val Thr Ile Met Ala Ile Leu
1265 1270 1275

Cys Val Pro Asn Ala Val Ile Leu Gln Asn Ala Trp Lys Val Ser
1280 1285 1290

Cys Thr Ile Leu Ala Val Val Ser Val Ser Pro Leu Phe Leu Thr
1295 1300 1305

Ser Ser Gln Gln Lys Thr Asp Trp Ile Pro Leu Ala Leu Thr Ile
1310 1315 1320

Lys Gly Leu Asn Pro Thr Ala Ile Phe Leu Thr Thr Leu Ser Arg

1325 1330 1335

Thr Ser Lys Lys Arg Ser Trp Pro Leu Asn Glu Ala Ile Met Ala
1340 1345 1350

Val Gly Met Val Ser Ile Leu Ala Ser Ser Leu Leu Lys Asn Asp
1355 1360 1365

Ile Pro Met Thr Gly Pro Leu Val Ala Gly Gly Leu Leu Thr Val
1370 1375 1380

Cys Tyr Val Leu Thr Gly Arg Ser Ala Asp Leu Glu Leu Glu Arg
1385 1390 1395

Ala Ala Asp Val Lys Trp Glu Asp Gln Ala Glu Ile Ser Gly Ser
1400 1405 1410

Ser Pro Ile Leu Ser Ile Thr Ile Ser Glu Asp Gly Ser Met Ser
1415 1420 1425

Ile Lys Asn Glu Glu Glu Glu Gln Thr Leu Thr Ile Leu Ile Arg
1430 1435 1440

Thr Gly Leu Leu Val Ile Ser Gly Leu Phe Pro Val Ser Ile Pro
1445 1450 1455

Ile Thr Ala Ala Ala Trp Tyr Leu Trp Glu Val Lys Lys Gln Arg
1460 1465 1470

Ala Gly Val Leu Trp Asp Val Pro Ser Pro Pro Pro Met Gly Lys
1475 1480 1485

Ala Glu Leu Glu Asp Gly Ala Tyr Arg Ile Lys Gln Lys Gly Ile
1490 1495 1500

Leu Gly Tyr Ser Gln Ile Gly Ala Gly Val Tyr Lys Glu Gly Thr
1505 1510 1515

Phe His Thr Met Trp His Val Thr Arg Gly Ala Val Leu Met His
1520 1525 1530

Lys Gly Lys Arg Ile Glu Pro Ser Trp Ala Asp Val Lys Lys Asp
1535 1540 1545

Leu Ile Ser Tyr Gly Gly Gly Trp Lys Leu Glu Gly Glu Trp Lys
1550 1555 1560

Glu Gly Glu Glu Val Gln Val Leu Ala Leu Glu Pro Gly Lys Asn
1565 1570 1575

Pro Arg Ala Val Gln Thr Lys Pro Gly Leu Phe Lys Thr Asn Ala

1580 1585 1590

Gly Thr Ile Gly Ala Val Ser Leu Asp Phe Ser Pro Gly Thr Ser
1595 1600 1605

Gly Ser Pro Ile Ile Asp Lys Lys Gly Lys Val Val Gly Leu Tyr
1610 1615 1620

Gly Asn Gly Val Val Thr Arg Ser Gly Ala Tyr Val Ser Ala Ile
1625 1630 1635

Ala Gln Thr Glu Lys Ser Ile Glu Asp Asn Pro Glu Ile Glu Asp
1640 1645 1650

Asp Ile Phe Arg Lys Arg Arg Leu Thr Ile Met Asp Leu His Pro
1655 1660 1665

Gly Ala Gly Lys Thr Lys Arg Tyr Leu Pro Ala Ile Val Arg Glu
1670 1675 1680

Ala Ile Lys Arg Gly Leu Arg Thr Leu Ile Leu Ala Pro Thr Arg
1685 1690 1695

Val Val Ala Ala Glu Met Glu Glu Ala Leu Arg Gly Leu Pro Ile
1700 1705 1710

Arg Tyr Gln Thr Pro Ala Ile Arg Ala Val His Thr Gly Arg Glu
1715 1720 1725

Ile Val Asp Leu Met Cys His Ala Thr Phe Thr Met Arg Leu Leu
1730 1735 1740

Ser Pro Val Arg Val Pro Asn Tyr Asn Leu Ile Ile Met Asp Glu
1745 1750 1755

Ala His Phe Thr Asp Pro Ala Ser Ile Ala Ala Arg Gly Tyr Ile
1760 1765 1770

Ser Thr Arg Val Glu Met Gly Glu Ala Ala Gly Ile Phe Met Thr
1775 1780 1785

Ala Thr Pro Pro Gly Ser Arg Asp Pro Phe Pro Gln Ser Asn Ala
1790 1795 1800

Pro Ile Ile Asp Glu Glu Arg Glu Ile Pro Glu Arg Ser Trp Asn
1805 1810 1815

Ser Gly His Glu Trp Val Thr Asp Phe Lys Gly Lys Thr Val Trp
1820 1825 1830

Phe Val Pro Ser Ile Lys Ala Gly Asn Asp Ile Ala Ala Cys Leu

1835 1840 1845

Arg Lys Asn Gly Lys Lys Val Ile Gln Leu Ser Arg Lys Thr Phe
1850 1855 1860

Asp Ser Glu Tyr Val Lys Thr Arg Thr Asn Asp Trp Asp Phe Val
1865 1870 1875

Val Thr Thr Asp Ile Ser Glu Met Gly Ala Asn Phe Lys Ala Glu
1880 1885 1890

Arg Val Ile Asp Pro Arg Arg Cys Met Lys Pro Val Ile Leu Thr
1895 1900 1905

Asp Gly Glu Glu Arg Val Ile Leu Ala Gly Pro Met Pro Val Thr
1910 1915 1920

His Ser Ser Ala Ala Gln Arg Arg Gly Arg Ile Gly Arg Asn Pro
1925 1930 1935

Lys Asn Glu Asn Asp Gln Tyr Ile Tyr Met Gly Glu Pro Leu Glu
1940 1945 1950

Asn Asp Glu Asp Cys Ala His Trp Lys Glu Ala Lys Met Leu Leu
1955 1960 1965

Asp Asn Ile Asn Thr Pro Glu Gly Ile Ile Pro Ser Met Phe Glu
1970 1975 1980

Pro Glu Arg Glu Lys Val Asp Ala Ile Asp Gly Glu Tyr Arg Leu
1985 1990 1995

Arg Gly Glu Ala Arg Lys Thr Phe Val Asp Leu Met Arg Arg Gly
2000 2005 2010

Asp Leu Pro Val Trp Leu Ala Tyr Arg Val Ala Ala Glu Gly Ile
2015 2020 2025

Asn Tyr Ala Asp Arg Arg Trp Cys Phe Asp Gly Val Lys Asn Asn
2030 2035 2040

Gln Ile Leu Glu Glu Asn Val Glu Val Glu Ile Trp Thr Lys Glu
2045 2050 2055

Gly Glu Arg Lys Lys Leu Lys Pro Arg Trp Leu Asp Ala Arg Ile
2060 2065 2070

Tyr Ser Asp Pro Leu Ala Leu Lys Glu Phe Lys Glu Phe Ala Ala
2075 2080 2085

Gly Arg Lys Ser Leu Thr Leu Asn Leu Ile Thr Glu Met Gly Arg

2090 2095 2100

Leu Pro Thr Phe Met Thr Gln Lys Ala Arg Asp Ala Leu Asp Asn
2105 2110 2115

Leu Ala Val Leu His Thr Ala Glu Ala Gly Gly Arg Ala Tyr Asn
2120 2125 2130

His Ala Leu Ser Glu Leu Pro Glu Thr Leu Glu Thr Leu Leu Leu
2135 2140 2145

Leu Thr Leu Leu Ala Thr Val Thr Gly Gly Ile Phe Leu Phe Leu
2150 2155 2160

Met Ser Ala Arg Gly Ile Gly Lys Met Thr Leu Gly Met Cys Cys
2165 2170 2175

Ile Ile Thr Ala Ser Ile Leu Leu Trp Tyr Ala Gln Ile Gln Pro
2180 2185 2190

His Trp Ile Ala Ala Ser Ile Ile Leu Glu Phe Phe Leu Ile Val
2195 2200 2205

Leu Leu Ile Pro Glu Pro Glu Lys Gln Arg Thr Pro Gln Asp Asn
2210 2215 2220

Gln Leu Thr Tyr Val Val Ile Ala Ile Leu Thr Val Val Ala Ala
2225 2230 2235

Thr Met Ala Asn Glu Met Gly Phe Leu Glu Lys Thr Lys Lys Asp
2240 2245 2250

Leu Gly Leu Gly Ser Ile Ala Thr Gln Gln Pro Glu Ser Asn Ile
2255 2260 2265

Leu Asp Ile Asp Leu Arg Pro Ala Ser Ala Trp Thr Leu Tyr Ala
2270 2275 2280

Val Ala Thr Thr Phe Val Thr Pro Met Leu Arg His Ser Ile Glu
2285 2290 2295

Asn Ser Ser Val Asn Val Ser Leu Thr Ala Ile Ala Asn Gln Ala
2300 2305 2310

Thr Val Leu Met Gly Leu Gly Lys Gly Trp Pro Leu Ser Lys Met
2315 2320 2325

Asp Ile Gly Val Pro Leu Leu Ala Ile Gly Cys Tyr Ser Gln Val
2330 2335 2340

Asn Pro Ile Thr Leu Thr Ala Ala Leu Phe Leu Leu Val Ala His

2345 2350 2355

Tyr Ala Ile Ile Gly Pro Gly Leu Gln Ala Lys Ala Thr Arg Glu
2360 2365 2370

Ala Gln Lys Arg Ala Ala Ala Gly Ile Met Lys Asn Pro Thr Val
2375 2380 2385

Asp Gly Ile Thr Val Ile Asp Leu Asp Pro Ile Pro Tyr Asp Pro
2390 2395 2400

Lys Phe Glu Lys Gln Leu Gly Gln Val Met Leu Leu Val Leu Cys
2405 2410 2415

Val Thr Gln Val Leu Met Met Arg Thr Thr Trp Ala Leu Cys Glu
2420 2425 2430

Ala Leu Thr Leu Ala Thr Gly Pro Ile Ser Thr Leu Trp Glu Gly
2435 2440 2445

Asn Pro Gly Arg Phe Trp Asn Thr Thr Ile Ala Val Ser Met Ala
2450 2455 2460

Asn Ile Phe Arg Gly Ser Tyr Leu Ala Gly Ala Gly Leu Leu Phe
2465 2470 2475

Ser Ile Met Lys Asn Thr Thr Asn Thr Arg Arg Gly Thr Gly Asn
2480 2485 2490

Ile Gly Glu Thr Leu Gly Glu Lys Trp Lys Ser Arg Leu Asn Ala
2495 2500 2505

Leu Gly Lys Ser Glu Phe Gln Ile Tyr Lys Lys Ser Gly Ile Gln
2510 2515 2520

Glu Val Asp Arg Thr Leu Ala Lys Glu Gly Ile Lys Arg Gly Glu
2525 2530 2535

Thr Asp His His Ala Val Ser Arg Gly Ser Ala Lys Leu Arg Trp
2540 2545 2550

Phe Val Glu Arg Asn Met Val Thr Pro Glu Gly Lys Val Val Asp
2555 2560 2565

Leu Gly Cys Gly Arg Gly Gly Trp Ser Tyr Tyr Cys Gly Gly Leu
2570 2575 2580

Lys Asn Val Arg Glu Val Lys Gly Leu Thr Lys Gly Gly Pro Gly
2585 2590 2595

His Glu Glu Pro Ile Pro Met Ser Thr Tyr Gly Trp Asn Leu Val

2600 2605 2610

Arg Leu Gln Ser Gly Val Asp Val Phe Phe Ile Pro Pro Glu Lys
2615 2620 2625

Cys Asp Thr Leu Leu Cys Asp Ile Gly Glu Ser Ser Pro Asn Pro
2630 2635 2640

Thr Val Glu Ala Gly Arg Thr Leu Arg Val Leu Asn Leu Val Glu
2645 2650 2655

Asn Trp Leu Asn Asn Asn Thr Gln Phe Cys Ile Lys Val Leu Asn
2660 2665 2670

Pro Tyr Met Pro Ser Val Ile Glu Lys Met Glu Ala Leu Gln Arg
2675 2680 2685

Lys Tyr Gly Gly Ala Leu Val Arg Asn Pro Leu Ser Arg Asn Ser
2690 2695 2700

Thr His Glu Met Tyr Trp Val Ser Asn Ala Ser Gly Asn Ile Val
2705 2710 2715

Ser Ser Val Asn Met Ile Ser Arg Met Leu Ile Asn Arg Phe Thr
2720 2725 2730

Met Arg Tyr Lys Lys Ala Thr Tyr Glu Pro Asp Val Asp Leu Gly
2735 2740 2745

Ser Gly Thr Arg Asn Ile Gly Ile Glu Ser Glu Ile Pro Asn Leu
2750 2755 2760

Asp Ile Ile Gly Lys Arg Ile Glu Lys Ile Lys Gln Glu His Glu
2765 2770 2775

Thr Ser Trp His Tyr Asp Gln Asp His Pro Tyr Lys Thr Trp Ala
2780 2785 2790

Tyr His Gly Ser Tyr Glu Thr Lys Gln Thr Gly Ser Ala Ser Ser
2795 2800 2805

Met Val Asn Gly Val Val Arg Leu Leu Thr Lys Pro Trp Asp Val
2810 2815 2820

Val Pro Met Val Thr Gln Met Ala Met Thr Asp Thr Thr Pro Phe
2825 2830 2835

Gly Gln Gln Arg Val Phe Lys Glu Lys Val Asp Thr Arg Thr Gln
2840 2845 2850

Glu Pro Lys Glu Gly Thr Lys Lys Leu Met Lys Ile Thr Ala Glu

2855 2860 2865

Trp Leu Trp Lys Glu Leu Gly Lys Lys Lys Thr Pro Arg Met Cys
2870 2875 2880

Thr Arg Glu Glu Phe Thr Arg Lys Val Arg Ser Asn Ala Ala Leu
2885 2890 2895

Gly Ala Ile Phe Thr Asp Glu Asn Lys Trp Lys Ser Ala Arg Glu
2900 2905 2910

Ala Val Glu Asp Ser Arg Phe Trp Glu Leu Val Asp Lys Glu Arg
2915 2920 2925

Asn Leu His Leu Glu Gly Lys Cys Glu Thr Cys Val Tyr Asn Met
2930 2935 2940

Met Gly Lys Arg Glu Lys Lys Leu Gly Glu Phe Gly Lys Ala Lys
2945 2950 2955

Gly Ser Arg Ala Ile Trp Tyr Met Trp Leu Gly Ala Arg Phe Leu
2960 2965 2970

Glu Phe Glu Ala Leu Gly Phe Leu Asn Glu Asp His Trp Phe Ser
2975 2980 2985

Arg Glu Asn Ser Leu Ser Gly Val Glu Gly Glu Gly Leu His Lys
2990 2995 3000

Leu Gly Tyr Ile Leu Arg Asp Val Ser Lys Lys Glu Gly Gly Ala
3005 3010 3015

Met Tyr Ala Asp Asp Thr Ala Gly Trp Asp Thr Arg Ile Thr Leu
3020 3025 3030

Glu Asp Leu Lys Asn Glu Glu Met Val Thr Asn His Met Glu Gly
3035 3040 3045

Glu His Lys Lys Leu Ala Glu Ala Ile Phe Lys Leu Thr Tyr Gln
3050 3055 3060

Asn Lys Val Val Arg Val Gln Arg Pro Thr Pro Arg Gly Thr Val
3065 3070 3075

Met Asp Ile Ile Ser Arg Arg Asp Gln Arg Gly Ser Gly Gln Val
3080 3085 3090

Gly Thr Tyr Gly Leu Asn Thr Phe Thr Asn Met Glu Ala Gln Leu
3095 3100 3105

Ile Arg Gln Met Glu Gly Glu Gly Val Phe Lys Ser Ile Gln His

3110 3115 3120

Leu Thr Ile Thr Glu Glu Ile Ala Val Gln Asn Trp Leu Ala Arg
3125 3130 3135

Val Gly Arg Glu Arg Leu Ser Arg Met Ala Ile Ser Gly Asp Asp
3140 3145 3150

Cys Val Val Lys Pro Leu Asp Asp Arg Phe Ala Ser Ala Leu Thr
3155 3160 3165

Ala Leu Asn Asp Met Gly Lys Ile Arg Lys Asp Ile Gln Gln Trp
3170 3175 3180

Glu Pro Ser Arg Gly Trp Asn Asp Trp Thr Gln Val Pro Phe Cys
3185 3190 3195

Ser His His Phe His Glu Leu Ile Met Lys Asp Gly Arg Val Leu
3200 3205 3210

Val Val Pro Cys Arg Asn Gln Asp Glu Leu Ile Gly Arg Ala Arg
3215 3220 3225

Ile Ser Gln Gly Ala Gly Trp Ser Leu Arg Glu Thr Ala Cys Leu
3230 3235 3240

Gly Lys Ser Tyr Ala Gln Met Trp Ser Leu Met Tyr Phe His Arg
3245 3250 3255

Arg Asp Leu Arg Leu Ala Ala Asn Ala Ile Cys Ser Ala Val Pro
3260 3265 3270

Ser His Trp Val Pro Thr Ser Arg Thr Thr Trp Ser Ile His Ala
3275 3280 3285

Lys His Glu Trp Met Thr Thr Glu Asp Met Leu Thr Val Trp Asn
3290 3295 3300

Arg Val Trp Ile Gln Glu Asn Pro Trp Met Glu Asp Lys Thr Pro
3305 3310 3315

Val Glu Ser Trp Glu Glu Ile Pro Tyr Leu Gly Lys Arg Glu Asp
3320 3325 3330

Gln Trp Cys Gly Ser Leu Ile Gly Leu Thr Ser Arg Ala Thr Trp
3335 3340 3345

Ala Lys Asn Ile Gln Ala Ala Ile Asn Gln Val Arg Ser Leu Ile
3350 3355 3360

Gly Asn Glu Glu Tyr Thr Asp Tyr Met Pro Ser Met Lys Arg Phe

3365 3370 3375

Arg Arg Glu Glu Glu Glu Ala Gly Val Leu Trp
3380 3385

<210> 10
<211> 10723
<212> DNA
<213> Artificial Sequence

<220>
<223> Dengue virus serotype 4, BVS

<400> 10
agttgttagt ctacgtggac cgacaaagac agattctttg agggagctaa gctcaatgta 60

gttctaacag ttttttaatt agagagcaga tctctgatga ataaccaacg gaaaaaggcg 120

aaaaacacgc ctttcaatat gctgaaacgc gagagaaacc gcgtgtcgac tgtgcaacag 180

ctgacaaaga gattctcact tggaatgctg cagggacgag gacctttaaa actgttcatg 240

gccctggtgg cgttccttcg tttcctaaca atcccaccaa cagcagggat attgaagaga 300

tggggaacaa ttaaaaaatc aaaagctatt aatgttttga gagggttcag gaaagagatt 360

ggaaggatgc tgaacatctt gaataggaga cgcagctctg caggcatgat cattatgctg 420

attccaacag tgatggcgtt ccatttaacc acgcgtgatg gcgaacccct catgatagtg 480

gcaaaacatg aaagggggag acctctcttg tttaagacaa cagaggggat caacaaatgc 540

actctcattg ccatggactt gggtgaaatg tgtgaggaca ctgtcacgta taaatgcccc 600

ttactggtca ataccgaacc tgaagacatt gattgctggt gcaatctcac gtctacctgg 660

gtcatgtatg ggacatgcac ccagagcgga gaacggagac gagagaagcg ctcagtagct 720

ttaacaccac attcaggaat gggattggaa acaagagctg agacatggat gtcatcggaa 780

ggggcttgga agcatgctca gagagtagag agctggatac tcagaaaccc aggattcgcg 840

ctcttggcag gatttatggc ttatatgatt gggcaaacag gaatccagcg aactgtcttc 900

tttgtcctaa tgatgctggt cgccccatcc tacggaatgc gatgcgtagg agtaggaaac 960

agagactttg tggaaggagt ctcaggtgga gcatgggtcg atctggtgct agaacatgga 1020

ggatgcgtca caaccatggc ccagggaaaa ccaaccttgg attttgaact gactaagaca 1080

acagccaagg aagtggctct gttaagaacc tattgcattg aagcctcaat atcaaacata 1140

accacggcaa caagatgtcc aacgcaagga gagccttatc taaaagagga acaagaccaa 1200

cagtacattt gccggagaga tgtggtagac agagggtggg gcaatggctg tggcttgttt 1260

ggaaaaggag gagttgtgac atgtgcgaag ttttcatgtt cggggaagat aacaggcaat 1320

ttggtccaaa ttgagaacct tgaatacaca gtggttgtaa cagtccacaa tggagacacc 1380

catgcagtag gaaatgacac gtccaatcat ggagttacag ccacgataac tcccaggtca 1440

ccatcggtgg aagtcaaatt gccggactat ggagaactaa cactcgattg tgaacccagg 1500

tctggaattg actttaatga gatgattctg atgaaaatga aaaagaaaac atggcttgtg 1560

cataagcaat ggtttttgga tctacctcta ccatggacag caggagcaga cacatcagag 1620

gttcactgga attacaaaga gagaatggtg acatttaagg ttcctcatgc caagagacag 1680

gatgtgacag tgctgggatc tcaggaagga gccatgcatt ctgccctcgc tggagccaca 1740

gaagtggact ccggtgatgg aaatcacatg tttgcaggac atctcaagtg caaagtccgt 1800

atggagaaat tgagaatcaa gggaatgtca tacacgatgt gttcaggaaa gttctcaatt 1860

gacaaagaga tggcagaaac acagcatggg acaacagtgg tgaaagtcaa gtatgaaggt 1920

gctggagctc cgtgtaaagt ccccatagag ataagagatg tgaacaagga aaaagtggtt 1980

gggcgtatca tctcatccac ccctttggct gagaatacca acagtgtaac caacatagag 2040

ttagaacccc cctttgggga cagctacata gtgataggtg ttggaaacag tgcattaaca 2100

ctccattggt tcaggaaagg gagttccatt ggcaagatgt ttgagtccac atacagaggt 2160

gcaaaacgaa tggccattct aggtgaaaca gcttgggatt ttggttccgt tggtggactg 2220

ttcacatcat tgggaaaggc tgtgcaccag gtttttggaa gtgtgtatac aaccctgttt 2280

ggaggagtct catggatgat tagaatccta attgggttcc tagtgttgtg gattggcacg 2340

aactcaagga acacttcaat ggctatgacg tgcatagctg ccggcattgt gacactgtat 2400

ttgggagtca tggtgcaggc cgatagtggt tgcgttgtga gctggaaaaa caaagaactg 2460

aaatgtggca gtgggatttt catcacagac aacgtgcaca catggacaga acaatacaag 2520

ttccaaccag aatccccttc aaaactagct tcagctatcc agaaagccca tgaagaggac 2580

atttgtggaa tccgctcagt aacaagactg gagaatctga tgtggaaaca aataacacca 2640

gaattgaatc acattctatc agaaaatgag gtgaagttaa ctattatgac aggagacatc 2700

aaaggaatca tgcaggcagg aaaacgatct ctgcggcctc agcccactga gctgaagtat 2760

tcatggaaaa catggggcaa agcaaaaatg ctctctacag agtctcataa ccagaccttt 2820

ctcattgatg gccccgaaac agcagaatgc cccaacacaa atagagcttg gaattcgttg 2880

gaagttgaag actatggctt tggagtattc accaccaata tatggctaaa attgaaagaa 2940

aaacaggatg tattctgcga ctcaaaactc atgtcagcgg ccataaaaga caacagagcc 3000

gtccatgccg atatgggtta ttggatagaa agtgcactca atgacacatg gaagatagag 3060

aaagcctctt tcattgaagt taaaaactgc cactggccaa aatcacacac cctctggagc 3120

aatggagtgc tagaaagtga gatgataatt ccaaagaatc tcgctggacc agtgtctcaa 3180

cacaactata gaccaggcta ccatacacaa ataacaggac catggcatct aggtaagctt 3240

gagatggact ttgatttctg tgatggaaca acagtggtag tgactgagga ctgcggaaat 3300

agaggaccct ctttgagaac aaccactgcc tctggaaaac tcataacaga atggtgctgc 3360

cgatcttgca cattaccacc gctaagatac agaggtgagg atgggtgctg gtacgggatg 3420

gaaatcagac cattgaagga gaaagaagag aatttggtca actccttggt cacagctgga 3480

catgggcagg tcgacaactt ttcactagga gtcttgggaa tggcattgtt cctggaggaa 3540

atgcttagga cccgagtagg aacgaaacat gcaatactac tagttgcagt ttcttttgtg 3600

acattgatca cagggaacat gtcctttaga gacctgggaa gagtgatggt tatggtaggc 3660
gccactatga cgggtgacat aggtatgggc gtgacttatc ttgccctact agcagccttc 3720
aaagtcagac caacttttgc agctggacta ctcttgagaa agctgacctc cagggaattg 3780
atgatgacta ctataggaat tgtactcctc tcccagagca ccataccaga gaccattctt 3840
gagttgactg atgcgttagc cttaggcatg atggtcctca aaatggtgag aaatatggaa 3900
aagtatcaat tggcagtgac tatcatggct atcttgtgcg tcccaaacgc agtgatatta 3960
caaaacgcat ggaaagtgag ttgcacaata ttggcagtgg tgtccgtttc cccactgttc 4020
ttaacatcct cacagcaaaa aacagattgg ataccattag cattgacgat caaaggtctc 4080
aatccaacag ctatttttct aacaaccctc tcaagaacca gcaagaaaag gagctggcca 4140
ttaaatgagg ctatcatggc agtcgggatg gtgagcattt tagccagttc tctcctaaaa 4200
aatgatattc ccatgacagg accattagtg gctggagggc tcctcactgt gtgctacgtg 4260
ctcactggac gatcggccga tttggaactg gagagagcag ccgatgtcaa atgggaagac 4320
caggcagaga tatcaggaag cagtccaatc ctgtcaataa caatatcaga agatggtagc 4380
atgtcgataa aaaatgaaga ggaagaacaa acactgacca tactcattag aacaggattg 4440
ctggtgatct caggactttt tcctgtatca ataccaatca cggcagcagc atggtacctg 4500
tgggaagtga agaaacaacg ggccggagta ttgtgggatg ttccttcacc cccacccatg 4560
ggaaaggctg aactggaaga tggagcctat agaattaagc aaaaagggat tcttggatat 4620
tcccagatcg gagccggagt ttacaaagaa ggaacattcc atacaatgtg gcatgtcaca 4680
cgtggcgctg ttctaatgca taaaggaaag aggattgaac catcatgggc ggacgtcaag 4740
aaagacctaa tatcatatgg aggaggctgg aagttagaag gagaatggaa ggaaggagaa 4800
gaagtccagg tattggcact ggagcctgga aaaaatccaa gagccgtcca aacgaaacct 4860
ggtcttttca aaaccaacgc cggaacaata ggtgctgtat ctctggactt ttctcctgga 4920
acgtcaggat ctccaattat cgacaaaaaa ggaaaagttg tgggtcttta tggtaatggt 4980
gttgttacaa ggagtggagc atatgtgagt gctatagccc agactgaaaa aagcattgaa 5040
gacaacccag agatcgaaga tgacattttc cgaaagagaa gactgaccat catggacctc 5100
cacccaggag cgggaaagac gaagagatac cttccggcca tagtcagaga agctataaaa 5160
cggggtttga gaacattaat cttggccccc actagagttg tggcagctga aatggaggaa 5220
gcccttagag gacttccaat aagataccag accccagcca tcagagctgt gcacaccggg 5280
cgggagattg tggacctaat gtgtcatgcc acatttacca tgaggctgct atcaccagtt 5340
agagtgccaa actacaacct gattatcatg gacgaagccc atttcacaga tccagcaagt 5400
atagcagcta gaggatacat ctcaactcga gtggagatgg gtgaggcagc tgggattttt 5460
atgacagcca ctcccccggg aagcagagac ccatttcctc agagcaatgc accaatcata 5520
gatgaagaaa gagaaatccc tgaacgctcg tggaattccg gacatgaatg ggtcacggat 5580
tttaaaggga agactgtttg gttcgttcca agtataaaag caggaaatga tatagcagct 5640

tgcctgagga aaaatggaaa gaaagtgata caactcagta ggaagacctt tgattctgag 5700

tatgtcaaga ctagaaccaa tgattgggac ttcgtggtta caactgacat ttcagaaatg 5760

ggtgccaatt tcaaggctga gagggttata gaccccagac gctgcatgaa accagtcata 5820

ctaacagatg gtgaagagcg ggtgattctg gcaggaccta tgccagtgac ccactctagt 5880

gcagcacaaa gaagagggag aataggaaga aatccaaaaa atgagaatga ccagtacata 5940

tacatggggg aacctctgga aaatgatgaa gactgtgcac actggaaaga agctaaaatg 6000

ctcctagata acatcaacac gccagaagga atcattccta gcatgttcga accagagcgt 6060

gaaaaggtgg atgccattga tggcgaatac cgcttgagag gagaagcaag gaaaaccttt 6120

gtagacttaa tgagaagagg agacctacca gtctggttgg cctacagagt ggcagctgaa 6180

ggcatcaact acgcagacag aaggtggtgt tttgatggag tcaagaacaa ccaaatccta 6240

gaagaaaacg tggaagttga aatctggaca aaagaagggg aaaggaagaa attgaaaccc 6300

agatggttgg atgctaggat ctattctgac ccactggcgc taaaagaatt taaggaattt 6360

gcagccggaa gaaagtctct gaccctgaac ctaatcacag aaatgggtag gctcccaacc 6420

ttcatgactc agaaggtaag agacgcactg gacaacttag cagtgctgca cacggctgag 6480

gcaggtggaa gggcgtacaa ccatgctctc agtgaactgc cggagaccct ggagacattg 6540

cttttactga cacttctggc tacagtcacg ggagggatct ttttattctt gatgagcgca 6600

aggggcatag ggaagatgac cctgggaatg tgctgcataa tcacggctag catcctccta 6660

tggtacgcac aaatacagcc acactggata gcagcttcaa taatactgga gtttttttctc 6720

atagttttgc ttattccaga acctgaaaaa cagagaacac cccaagacaa ccaactgacc 6780

tacgttgtca tagccatcct cacagtggtg gccgcaacca tggcaaacga gatgggtttc 6840

ctagaaaaaa cgaagaaaga tctcggattg ggaagcattg caacccagca acccgagagc 6900

aacatcctgg acatagatct acgtcctgca tcagcatgga cgctgtatgc cgtggccaca 6960

acatttgtta caccaatgtt gagacatagc attgaaaatt cctcagtgaa tgtgtcccta 7020

acagccatag ccaaccaagc cacagtgtta atgggtctcg ggaaaggatg gccattgtca 7080

aagatggaca tcggagttcc ccttctcgcc attggatgct actcacaagt caaccccata 7140

actctcacag cagctctttt cttattggta gcacattatg ccatcatagg gccaggactc 7200

caagcaaaag caaccagaga agctcagaaa agagcagcgg cgggcatcat gaaaaaccca 7260

actgtcgatg gaataacagt gattgaccta gatccaatac cttatgatcc aaagtttgaa 7320

aagcagttgg gacaagtaat gctcctagtc ctctgcgtga ctcaagtatt gatgatgagg 7380

actacatggg ctctgtgtga ggctttaacc ttagctaccg ggcccatctc cacattgtgg 7440

gaaggaaatc cagggaggtt ttggaacact accattgcgg tgtcaatggc taacattttt 7500

agagggagtt acttggccgg agctggactt ctcttttta ttatgaagaa cacaaccaac 7560

acaagaaggg gaactggcaa cataggagag acgcttggag agaaatggaa aagccgattg 7620

aacgcattgg gaaaaagtga attccagatc tacaagaaaa gtggaatcca ggaagtggat 7680

agaaccttag caaaagaagg cattaaaaga ggagaaacgg accatcacgc tgtgtcgcga 7740

ggctcagcaa aactgagatg gttcgttgag agaaacatgg tcacaccaga agggaaagta 7800

gtggacctcg gttgtggcag aggaggctgg tcatactatt gtggaggact aaagaatgta 7860

agagaagtca aaggcctaac aaaaggagga ccaggacacg aagaacccat ccccatgtca 7920

acatatgggt ggaatctagt gcgtcttcaa agtggagttg acgttttctt catcccgcca 7980

gaaaagtgtg acacattatt gtgtgacata ggggagtcat caccaaatcc cacagtggaa 8040

gcaggacgaa cactcagagt ccttaactta gtagaaaatt ggttgaacaa caacactcaa 8100

ttttgcataa aggttctcaa cccatatatg ccctcagtca tagaaaaaat ggaagcacta 8160

caaaggaaat atggaggagc cttagtgagg aatccactct cacgaaactc cacacatgag 8220

atgtactggg tatccaatgc ttccgggaac atagtgtcat cagtgaacat gatttcaagg 8280

atgttgatca acagatttac aatgagatac aagaaagcca cttacgagcc ggatgttgac 8340

ctcggaagcg gaacccgtaa catcgggatt gaaagtgaga taccaaacct agatataatt 8400

gggaaaagaa tagaaaaaat aaagcaagag catgaaacat catggcacta tgaccaagac 8460

cacccataca aaacgtgggc ataccatggt agctatgaaa caaaacagac tggatcagca 8520

tcatccatgg tcaacggagt ggtcaggctg ctgacaaaac cttgggacgt cgtccccatg 8580

gtgacacaga tggcaatgac agacacgact ccatttggac aacagcgcgt ttttaaagag 8640

aaagtggaca cgagaaccca agaaccgaaa gaaggcacga agaaactaat gaaaataaca 8700

gcagagtggc tttggaaaga attagggaag aaaaagacac ccaggatgtg caccagagaa 8760

gaattcacaa gaaaggtgag aagcaatgca gccttggggg ccatattcac tgatgagaac 8820

aagtggaagt cggcacgtga ggctgttgaa gatagtaggt tttgggagct ggttgacaag 8880

gaaaggaatc tccatcttga aggaaagtgt gaaacatgtg tgtacaacat gatgggaaaa 8940

agagagaaga agctagggga attcggcaag gcaaaaggca gcagagccat atggtacatg 9000

tggcttggag cacgcttctt agagtttgaa gccctaggat tcttaaatga agatcactgg 9060

ttctccagag agaactccct gagtggagtg gaaggagaag ggctgcacaa gctaggttac 9120

attctaagag acgtgagcaa gaaagaggga ggagcaatgt atgccgatga caccgcagga 9180

tgggatacaa gaatcacact agaagaccta aaaaatgaag aaatggtaac aaaccacatg 9240

gaaggagaac acaagaaact agccgaggcc attttcaaac taacgtacca aaacaaggtg 9300

gtgcgtgtgc aaagaccaac accaagaggc acagtaatgg acatcatatc gagaagagac 9360

caaagaggta gtggacaagt tggcacctat ggactcaata ctttcaccaa tatggaagcc 9420

caactaatca gacagatgga gggagaagga gtctttaaaa gcattcagca cctaacaatc 9480

acagaagaaa tcgctgtgca aaactggtta gcaagagtgg ggcgcgaaag gttatcaaga 9540

atggccatca gtggagatga ttgtgttgtg aaacctttag atgacaggtt cgcaagcgct 9600

ttaacagctc taaatgacat gggaaagatt aggaaagaca tacaacaatg ggaaccttca 9660

agaggatgga atgattggac acaagtgccc ttctgttcac accatttcca tgagttaatc 9720

atgaaagacg gtcgcgtact cgttgttccc tgtagaaacc aagatgaact gattggcaga 9780

gcccgaatct cccaaggagc agggtggtct ttgcgggaga cggcctgttt ggggaagtct 9840

tacgcccaaa tgtggagctt gatgtacttc cacagacgcg acctcaggct ggcggcaaat 9900

gctatttgct cggcagtacc atcacattgg gttccaacaa gtcgaacaac ctggtccata 9960

catgctaaac atgaatggat gacaacggaa gacatgctga cagtctggaa cagggtgtgg 10020

attcaagaaa acccatggat ggaagacaaa actccagtgg aatcatggga ggaaatccca 10080

tacttgggga aaagagaaga ccaatggtgc ggctcattga ttgggttaac aagcagggcc 10140

acctgggcaa agaacatcca agcagcaata aatcaagtta gatcccttat aggcaatgaa 10200

gaatacacag attacatgcc atccatgaaa agattcagaa gagaagagga agaagcagga 10260

gttctgtggt agaaagcaaa actaacatga aacaaggcta gaagtcaggt cggattaagc 10320

catagtacgg aaaaaactat gctacctgtg agccccgtcc aaggacgtta aaagaagtca 10380

ggccatcata aatgccatag cttgagtaaa ctatgcagcc tgtagctcca cctgagaagg 10440

tgtaaaaaat ccgggaggcc acaaaccatg gaagctgtac gcatggcgta gtggactagc 10500

ggttagagga gacccctccc ttacaaatcg cagcaacaat gggggcccaa ggcgagatga 10560

agctgtagtc tcgctggaag gactagaggt tagaggagac ccccccgaaa caaaaaacag 10620

catattgacg ctgggaaaga ccagagatcc tgctgtctcc tcagcatcat tccaggcaca 10680

gaacgccaga aaatggaatg gtgctgttga atcaacaggt tct 10723

<210> 11
<211> 3391
<212> PRT
<213> Artificial Sequence

<220>
<223> Dengue virus serotype 4, BVS

<400> 11

Met Asn Asn Gln Arg Lys Lys Ala Lys Asn Thr Pro Phe Asn Met Leu
1 5 10 15

Lys Arg Glu Arg Asn Arg Val Ser Thr Val Gln Gln Leu Thr Lys Arg
20 25 30

Phe Ser Leu Gly Met Leu Gln Gly Arg Gly Pro Leu Lys Leu Phe Met
35 40 45

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

Ile Leu Lys Arg Trp Gly Thr Ile Lys Lys Ser Lys Ala Ile Asn Val
65 70 75 80

Leu Arg Gly Phe Arg Lys Glu Ile Gly Arg Met Leu Asn Ile Leu Asn
85 90 95

```
Arg Arg Arg Ser Ser Ala Gly Met Ile Ile Met Leu Ile Pro Thr Val
            100                 105                 110

Met Ala Phe His Leu Thr Thr Arg Asp Gly Glu Pro Leu Met Ile Val
        115                 120                 125

Ala Lys His Glu Arg Gly Arg Pro Leu Leu Phe Lys Thr Thr Glu Gly
    130                 135                 140

Ile Asn Lys Cys Thr Leu Ile Ala Met Asp Leu Gly Glu Met Cys Glu
145                 150                 155                 160

Asp Thr Val Thr Tyr Lys Cys Pro Leu Leu Val Asn Thr Glu Pro Glu
                165                 170                 175

Asp Ile Asp Cys Trp Cys Asn Leu Thr Ser Thr Trp Val Met Tyr Gly
            180                 185                 190

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

Leu Thr Pro His Ser Gly Met Gly Leu Glu Thr Arg Ala Glu Thr Trp
    210                 215                 220

Met Ser Ser Glu Gly Ala Trp Lys His Ala Gln Arg Val Glu Ser Trp
225                 230                 235                 240

Ile Leu Arg Asn Pro Gly Phe Ala Leu Leu Ala Gly Phe Met Ala Tyr
                245                 250                 255

Met Ile Gly Gln Thr Gly Ile Gln Arg Thr Val Phe Phe Val Leu Met
            260                 265                 270

Met Leu Val Ala Pro Ser Tyr Gly Met Arg Cys Val Gly Val Gly Asn
        275                 280                 285

Arg Asp Phe Val Glu Gly Val Ser Gly Gly Ala Trp Val Asp Leu Val
    290                 295                 300

Leu Glu His Gly Gly Cys Val Thr Thr Met Ala Gln Gly Lys Pro Thr
305                 310                 315                 320

Leu Asp Phe Glu Leu Thr Lys Thr Thr Ala Lys Glu Val Ala Leu Leu
                325                 330                 335

Arg Thr Tyr Cys Ile Glu Ala Ser Ile Ser Asn Ile Thr Thr Ala Thr
            340                 345                 350

Arg Cys Pro Thr Gln Gly Glu Pro Tyr Leu Lys Glu Glu Gln Asp Gln
        355                 360                 365
```

```
Gln Tyr Ile Cys Arg Arg Asp Val Val Asp Arg Gly Trp Gly Asn Gly
    370                 375                 380

Cys Gly Leu Phe Gly Lys Gly Gly Val Val Thr Cys Ala Lys Phe Ser
385                 390                 395                 400

Cys Ser Gly Lys Ile Thr Gly Asn Leu Val Gln Ile Glu Asn Leu Glu
                405                 410                 415

Tyr Thr Val Val Val Thr Val His Asn Gly Asp Thr His Ala Val Gly
            420                 425                 430

Asn Asp Thr Ser Asn His Gly Val Thr Ala Thr Ile Thr Pro Arg Ser
        435                 440                 445

Pro Ser Val Glu Val Lys Leu Pro Asp Tyr Gly Glu Leu Thr Leu Asp
    450                 455                 460

Cys Glu Pro Arg Ser Gly Ile Asp Phe Asn Glu Met Ile Leu Met Lys
465                 470                 475                 480

Met Lys Lys Lys Thr Trp Leu Val His Lys Gln Trp Phe Leu Asp Leu
                485                 490                 495

Pro Leu Pro Trp Thr Ala Gly Ala Asp Thr Ser Glu Val His Trp Asn
            500                 505                 510

Tyr Lys Glu Arg Met Val Thr Phe Lys Val Pro His Ala Lys Arg Gln
        515                 520                 525

Asp Val Thr Val Leu Gly Ser Gln Glu Gly Ala Met His Ser Ala Leu
    530                 535                 540

Ala Gly Ala Thr Glu Val Asp Ser Gly Asp Gly Asn His Met Phe Ala
545                 550                 555                 560

Gly His Leu Lys Cys Lys Val Arg Met Glu Lys Leu Arg Ile Lys Gly
                565                 570                 575

Met Ser Tyr Thr Met Cys Ser Gly Lys Phe Ser Ile Asp Lys Glu Met
            580                 585                 590

Ala Glu Thr Gln His Gly Thr Thr Val Val Lys Val Lys Tyr Glu Gly
        595                 600                 605

Ala Gly Ala Pro Cys Lys Val Pro Ile Glu Ile Arg Asp Val Asn Lys
    610                 615                 620

Glu Lys Val Val Gly Arg Ile Ile Ser Ser Thr Pro Leu Ala Glu Asn
625                 630                 635                 640
```

```
Thr Asn Ser Val Thr Asn Ile Glu Leu Glu Pro Pro Phe Gly Asp Ser
                645                 650                 655

Tyr Ile Val Ile Gly Val Gly Asn Ser Ala Leu Thr Leu His Trp Phe
            660                 665                 670

Arg Lys Gly Ser Ser Ile Gly Lys Met Phe Glu Ser Thr Tyr Arg Gly
        675                 680                 685

Ala Lys Arg Met Ala Ile Leu Gly Glu Thr Ala Trp Asp Phe Gly Ser
    690                 695                 700

Val Gly Gly Leu Phe Thr Ser Leu Gly Lys Ala Val His Gln Val Phe
705                 710                 715                 720

Gly Ser Val Tyr Thr Thr Leu Phe Gly Gly Val Ser Trp Met Ile Arg
                725                 730                 735

Ile Leu Ile Gly Phe Leu Val Leu Trp Ile Gly Thr Asn Ser Arg Asn
            740                 745                 750

Thr Ser Met Ala Met Thr Cys Ile Ala Ala Gly Ile Val Thr Leu Tyr
        755                 760                 765

Leu Gly Val Met Val Gln Ala Asp Ser Gly Cys Val Val Ser Trp Lys
    770                 775                 780

Asn Lys Glu Leu Lys Cys Gly Ser Gly Ile Phe Ile Thr Asp Asn Val
785                 790                 795                 800

His Thr Trp Thr Glu Gln Tyr Lys Phe Gln Pro Glu Ser Pro Ser Lys
                805                 810                 815

Leu Ala Ser Ala Ile Gln Lys Ala His Glu Glu Asp Ile Cys Gly Ile
            820                 825                 830

Arg Ser Val Thr Arg Leu Glu Asn Leu Met Trp Lys Gln Ile Thr Pro
        835                 840                 845

Glu Leu Asn His Ile Leu Ser Glu Asn Glu Val Lys Leu Thr Ile Met
    850                 855                 860

Thr Gly Asp Ile Lys Gly Ile Met Gln Ala Gly Lys Arg Ser Leu Arg
865                 870                 875                 880

Pro Gln Pro Thr Glu Leu Lys Tyr Ser Trp Lys Thr Trp Gly Lys Ala
                885                 890                 895

Lys Met Leu Ser Thr Glu Ser His Asn Gln Thr Phe Leu Ile Asp Gly
            900                 905                 910
```

```
Pro Glu Thr Ala Glu Cys Pro Asn Thr Asn Arg Ala Trp Asn Ser Leu
        915                 920                 925

Glu Val Glu Asp Tyr Gly Phe Gly Val Phe Thr Thr Asn Ile Trp Leu
    930                 935                 940

Lys Leu Lys Glu Lys Gln Asp Val Phe Cys Asp Ser Lys Leu Met Ser
945                 950                 955                 960

Ala Ala Ile Lys Asp Asn Arg Ala Val His Ala Asp Met Gly Tyr Trp
                965                 970                 975

Ile Glu Ser Ala Leu Asn Asp Thr Trp Lys Ile Glu Lys Ala Ser Phe
            980                 985                 990

Ile Glu Val Lys Asn Cys His Trp  Pro Lys Ser His Thr  Leu Trp Ser
        995                 1000                1005

Asn Gly  Val Leu Glu Ser Glu  Met Ile Ile Pro Lys  Asn Leu Ala
    1010                1015                1020

Gly Pro  Val Ser Gln His Asn  Tyr Arg Pro Gly Tyr  His Thr Gln
    1025                1030                1035

Ile Thr  Gly Pro Trp His Leu  Gly Lys Leu Glu Met  Asp Phe Asp
    1040                1045                1050

Phe Cys  Asp Gly Thr Thr Val  Val Val Thr Glu Asp  Cys Gly Asn
    1055                1060                1065

Arg Gly  Pro Ser Leu Arg Thr  Thr Thr Ala Ser Gly  Lys Leu Ile
    1070                1075                1080

Thr Glu  Trp Cys Cys Arg Ser  Cys Thr Leu Pro Pro  Leu Arg Tyr
    1085                1090                1095

Arg Gly  Glu Asp Gly Cys Trp  Tyr Gly Met Glu Ile  Arg Pro Leu
    1100                1105                1110

Lys Glu  Lys Glu Glu Asn Leu  Val Asn Ser Leu Val  Thr Ala Gly
    1115                1120                1125

His Gly  Gln Val Asp Asn Phe  Ser Leu Gly Val Leu  Gly Met Ala
    1130                1135                1140

Leu Phe  Leu Glu Glu Met Leu  Arg Thr Arg Val Gly  Thr Lys His
    1145                1150                1155

Ala Ile  Leu Leu Val Ala Val  Ser Phe Val Thr Leu  Ile Thr Gly
    1160                1165                1170
```

```
Asn Met  Ser Phe Arg Asp Leu  Gly Arg Val Met Val  Met Val Gly
    1175                1180                1185

Ala Thr  Met Thr Gly Asp Ile  Gly Met Gly Val Thr  Tyr Leu Ala
    1190                1195                1200

Leu Leu  Ala Ala Phe Lys Val  Arg Pro Thr Phe Ala  Ala Gly Leu
    1205                1210                1215

Leu Leu  Arg Lys Leu Thr Ser  Arg Glu Leu Met Met  Thr Thr Ile
    1220                1225                1230

Gly Ile  Val Leu Leu Ser Gln  Ser Thr Ile Pro Glu  Thr Ile Leu
    1235                1240                1245

Glu Leu  Thr Asp Ala Leu Ala  Leu Gly Met Met Val  Leu Lys Met
    1250                1255                1260

Val Arg  Asn Met Glu Lys Tyr  Gln Leu Ala Val Thr  Ile Met Ala
    1265                1270                1275

Ile Leu  Cys Val Pro Asn Ala  Val Ile Leu Gln Asn  Ala Trp Lys
    1280                1285                1290

Val Ser  Cys Thr Ile Leu Ala  Val Val Ser Val Ser  Pro Leu Phe
    1295                1300                1305

Leu Thr  Ser Ser Gln Gln Lys  Thr Asp Trp Ile Pro  Leu Ala Leu
    1310                1315                1320

Thr Ile  Lys Gly Leu Asn Pro  Thr Ala Ile Phe Leu  Thr Thr Leu
    1325                1330                1335

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

Met Ala  Val Gly Met Val Ser  Ile Leu Ala Ser Ser  Leu Leu Lys
    1355                1360                1365

Asn Asp  Ile Pro Met Thr Gly  Pro Leu Val Ala Gly  Gly Leu Leu
    1370                1375                1380

Thr Val  Cys Tyr Val Leu Thr  Gly Arg Ser Ala Asp  Leu Glu Leu
    1385                1390                1395

Glu Arg  Ala Ala Asp Val Lys  Trp Glu Asp Gln Ala  Glu Ile Ser
    1400                1405                1410

Gly Ser  Ser Pro Ile Leu Ser  Ile Thr Ile Ser Glu  Asp Gly Ser
    1415                1420                1425
```

```
Met Ser  Ile Lys Asn Glu Glu  Glu Glu Gln Thr Leu  Thr Ile Leu
    1430                 1435                 1440

Ile Arg  Thr Gly Leu Leu Val  Ile Ser Gly Leu Phe  Pro Val Ser
    1445                 1450                 1455

Ile Pro  Ile Thr Ala Ala Ala  Trp Tyr Leu Trp Glu  Val Lys Lys
    1460                 1465                 1470

Gln Arg  Ala Gly Val Leu Trp  Asp Val Pro Ser Pro  Pro Pro Met
    1475                 1480                 1485

Gly Lys  Ala Glu Leu Glu Asp  Gly Ala Tyr Arg Ile  Lys Gln Lys
    1490                 1495                 1500

Gly Ile  Leu Gly Tyr Ser Gln  Ile Gly Ala Gly Val  Tyr Lys Glu
    1505                 1510                 1515

Gly Thr  Phe His Thr Met Trp  His Val Thr Arg Gly  Ala Val Leu
    1520                 1525                 1530

Met His  Lys Gly Lys Arg Ile  Glu Pro Ser Trp Ala  Asp Val Lys
    1535                 1540                 1545

Lys Asp  Leu Ile Ser Tyr Gly  Gly Gly Trp Lys Leu  Glu Gly Glu
    1550                 1555                 1560

Trp Lys  Glu Gly Glu Glu Val  Gln Val Leu Ala Leu  Glu Pro Gly
    1565                 1570                 1575

Lys Asn  Pro Arg Ala Val Gln  Thr Lys Pro Gly Leu  Phe Lys Thr
    1580                 1585                 1590

Asn Ala  Gly Thr Ile Gly Ala  Val Ser Leu Asp Phe  Ser Pro Gly
    1595                 1600                 1605

Thr Ser  Gly Ser Pro Ile Ile  Asp Lys Lys Gly Lys  Val Val Gly
    1610                 1615                 1620

Leu Tyr  Gly Asn Gly Val Val  Thr Arg Ser Gly Ala  Tyr Val Ser
    1625                 1630                 1635

Ala Ile  Ala Gln Thr Glu Lys  Ser Ile Glu Asp Asn  Pro Glu Ile
    1640                 1645                 1650

Glu Asp  Asp Ile Phe Arg Lys  Arg Arg Leu Thr Ile  Met Asp Leu
    1655                 1660                 1665

His Pro  Gly Ala Gly Lys Thr  Lys Arg Tyr Leu Pro  Ala Ile Val
    1670                 1675                 1680
```

```
Arg Glu  Ala Ile Lys Arg Gly  Leu Arg Thr Leu Ile  Leu Ala Pro
    1685                 1690                 1695

Thr Arg  Val Val Ala Ala Glu  Met Glu Glu Ala Leu  Arg Gly Leu
    1700                 1705                 1710

Pro Ile  Arg Tyr Gln Thr Pro  Ala Ile Arg Ala Val  His Thr Gly
    1715                 1720                 1725

Arg Glu  Ile Val Asp Leu Met  Cys His Ala Thr Phe  Thr Met Arg
    1730                 1735                 1740

Leu Leu  Ser Pro Val Arg Val  Pro Asn Tyr Asn Leu  Ile Ile Met
    1745                 1750                 1755

Asp Glu  Ala His Phe Thr Asp  Pro Ala Ser Ile Ala  Ala Arg Gly
    1760                 1765                 1770

Tyr Ile  Ser Thr Arg Val Glu  Met Gly Glu Ala Ala  Gly Ile Phe
    1775                 1780                 1785

Met Thr  Ala Thr Pro Pro Gly  Ser Arg Asp Pro Phe  Pro Gln Ser
    1790                 1795                 1800

Asn Ala  Pro Ile Ile Asp Glu  Glu Arg Glu Ile Pro  Glu Arg Ser
    1805                 1810                 1815

Trp Asn  Ser Gly His Glu Trp  Val Thr Asp Phe Lys  Gly Lys Thr
    1820                 1825                 1830

Val Trp  Phe Val Pro Ser Ile  Lys Ala Gly Asn Asp  Ile Ala Ala
    1835                 1840                 1845

Cys Leu  Arg Lys Asn Gly Lys  Lys Val Ile Gln Leu  Ser Arg Lys
    1850                 1855                 1860

Thr Phe  Asp Ser Glu Tyr Val  Lys Thr Arg Thr Asn  Asp Trp Asp
    1865                 1870                 1875

Phe Val  Val Thr Thr Asp Ile  Ser Glu Met Gly Ala  Asn Phe Lys
    1880                 1885                 1890

Ala Glu  Arg Val Ile Asp Pro  Arg Arg Cys Met Lys  Pro Val Ile
    1895                 1900                 1905

Leu Thr  Asp Gly Glu Glu Arg  Val Ile Leu Ala Gly  Pro Met Pro
    1910                 1915                 1920

Val Thr  His Ser Ser Ala Ala  Gln Arg Arg Gly Arg  Ile Gly Arg
    1925                 1930                 1935
```

```
Asn Pro  Lys Asn Glu Asn Asp  Gln Tyr Ile Tyr Met  Gly Glu Pro
    1940                1945                1950

Leu Glu  Asn Asp Glu Asp Cys  Ala His Trp Lys Glu  Ala Lys Met
    1955                1960                1965

Leu Leu  Asp Asn Ile Asn Thr  Pro Glu Gly Ile Ile  Pro Ser Met
    1970                1975                1980

Phe Glu  Pro Glu Arg Glu Lys  Val Asp Ala Ile Asp  Gly Glu Tyr
    1985                1990                1995

Arg Leu  Arg Gly Glu Ala Arg  Lys Thr Phe Val Asp  Leu Met Arg
    2000                2005                2010

Arg Gly  Asp Leu Pro Val Trp  Leu Ala Tyr Arg Val  Ala Ala Glu
    2015                2020                2025

Gly Ile  Asn Tyr Ala Asp Arg  Arg Trp Cys Phe Asp  Gly Val Lys
    2030                2035                2040

Asn Asn  Gln Ile Leu Glu Glu  Asn Val Glu Val Glu  Ile Trp Thr
    2045                2050                2055

Lys Glu  Gly Glu Arg Lys Lys  Leu Lys Pro Arg Trp  Leu Asp Ala
    2060                2065                2070

Arg Ile  Tyr Ser Asp Pro Leu  Ala Leu Lys Glu Phe  Lys Glu Phe
    2075                2080                2085

Ala Ala  Gly Arg Lys Ser Leu  Thr Leu Asn Leu Ile  Thr Glu Met
    2090                2095                2100

Gly Arg  Leu Pro Thr Phe Met  Thr Gln Lys Val Arg  Asp Ala Leu
    2105                2110                2115

Asp Asn  Leu Ala Val Leu His  Thr Ala Glu Ala Gly  Gly Arg Ala
    2120                2125                2130

Tyr Asn  His Ala Leu Ser Glu  Leu Pro Glu Thr Leu  Glu Thr Leu
    2135                2140                2145

Leu Leu  Leu Thr Leu Leu Ala  Thr Val Thr Gly Gly  Ile Phe Leu
    2150                2155                2160

Phe Leu  Met Ser Ala Arg Gly  Ile Gly Lys Met Thr  Leu Gly Met
    2165                2170                2175

Cys Cys  Ile Ile Thr Ala Ser  Ile Leu Leu Trp Tyr  Ala Gln Ile
    2180                2185                2190
```

```
Gln Pro  His Trp Ile Ala Ala  Ser Ile Ile Leu Glu  Phe Phe Leu
    2195                 2200                 2205

Ile Val  Leu Leu Ile Pro Glu  Pro Glu Lys Gln Arg  Thr Pro Gln
    2210                 2215                 2220

Asp Asn  Gln Leu Thr Tyr Val  Val Ile Ala Ile Leu  Thr Val Val
    2225                 2230                 2235

Ala Ala  Thr Met Ala Asn Glu  Met Gly Phe Leu Glu  Lys Thr Lys
    2240                 2245                 2250

Lys Asp  Leu Gly Leu Gly Ser  Ile Ala Thr Gln Gln  Pro Glu Ser
    2255                 2260                 2265

Asn Ile  Leu Asp Ile Asp Leu  Arg Pro Ala Ser Ala  Trp Thr Leu
    2270                 2275                 2280

Tyr Ala  Val Ala Thr Thr Phe  Val Thr Pro Met Leu  Arg His Ser
    2285                 2290                 2295

Ile Glu  Asn Ser Ser Val Asn  Val Ser Leu Thr Ala  Ile Ala Asn
    2300                 2305                 2310

Gln Ala  Thr Val Leu Met Gly  Leu Gly Lys Gly Trp  Pro Leu Ser
    2315                 2320                 2325

Lys Met  Asp Ile Gly Val Pro  Leu Leu Ala Ile Gly  Cys Tyr Ser
    2330                 2335                 2340

Gln Val  Asn Pro Ile Thr Leu  Thr Ala Ala Leu Phe  Leu Leu Val
    2345                 2350                 2355

Ala His  Tyr Ala Ile Ile Gly  Pro Gly Leu Gln Ala  Lys Ala Thr
    2360                 2365                 2370

Arg Glu  Ala Gln Lys Arg Ala  Ala Ala Gly Ile Met  Lys Asn Pro
    2375                 2380                 2385

Thr Val  Asp Gly Ile Thr Val  Ile Asp Leu Asp Pro  Ile Pro Tyr
    2390                 2395                 2400

Asp Pro  Lys Phe Glu Lys Gln  Leu Gly Gln Val Met  Leu Leu Val
    2405                 2410                 2415

Leu Cys  Val Thr Gln Val Leu  Met Met Arg Thr Thr  Trp Ala Leu
    2420                 2425                 2430

Cys Glu  Ala Leu Thr Leu Ala  Thr Gly Pro Ile Ser  Thr Leu Trp
    2435                 2440                 2445
```

```
Glu Gly  Asn Pro Gly Arg Phe  Trp Asn Thr Thr Ile  Ala Val Ser
    2450                2455                2460

Met Ala  Asn Ile Phe Arg Gly  Ser Tyr Leu Ala Gly  Ala Gly Leu
    2465                2470                2475

Leu Phe  Phe Ile Met Lys Asn  Thr Thr Asn Thr Arg  Arg Gly Thr
    2480                2485                2490

Gly Asn  Ile Gly Glu Thr Leu  Gly Glu Lys Trp Lys  Ser Arg Leu
    2495                2500                2505

Asn Ala  Leu Gly Lys Ser Glu  Phe Gln Ile Tyr Lys  Lys Ser Gly
    2510                2515                2520

Ile Gln  Glu Val Asp Arg Thr  Leu Ala Lys Glu Gly  Ile Lys Arg
    2525                2530                2535

Gly Glu  Thr Asp His His Ala  Val Ser Arg Gly Ser  Ala Lys Leu
    2540                2545                2550

Arg Trp  Phe Val Glu Arg Asn  Met Val Thr Pro Glu  Gly Lys Val
    2555                2560                2565

Val Asp  Leu Gly Cys Gly Arg  Gly Gly Trp Ser Tyr  Tyr Cys Gly
    2570                2575                2580

Gly Leu  Lys Asn Val Arg Glu  Val Lys Gly Leu Thr  Lys Gly Gly
    2585                2590                2595

Pro Gly  His Glu Glu Pro Ile  Pro Met Ser Thr Tyr  Gly Trp Asn
    2600                2605                2610

Leu Val  Arg Leu Gln Ser Gly  Val Asp Val Phe Phe  Ile Pro Pro
    2615                2620                2625

Glu Lys  Cys Asp Thr Leu Leu  Cys Asp Ile Gly Glu  Ser Ser Pro
    2630                2635                2640

Asn Pro  Thr Val Glu Ala Gly  Arg Thr Leu Arg Val  Leu Asn Leu
    2645                2650                2655

Val Glu  Asn Trp Leu Asn Asn  Asn Thr Gln Phe Cys  Ile Lys Val
    2660                2665                2670

Leu Asn  Pro Tyr Met Pro Ser  Val Ile Glu Lys Met  Glu Ala Leu
    2675                2680                2685

Gln Arg  Lys Tyr Gly Gly Ala  Leu Val Arg Asn Pro  Leu Ser Arg
    2690                2695                2700
```

```
Asn Ser  Thr His Glu Met Tyr  Trp Val Ser Asn Ala  Ser Gly Asn
    2705                 2710                 2715

Ile Val  Ser Ser Val Asn Met  Ile Ser Arg Met Leu  Ile Asn Arg
    2720                 2725                 2730

Phe Thr  Met Arg Tyr Lys Lys  Ala Thr Tyr Glu Pro  Asp Val Asp
    2735                 2740                 2745

Leu Gly  Ser Gly Thr Arg Asn  Ile Gly Ile Glu Ser  Glu Ile Pro
    2750                 2755                 2760

Asn Leu  Asp Ile Ile Gly Lys  Arg Ile Glu Lys Ile  Lys Gln Glu
    2765                 2770                 2775

His Glu  Thr Ser Trp His Tyr  Asp Gln Asp His Pro  Tyr Lys Thr
    2780                 2785                 2790

Trp Ala  Tyr His Gly Ser Tyr  Glu Thr Lys Gln Thr  Gly Ser Ala
    2795                 2800                 2805

Ser Ser  Met Val Asn Gly Val  Val Arg Leu Leu Thr  Lys Pro Trp
    2810                 2815                 2820

Asp Val  Val Pro Met Val Thr  Gln Met Ala Met Thr  Asp Thr Thr
    2825                 2830                 2835

Pro Phe  Gly Gln Gln Arg Val  Phe Lys Glu Lys Val  Asp Thr Arg
    2840                 2845                 2850

Thr Gln  Glu Pro Lys Glu Gly  Thr Lys Lys Leu Met  Lys Ile Thr
    2855                 2860                 2865

Ala Glu  Trp Leu Trp Lys Glu  Leu Gly Lys Lys Lys  Thr Pro Arg
    2870                 2875                 2880

Met Cys  Thr Arg Glu Glu Phe  Thr Arg Lys Val Arg  Ser Asn Ala
    2885                 2890                 2895

Ala Leu  Gly Ala Ile Phe Thr  Asp Glu Asn Lys Trp  Lys Ser Ala
    2900                 2905                 2910

Arg Glu  Ala Val Glu Asp Ser  Arg Phe Trp Glu Leu  Val Asp Lys
    2915                 2920                 2925

Glu Arg  Asn Leu His Leu Glu  Gly Lys Cys Glu Thr  Cys Val Tyr
    2930                 2935                 2940

Asn Met  Met Gly Lys Arg Glu  Lys Lys Leu Gly Glu  Phe Gly Lys
    2945                 2950                 2955
```

```
Ala Lys  Gly Ser Arg Ala Ile  Trp Tyr Met Trp Leu  Gly Ala Arg
    2960                2965                2970

Phe Leu  Glu Phe Glu Ala Leu  Gly Phe Leu Asn Glu  Asp His Trp
    2975                2980                2985

Phe Ser  Arg Glu Asn Ser Leu  Ser Gly Val Glu Gly  Glu Gly Leu
    2990                2995                3000

His Lys  Leu Gly Tyr Ile Leu  Arg Asp Val Ser Lys  Lys Glu Gly
    3005                3010                3015

Gly Ala  Met Tyr Ala Asp Asp  Thr Ala Gly Trp Asp  Thr Arg Ile
    3020                3025                3030

Thr Leu  Glu Asp Leu Lys Asn  Glu Glu Met Val Thr  Asn His Met
    3035                3040                3045

Glu Gly  Glu His Lys Lys Leu  Ala Glu Ala Ile Phe  Lys Leu Thr
    3050                3055                3060

Tyr Gln  Asn Lys Val Val Arg  Val Gln Arg Pro Thr  Pro Arg Gly
    3065                3070                3075

Thr Val  Met Asp Ile Ile Ser  Arg Arg Asp Gln Arg  Gly Ser Gly
    3080                3085                3090

Gln Val  Gly Thr Tyr Gly Leu  Asn Thr Phe Thr Asn  Met Glu Ala
    3095                3100                3105

Gln Leu  Ile Arg Gln Met Glu  Gly Glu Gly Val Phe  Lys Ser Ile
    3110                3115                3120

Gln His  Leu Thr Ile Thr Glu  Glu Ile Ala Val Gln  Asn Trp Leu
    3125                3130                3135

Ala Arg  Val Gly Arg Glu Arg  Leu Ser Arg Met Ala  Ile Ser Gly
    3140                3145                3150

Asp Asp  Cys Val Val Lys Pro  Leu Asp Asp Arg Phe  Ala Ser Ala
    3155                3160                3165

Leu Thr  Ala Leu Asn Asp Met  Gly Lys Ile Arg Lys  Asp Ile Gln
    3170                3175                3180

Gln Trp  Glu Pro Ser Arg Gly  Trp Asn Asp Trp Thr  Gln Val Pro
    3185                3190                3195

Phe Cys  Ser His His Phe His  Glu Leu Ile Met Lys  Asp Gly Arg
    3200                3205                3210
```

Val Leu Val Val Pro Cys Arg Asn Gln Asp Glu Leu Ile Gly Arg
3215 3220 3225

Ala Arg Ile Ser Gln Gly Ala Gly Trp Ser Leu Arg Glu Thr Ala
3230 3235 3240

Cys Leu Gly Lys Ser Tyr Ala Gln Met Trp Ser Leu Met Tyr Phe
3245 3250 3255

His Arg Arg Asp Leu Arg Leu Ala Ala Asn Ala Ile Cys Ser Ala
3260 3265 3270

Val Pro Ser His Trp Val Pro Thr Ser Arg Thr Thr Trp Ser Ile
3275 3280 3285

His Ala Lys His Glu Trp Met Thr Thr Glu Asp Met Leu Thr Val
3290 3295 3300

Trp Asn Arg Val Trp Ile Gln Glu Asn Pro Trp Met Glu Asp Lys
3305 3310 3315

Thr Pro Val Glu Ser Trp Glu Glu Ile Pro Tyr Leu Gly Lys Arg
3320 3325 3330

Glu Asp Gln Trp Cys Gly Ser Leu Ile Gly Leu Thr Ser Arg Ala
3335 3340 3345

Thr Trp Ala Lys Asn Ile Gln Ala Ala Ile Asn Gln Val Arg Ser
3350 3355 3360

Leu Ile Gly Asn Glu Glu Tyr Thr Asp Tyr Met Pro Ser Met Lys
3365 3370 3375

Arg Phe Arg Arg Glu Glu Glu Glu Ala Gly Val Leu Trp
3380 3385 3390

<210> 12
<211> 3391
<212> PRT
<213> Artificial Sequence

<220>
<223> Dengue virus serotype 4, MVS

<400> 12

Met Asn Asn Gln Arg Lys Lys Ala Lys Asn Thr Pro Phe Asn Met Leu
1 5 10 15

Lys Arg Glu Arg Asn Arg Val Ser Thr Val Gln Gln Leu Thr Lys Arg
20 25 30

Phe Ser Leu Gly Met Leu Gln Gly Arg Gly Pro Leu Lys Leu Phe Met

```
        35                  40                  45

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

Ile Leu Lys Arg Trp Gly Thr Ile Lys Lys Ser Lys Ala Ile Asn Val
65                  70                  75                  80

Leu Arg Gly Phe Arg Lys Glu Ile Gly Arg Met Leu Asn Ile Leu Asn
                85                  90                  95

Arg Arg Arg Ser Ser Ala Gly Met Ile Ile Met Leu Ile Pro Thr Val
            100                 105                 110

Met Ala Phe His Leu Thr Thr Arg Asp Gly Glu Pro Leu Met Ile Val
        115                 120                 125

Ala Lys His Glu Arg Gly Arg Pro Leu Leu Phe Lys Thr Thr Glu Gly
    130                 135                 140

Ile Asn Lys Cys Thr Leu Ile Ala Met Asp Leu Gly Glu Met Cys Glu
145                 150                 155                 160

Asp Thr Val Thr Tyr Lys Cys Pro Leu Leu Val Asn Thr Glu Pro Glu
                165                 170                 175

Asp Ile Asp Cys Trp Cys Asn Leu Thr Ser Thr Trp Val Met Tyr Gly
            180                 185                 190

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

Leu Thr Pro His Ser Gly Met Gly Leu Glu Thr Arg Ala Glu Thr Trp
    210                 215                 220

Met Ser Ser Glu Gly Ala Trp Lys His Ala Gln Arg Val Glu Ser Trp
225                 230                 235                 240

Ile Leu Arg Asn Pro Gly Phe Ala Leu Leu Ala Gly Phe Met Ala Tyr
                245                 250                 255

Met Ile Gly Gln Thr Gly Ile Gln Arg Thr Val Phe Phe Val Leu Met
            260                 265                 270

Met Leu Val Ala Pro Ser Tyr Gly Met Arg Cys Val Gly Val Gly Asn
        275                 280                 285

Arg Asp Phe Val Glu Gly Val Ser Gly Gly Ala Trp Val Asp Leu Val
    290                 295                 300

Leu Glu His Gly Gly Cys Val Thr Thr Met Ala Gln Gly Lys Pro Thr
```

305 310 315 320

Leu Asp Phe Glu Leu Thr Lys Thr Thr Ala Lys Glu Val Ala Leu Leu
325 330 335

Arg Thr Tyr Cys Ile Glu Ala Ser Ile Ser Asn Ile Thr Thr Ala Thr
340 345 350

Arg Cys Pro Thr Gln Gly Glu Pro Tyr Leu Lys Glu Glu Gln Asp Gln
355 360 365

Gln Tyr Ile Cys Arg Arg Asp Val Val Asp Arg Gly Trp Gly Asn Gly
370 375 380

Cys Gly Leu Phe Gly Lys Gly Gly Val Val Thr Cys Ala Lys Phe Ser
385 390 395 400

Cys Ser Gly Lys Ile Thr Gly Asn Leu Val Gln Ile Glu Asn Leu Glu
405 410 415

Tyr Thr Val Val Val Thr Val His Asn Gly Asp Thr His Ala Val Gly
420 425 430

Asn Asp Thr Ser Asn His Gly Val Thr Ala Thr Ile Thr Pro Arg Ser
435 440 445

Pro Ser Val Glu Val Lys Leu Pro Asp Tyr Gly Glu Leu Thr Leu Asp
450 455 460

Cys Glu Pro Arg Ser Gly Ile Asp Phe Asn Glu Met Ile Leu Met Lys
465 470 475 480

Met Lys Lys Lys Thr Trp Leu Val His Lys Gln Trp Phe Leu Asp Leu
485 490 495

Pro Leu Pro Trp Thr Ala Gly Ala Asp Thr Ser Glu Val His Trp Asn
500 505 510

Tyr Lys Glu Arg Met Val Thr Phe Lys Val Pro His Ala Lys Arg Gln
515 520 525

Asp Val Thr Val Leu Gly Ser Gln Glu Gly Ala Met His Ser Ala Leu
530 535 540

Ala Gly Ala Thr Glu Val Asp Ser Gly Asp Gly Asn His Met Phe Ala
545 550 555 560

Gly His Leu Lys Cys Lys Val Arg Met Glu Lys Leu Arg Ile Lys Gly
565 570 575

Met Ser Tyr Thr Met Cys Ser Gly Lys Phe Ser Ile Asp Lys Glu Met

580 585 590

Ala Glu Thr Gln His Gly Thr Thr Val Val Lys Val Lys Tyr Glu Gly
595 600 605

Ala Gly Ala Pro Cys Lys Val Pro Ile Glu Ile Arg Asp Val Asn Lys
610 615 620

Glu Lys Val Val Gly Arg Ile Ile Ser Ser Thr Pro Leu Ala Glu Asn
625 630 635 640

Thr Asn Ser Val Thr Asn Ile Glu Leu Glu Pro Pro Phe Gly Asp Ser
645 650 655

Tyr Ile Val Ile Gly Val Gly Asn Ser Ala Leu Thr Leu His Trp Phe
660 665 670

Arg Lys Gly Ser Ser Ile Gly Lys Met Phe Glu Ser Thr Tyr Arg Gly
675 680 685

Ala Lys Arg Met Ala Ile Leu Gly Glu Thr Ala Trp Asp Phe Gly Ser
690 695 700

Val Gly Gly Leu Phe Thr Ser Leu Gly Lys Ala Val His Gln Val Phe
705 710 715 720

Gly Ser Val Tyr Thr Thr Leu Phe Gly Gly Val Ser Trp Met Ile Arg
725 730 735

Ile Leu Ile Gly Phe Leu Val Leu Trp Ile Gly Thr Asn Ser Arg Asn
740 745 750

Thr Ser Met Ala Met Thr Cys Ile Ala Ala Gly Ile Val Thr Leu Tyr
755 760 765

Leu Gly Val Met Val Gln Ala Asp Ser Gly Cys Val Val Ser Trp Lys
770 775 780

Asn Lys Glu Leu Lys Cys Gly Ser Gly Ile Phe Ile Thr Asp Asn Val
785 790 795 800

His Thr Trp Thr Glu Gln Tyr Lys Phe Gln Pro Glu Ser Pro Ser Lys
805 810 815

Leu Ala Ser Ala Ile Gln Lys Ala His Glu Glu Asp Ile Cys Gly Ile
820 825 830

Arg Ser Val Thr Arg Leu Glu Asn Leu Met Trp Lys Gln Ile Thr Pro
835 840 845

Glu Leu Asn His Ile Leu Ser Glu Asn Glu Val Lys Leu Thr Ile Met

```
    850                 855                 860

Thr Gly Asp Ile Lys Gly Ile Met Gln Ala Gly Lys Arg Ser Leu Arg
865                 870                 875                 880

Pro Gln Pro Thr Glu Leu Lys Tyr Ser Trp Lys Thr Trp Gly Lys Ala
                885                 890                 895

Lys Met Leu Ser Thr Glu Ser His Asn Gln Thr Phe Leu Ile Asp Gly
            900                 905                 910

Pro Glu Thr Ala Glu Cys Pro Asn Thr Asn Arg Ala Trp Asn Ser Leu
        915                 920                 925

Glu Val Glu Asp Tyr Gly Phe Gly Val Phe Thr Thr Asn Ile Trp Leu
    930                 935                 940

Lys Leu Lys Glu Lys Gln Asp Val Phe Cys Asp Ser Lys Leu Met Ser
945                 950                 955                 960

Ala Ala Ile Lys Asp Asn Arg Ala Val His Ala Asp Met Gly Tyr Trp
                965                 970                 975

Ile Glu Ser Ala Leu Asn Asp Thr Trp Lys Ile Glu Lys Ala Ser Phe
            980                 985                 990

Ile Glu Val Lys Asn Cys His Trp  Pro Lys Ser His Thr  Leu Trp Ser
        995                 1000                1005

Asn Gly  Val Leu Glu Ser Glu  Met Ile Ile Pro Lys  Asn Leu Ala
    1010                1015                1020

Gly Pro  Val Ser Gln His Asn  Tyr Arg Pro Gly Tyr  His Thr Gln
    1025                1030                1035

Ile Thr  Gly Pro Trp His Leu  Gly Lys Leu Glu Met  Asp Phe Asp
    1040                1045                1050

Phe Cys  Asp Gly Thr Thr Val  Val Val Thr Glu Asp  Cys Gly Asn
    1055                1060                1065

Arg Gly  Pro Ser Leu Arg Thr  Thr Thr Ala Ser Gly  Lys Leu Ile
    1070                1075                1080

Thr Glu  Trp Cys Cys Arg Ser  Cys Thr Leu Pro Pro  Leu Arg Tyr
    1085                1090                1095

Arg Gly  Glu Asp Gly Cys Trp  Tyr Gly Met Glu Ile  Arg Pro Leu
    1100                1105                1110

Lys Glu  Lys Glu Glu Asn Leu  Val Asn Ser Leu Val  Thr Ala Gly
```

1115 1120 1125

His Gly Gln Val Asp Asn Phe Ser Leu Gly Val Leu Gly Met Ala
1130 1135 1140

Leu Phe Leu Glu Glu Met Leu Arg Thr Arg Val Gly Thr Lys His
1145 1150 1155

Ala Ile Leu Leu Val Ala Val Ser Phe Val Thr Leu Ile Thr Gly
1160 1165 1170

Asn Met Ser Phe Arg Asp Leu Gly Arg Val Met Val Met Val Gly
1175 1180 1185

Ala Thr Met Thr Gly Asp Ile Gly Met Gly Val Thr Tyr Leu Ala
1190 1195 1200

Leu Leu Ala Ala Phe Lys Val Arg Pro Thr Phe Ala Ala Gly Leu
1205 1210 1215

Leu Leu Arg Lys Leu Thr Ser Arg Glu Leu Met Met Thr Thr Ile
1220 1225 1230

Gly Ile Val Leu Leu Ser Gln Ser Thr Ile Pro Glu Thr Ile Leu
1235 1240 1245

Glu Leu Thr Asp Ala Leu Ala Leu Gly Met Met Val Leu Lys Met
1250 1255 1260

Val Arg Asn Met Glu Lys Tyr Gln Leu Ala Val Thr Ile Met Ala
1265 1270 1275

Ile Leu Cys Val Pro Asn Ala Val Ile Leu Gln Asn Ala Trp Lys
1280 1285 1290

Val Ser Cys Thr Ile Leu Ala Val Val Ser Val Ser Pro Leu Phe
1295 1300 1305

Leu Thr Ser Ser Gln Gln Lys Thr Asp Trp Ile Pro Leu Ala Leu
1310 1315 1320

Thr Ile Lys Gly Leu Asn Pro Thr Ala Ile Phe Leu Thr Thr Leu
1325 1330 1335

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

Met Ala Val Gly Met Val Ser Ile Leu Ala Ser Ser Leu Leu Lys
1355 1360 1365

Asn Asp Ile Pro Met Thr Gly Pro Leu Val Ala Gly Gly Leu Leu

1370 1375 1380

Thr Val Cys Tyr Val Leu Thr Gly Arg Ser Ala Asp Leu Glu Leu
1385 1390 1395

Glu Arg Ala Ala Asp Val Lys Trp Glu Asp Gln Ala Glu Ile Ser
1400 1405 1410

Gly Ser Ser Pro Ile Leu Ser Ile Thr Ile Ser Glu Asp Gly Ser
1415 1420 1425

Met Ser Ile Lys Asn Glu Glu Glu Glu Gln Thr Leu Thr Ile Leu
1430 1435 1440

Ile Arg Thr Gly Leu Leu Val Ile Ser Gly Leu Phe Pro Val Ser
1445 1450 1455

Ile Pro Ile Thr Ala Ala Ala Trp Tyr Leu Trp Glu Val Lys Lys
1460 1465 1470

Gln Arg Ala Gly Val Leu Trp Asp Val Pro Ser Pro Pro Pro Met
1475 1480 1485

Gly Lys Ala Glu Leu Glu Asp Gly Ala Tyr Arg Ile Lys Gln Lys
1490 1495 1500

Gly Ile Leu Gly Tyr Ser Gln Ile Gly Ala Gly Val Tyr Lys Glu
1505 1510 1515

Gly Thr Phe His Thr Met Trp His Val Thr Arg Gly Ala Val Leu
1520 1525 1530

Met His Lys Gly Lys Arg Ile Glu Pro Ser Trp Ala Asp Val Lys
1535 1540 1545

Lys Asp Leu Ile Ser Tyr Gly Gly Gly Trp Lys Leu Glu Gly Glu
1550 1555 1560

Trp Lys Glu Gly Glu Glu Val Gln Val Leu Ala Leu Glu Pro Gly
1565 1570 1575

Lys Asn Pro Arg Ala Val Gln Thr Lys Pro Gly Leu Phe Lys Thr
1580 1585 1590

Asn Ala Gly Thr Ile Gly Ala Val Ser Leu Asp Phe Ser Pro Gly
1595 1600 1605

Thr Ser Gly Ser Pro Ile Ile Asp Lys Lys Gly Lys Val Val Gly
1610 1615 1620

Leu Tyr Gly Asn Gly Val Val Thr Arg Ser Gly Ala Tyr Val Ser

1625 1630 1635

Ala Ile Ala Gln Thr Glu Lys Ser Ile Glu Asp Asn Pro Glu Ile
1640 1645 1650

Glu Asp Asp Ile Phe Arg Lys Arg Arg Leu Thr Ile Met Asp Leu
1655 1660 1665

His Pro Gly Ala Gly Lys Thr Lys Arg Tyr Leu Pro Ala Ile Val
1670 1675 1680

Arg Glu Ala Ile Lys Arg Gly Leu Arg Thr Leu Ile Leu Ala Pro
1685 1690 1695

Thr Arg Val Val Ala Ala Glu Met Glu Glu Ala Leu Arg Gly Leu
1700 1705 1710

Pro Ile Arg Tyr Gln Thr Pro Ala Ile Arg Ala Val His Thr Gly
1715 1720 1725

Arg Glu Ile Val Asp Leu Met Cys His Ala Thr Phe Thr Met Arg
1730 1735 1740

Leu Leu Ser Pro Val Arg Val Pro Asn Tyr Asn Leu Ile Ile Met
1745 1750 1755

Asp Glu Ala His Phe Thr Asp Pro Ala Ser Ile Ala Ala Arg Gly
1760 1765 1770

Tyr Ile Ser Thr Arg Val Glu Met Gly Glu Ala Ala Gly Ile Phe
1775 1780 1785

Met Thr Ala Thr Pro Pro Gly Ser Arg Asp Pro Phe Pro Gln Ser
1790 1795 1800

Asn Ala Pro Ile Ile Asp Glu Glu Arg Glu Ile Pro Glu Arg Ser
1805 1810 1815

Trp Asn Ser Gly His Glu Trp Val Thr Asp Phe Lys Gly Lys Thr
1820 1825 1830

Val Trp Phe Val Pro Ser Ile Lys Ala Gly Asn Asp Ile Ala Ala
1835 1840 1845

Cys Leu Arg Lys Asn Gly Lys Lys Val Ile Gln Leu Ser Arg Lys
1850 1855 1860

Thr Phe Asp Ser Glu Tyr Val Lys Thr Arg Thr Asn Asp Trp Asp
1865 1870 1875

Phe Val Val Thr Thr Asp Ile Ser Glu Met Gly Ala Asn Phe Lys

1880 1885 1890

Ala Glu Arg Val Ile Asp Pro Arg Arg Cys Met Lys Pro Val Ile
1895 1900 1905

Leu Thr Asp Gly Glu Glu Arg Val Ile Leu Ala Gly Pro Met Pro
1910 1915 1920

Val Thr His Ser Ser Ala Ala Gln Arg Arg Gly Arg Ile Gly Arg
1925 1930 1935

Asn Pro Lys Asn Glu Asn Asp Gln Tyr Ile Tyr Met Gly Glu Pro
1940 1945 1950

Leu Glu Asn Asp Glu Asp Cys Ala His Trp Lys Glu Ala Lys Met
1955 1960 1965

Leu Leu Asp Asn Ile Asn Thr Pro Glu Gly Ile Ile Pro Ser Met
1970 1975 1980

Phe Glu Pro Glu Arg Glu Lys Val Asp Ala Ile Asp Gly Glu Tyr
1985 1990 1995

Arg Leu Arg Gly Glu Ala Arg Lys Thr Phe Val Asp Leu Met Arg
2000 2005 2010

Arg Gly Asp Leu Pro Val Trp Leu Ala Tyr Arg Val Ala Ala Glu
2015 2020 2025

Gly Ile Asn Tyr Ala Asp Arg Arg Trp Cys Phe Asp Gly Val Lys
2030 2035 2040

Asn Asn Gln Ile Leu Glu Glu Asn Val Glu Val Glu Ile Trp Thr
2045 2050 2055

Lys Glu Gly Glu Arg Lys Lys Leu Lys Pro Arg Trp Leu Asp Ala
2060 2065 2070

Arg Ile Tyr Ser Asp Pro Leu Ala Leu Lys Glu Phe Lys Glu Phe
2075 2080 2085

Ala Ala Gly Arg Lys Ser Leu Thr Leu Asn Leu Ile Thr Glu Met
2090 2095 2100

Gly Arg Leu Pro Thr Phe Met Thr Gln Lys Val Arg Asp Ala Leu
2105 2110 2115

Asp Asn Leu Ala Val Leu His Thr Ala Glu Ala Gly Gly Arg Ala
2120 2125 2130

Tyr Asn His Ala Leu Ser Glu Leu Pro Glu Thr Leu Glu Thr Leu

2135 2140 2145

Leu Leu Leu Thr Leu Leu Ala Thr Val Thr Gly Gly Ile Phe Leu
2150 2155 2160

Phe Leu Met Ser Ala Arg Gly Ile Gly Lys Met Thr Leu Gly Met
2165 2170 2175

Cys Cys Ile Ile Thr Ala Ser Ile Leu Leu Trp Tyr Ala Gln Ile
2180 2185 2190

Gln Pro His Trp Ile Ala Ala Ser Ile Ile Leu Glu Phe Phe Leu
2195 2200 2205

Ile Val Leu Leu Ile Pro Glu Pro Glu Lys Gln Arg Thr Pro Gln
2210 2215 2220

Asp Asn Gln Leu Thr Tyr Val Val Ile Ala Ile Leu Thr Val Val
2225 2230 2235

Ala Ala Thr Met Ala Asn Glu Met Gly Phe Leu Glu Lys Thr Lys
2240 2245 2250

Lys Asp Leu Gly Leu Gly Ser Ile Ala Thr Gln Gln Pro Glu Ser
2255 2260 2265

Asn Ile Leu Asp Ile Asp Leu Arg Pro Ala Ser Ala Trp Thr Leu
2270 2275 2280

Tyr Ala Val Ala Thr Thr Phe Val Thr Pro Met Leu Arg His Ser
2285 2290 2295

Ile Glu Asn Ser Ser Val Asn Val Ser Leu Thr Ala Ile Ala Asn
2300 2305 2310

Gln Ala Thr Val Leu Met Gly Leu Gly Lys Gly Trp Pro Leu Ser
2315 2320 2325

Lys Met Asp Ile Gly Val Pro Leu Leu Ala Ile Gly Cys Tyr Ser
2330 2335 2340

Gln Val Asn Pro Ile Thr Leu Thr Ala Ala Leu Phe Leu Leu Val
2345 2350 2355

Ala His Tyr Ala Ile Ile Gly Pro Gly Leu Gln Ala Lys Ala Thr
2360 2365 2370

Arg Glu Ala Gln Lys Arg Ala Ala Ala Gly Ile Met Lys Asn Pro
2375 2380 2385

Thr Val Asp Gly Ile Thr Val Ile Asp Leu Asp Pro Ile Pro Tyr

2390 2395 2400

Asp Pro Lys Phe Glu Lys Gln Leu Gly Gln Val Met Leu Leu Val
2405 2410 2415

Leu Cys Val Thr Gln Val Leu Met Met Arg Thr Thr Trp Ala Leu
2420 2425 2430

Cys Glu Ala Leu Thr Leu Ala Thr Gly Pro Ile Ser Thr Leu Trp
2435 2440 2445

Glu Gly Asn Pro Gly Arg Phe Trp Asn Thr Thr Ile Ala Val Ser
2450 2455 2460

Met Ala Asn Ile Phe Arg Gly Ser Tyr Leu Ala Gly Ala Gly Leu
2465 2470 2475

Leu Phe Ser Ile Met Lys Asn Thr Thr Asn Thr Arg Arg Gly Thr
2480 2485 2490

Gly Asn Ile Gly Glu Thr Leu Gly Glu Lys Trp Lys Ser Arg Leu
2495 2500 2505

Asn Ala Leu Gly Lys Ser Glu Phe Gln Ile Tyr Lys Lys Ser Gly
2510 2515 2520

Ile Gln Glu Val Asp Arg Thr Leu Ala Lys Glu Gly Ile Lys Arg
2525 2530 2535

Gly Glu Thr Asp His His Ala Val Ser Arg Gly Ser Ala Lys Leu
2540 2545 2550

Arg Trp Phe Val Glu Arg Asn Met Val Thr Pro Glu Gly Lys Val
2555 2560 2565

Val Asp Leu Gly Cys Gly Arg Gly Gly Trp Ser Tyr Tyr Cys Gly
2570 2575 2580

Gly Leu Lys Asn Val Arg Glu Val Lys Gly Leu Thr Lys Gly Gly
2585 2590 2595

Pro Gly His Glu Glu Pro Ile Pro Met Ser Thr Tyr Gly Trp Asn
2600 2605 2610

Leu Val Arg Leu Gln Ser Gly Val Asp Val Phe Phe Ile Pro Pro
2615 2620 2625

Glu Lys Cys Asp Thr Leu Leu Cys Asp Ile Gly Glu Ser Ser Pro
2630 2635 2640

Asn Pro Thr Val Glu Ala Gly Arg Thr Leu Arg Val Leu Asn Leu

2645 2650 2655

Val Glu Asn Trp Leu Asn Asn Asn Thr Gln Phe Cys Ile Lys Val
2660 2665 2670

Leu Asn Pro Tyr Met Pro Ser Val Ile Glu Lys Met Glu Ala Leu
2675 2680 2685

Gln Arg Lys Tyr Gly Gly Ala Leu Val Arg Asn Pro Leu Ser Arg
2690 2695 2700

Asn Ser Thr His Glu Met Tyr Trp Val Ser Asn Ala Ser Gly Asn
2705 2710 2715

Ile Val Ser Ser Val Asn Met Ile Ser Arg Met Leu Ile Asn Arg
2720 2725 2730

Phe Thr Met Arg Tyr Lys Lys Ala Thr Tyr Glu Pro Asp Val Asp
2735 2740 2745

Leu Gly Ser Gly Thr Arg Asn Ile Gly Ile Glu Ser Glu Ile Pro
2750 2755 2760

Asn Leu Asp Ile Ile Gly Lys Arg Ile Glu Lys Ile Lys Gln Glu
2765 2770 2775

His Glu Thr Ser Trp His Tyr Asp Gln Asp His Pro Tyr Lys Thr
2780 2785 2790

Trp Ala Tyr His Gly Ser Tyr Glu Thr Lys Gln Thr Gly Ser Ala
2795 2800 2805

Ser Ser Met Val Asn Gly Val Val Arg Leu Leu Thr Lys Pro Trp
2810 2815 2820

Asp Val Val Pro Met Val Thr Gln Met Ala Met Thr Asp Thr Thr
2825 2830 2835

Pro Phe Gly Gln Gln Arg Val Phe Lys Glu Lys Val Asp Thr Arg
2840 2845 2850

Thr Gln Glu Pro Lys Glu Gly Thr Lys Lys Leu Met Lys Ile Thr
2855 2860 2865

Ala Glu Trp Leu Trp Lys Glu Leu Gly Lys Lys Lys Thr Pro Arg
2870 2875 2880

Met Cys Thr Arg Glu Glu Phe Thr Arg Lys Val Arg Ser Asn Ala
2885 2890 2895

Ala Leu Gly Ala Ile Phe Thr Asp Glu Asn Lys Trp Lys Ser Ala

2900 2905 2910

Arg Glu Ala Val Glu Asp Ser Arg Phe Trp Glu Leu Val Asp Lys
2915 2920 2925

Glu Arg Asn Leu His Leu Glu Gly Lys Cys Glu Thr Cys Val Tyr
2930 2935 2940

Asn Met Met Gly Lys Arg Glu Lys Lys Leu Gly Glu Phe Gly Lys
2945 2950 2955

Ala Lys Gly Ser Arg Ala Ile Trp Tyr Met Trp Leu Gly Ala Arg
2960 2965 2970

Phe Leu Glu Phe Glu Ala Leu Gly Phe Leu Asn Glu Asp His Trp
2975 2980 2985

Phe Ser Arg Glu Asn Ser Leu Ser Gly Val Glu Gly Glu Gly Leu
2990 2995 3000

His Lys Leu Gly Tyr Ile Leu Arg Asp Val Ser Lys Lys Glu Gly
3005 3010 3015

Gly Ala Met Tyr Ala Asp Asp Thr Ala Gly Trp Asp Thr Arg Ile
3020 3025 3030

Thr Leu Glu Asp Leu Lys Asn Glu Glu Met Val Thr Asn His Met
3035 3040 3045

Glu Gly Glu His Lys Lys Leu Ala Glu Ala Ile Phe Lys Leu Thr
3050 3055 3060

Tyr Gln Asn Lys Val Val Arg Val Gln Arg Pro Thr Pro Arg Gly
3065 3070 3075

Thr Val Met Asp Ile Ile Ser Arg Arg Asp Gln Arg Gly Ser Gly
3080 3085 3090

Gln Val Gly Thr Tyr Gly Leu Asn Thr Phe Thr Asn Met Glu Ala
3095 3100 3105

Gln Leu Ile Arg Gln Met Glu Gly Glu Gly Val Phe Lys Ser Ile
3110 3115 3120

Gln His Leu Thr Ile Thr Glu Glu Ile Ala Val Gln Asn Trp Leu
3125 3130 3135

Ala Arg Val Gly Arg Glu Arg Leu Ser Arg Met Ala Ile Ser Gly
3140 3145 3150

Asp Asp Cys Val Val Lys Pro Leu Asp Asp Arg Phe Ala Ser Ala

3155 3160 3165

Leu Thr Ala Leu Asn Asp Met Gly Lys Ile Arg Lys Asp Ile Gln
3170 3175 3180

Gln Trp Glu Pro Ser Arg Gly Trp Asn Asp Trp Thr Gln Val Pro
3185 3190 3195

Phe Cys Ser His His Phe His Glu Leu Ile Met Lys Asp Gly Arg
3200 3205 3210

Val Leu Val Val Pro Cys Arg Asn Gln Asp Glu Leu Ile Gly Arg
3215 3220 3225

Ala Arg Ile Ser Gln Gly Ala Gly Trp Ser Leu Arg Glu Thr Ala
3230 3235 3240

Cys Leu Gly Lys Ser Tyr Ala Gln Met Trp Ser Leu Met Tyr Phe
3245 3250 3255

His Arg Arg Asp Leu Arg Leu Ala Ala Asn Ala Ile Cys Ser Ala
3260 3265 3270

Val Pro Ser His Trp Val Pro Thr Ser Arg Thr Thr Trp Ser Ile
3275 3280 3285

His Ala Lys His Glu Trp Met Thr Thr Glu Asp Met Leu Thr Val
3290 3295 3300

Trp Asn Arg Val Trp Ile Gln Glu Asn Pro Trp Met Glu Asp Lys
3305 3310 3315

Thr Pro Val Glu Ser Trp Glu Glu Ile Pro Tyr Leu Gly Lys Arg
3320 3325 3330

Glu Asp Gln Trp Cys Gly Ser Leu Ile Gly Leu Thr Ser Arg Ala
3335 3340 3345

Thr Trp Ala Lys Asn Ile Gln Ala Ala Ile Asn Gln Val Arg Ser
3350 3355 3360

Leu Ile Gly Asn Glu Glu Tyr Thr Asp Tyr Met Pro Ser Met Lys
3365 3370 3375

Arg Phe Arg Arg Glu Glu Glu Glu Ala Gly Val Leu Trp
3380 3385 3390