

50189A_SeqListing
SEQUENCE LISTING

<110> Immune Design Corporation
<120> COMPOSITIONS COMPRISING LENTIVIRAL VECTORS EXPRESSING IL-12 AND
METHODS OF USE THEREOF
<130> 31943/50189A/PC
<160> 28
<170> PatentIn version 3.5
<210> 1
<211> 423
<212> PRT
<213> Sindbis virus

<220>
<221> misc_feature
<222> (160)..(160)
<223> Xaa can be any naturally occurring amino acid

<400> 1

Ser	Val	Ile	Asp	Asp	Phe	Thr	Leu	Thr	Ser	Pro	Tyr	Leu	Gly	Thr	Cys
1				5					10					15	

Ser	Tyr	Cys	His	His	Thr	Val	Pro	Cys	Phe	Ser	Pro	Val	Lys	Ile	Glu
			20					25					30		

Gln	Val	Trp	Asp	Glu	Ala	Asp	Asp	Asn	Thr	Ile	Arg	Ile	Gln	Thr	Ser
		35					40					45			

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

Arg	Tyr	Met	Ser	Leu	Lys	Gln	Asp	His	Thr	Val	Lys	Glu	Gly	Thr	Met
65					70					75					80

Asp	Asp	Ile	Lys	Ile	Ser	Thr	Ser	Gly	Pro	Cys	Arg	Arg	Leu	Ser	Tyr
			85						90					95	

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Lys Gly Tyr Phe Leu Leu Ala Lys Cys Pro Pro Gly Asp Ser Val Thr
100 105 110

Val Ser Ile Val Ser Ser Asn Ser Ala Thr Ser Cys Thr Leu Ala Arg
115 120 125

Lys Ile Lys Pro Lys Phe Val Gly Arg Glu Lys Tyr Asp Leu Pro Pro
130 135 140

Val His Gly Lys Lys Ile Pro Cys Thr Val Tyr Asp Arg Leu Lys Xaa
145 150 155 160

Thr Thr Ala Gly Tyr Ile Thr Met His Arg Pro Arg Pro His Ala Tyr
165 170 175

Thr Ser Tyr Leu Glu Glu Ser Ser Gly Lys Val Tyr Ala Lys Pro Pro
180 185 190

Ser Gly Lys Asn Ile Thr Tyr Glu Cys Lys Cys Gly Asp Tyr Lys Thr
195 200 205

Gly Thr Val Ser Thr Arg Thr Glu Ile Thr Gly Cys Thr Ala Ile Lys
210 215 220

Gln Cys Val Ala Tyr Lys Ser Asp Gln Thr Lys Trp Val Phe Asn Ser
225 230 235 240

Pro Asp Leu Ile Arg His Asp Asp His Thr Ala Gln Gly Lys Leu His
245 250 255

Leu Pro Phe Lys Leu Ile Pro Ser Thr Cys Met Val Pro Val Ala His
260 265 270

Ala Pro Asn Val Ile His Gly Phe Lys His Ile Ser Leu Gln Leu Asp
275 280 285

50189A_SeqListing

Thr Asp His Leu Thr Leu Leu Thr Thr Arg Arg Leu Gly Ala Asn Pro
 290 295 300

Glu Pro Thr Thr Glu Trp Ile Val Gly Lys Thr Val Arg Asn Phe Thr
 305 310 315 320

Val Asp Arg Asp Gly Leu Glu Tyr Ile Trp Gly Asn His Glu Pro Val
 325 330 335

Arg Val Tyr Ala Gln Glu Ser Ala Pro Gly Asp Pro His Gly Trp Pro
 340 345 350

His Glu Ile Val Gln His Tyr Tyr His Arg His Pro Val Tyr Thr Ile
 355 360 365

Leu Ala Val Ala Ser Ala Thr Val Ala Met Met Ile Gly Val Thr Val
 370 375 380

Ala Val Leu Cys Ala Cys Lys Ala Arg Arg Glu Cys Leu Thr Pro Tyr
 385 390 395 400

Ala Leu Ala Pro Asn Ala Val Ile Pro Thr Ser Leu Ala Leu Leu Cys
 405 410 415

Cys Val Arg Ser Ala Asn Ala
 420

<210> 2
 <211> 986
 <212> PRT
 <213> Sindbis virus

<400> 2

Met Ser Ala Ala Pro Leu Val Thr Ala Met Cys Leu Leu Gly Asn Val
 1 5 10 15

50189A_SeqListing

Ser Phe Pro Cys Asp Arg Pro Pro Thr Cys Tyr Thr Arg Glu Pro Ser
20 25 30

Arg Ala Leu Asp Ile Leu Glu Glu Asn Val Asn His Glu Ala Tyr Asp
35 40 45

Thr Leu Leu Asn Ala Ile Leu Arg Cys Gly Ser Ser Gly Ser Val Ile
50 55 60

Asp Asp Phe Thr Leu Thr Ser Pro Tyr Leu Gly Thr Cys Ser Tyr Cys
65 70 75 80

His His Thr Val Pro Cys Phe Ser Pro Val Lys Ile Glu Gln Val Trp
85 90 95

Asp Glu Ala Asp Asp Asn Thr Ile Arg Ile Gln Thr Ser Ala Gln Phe
100 105 110

Gly Tyr Asp Gln Ser Gly Ala Ala Ser Ala Asn Lys Tyr Arg Tyr Met
115 120 125

Ser Leu Lys Gln Met Tyr Pro Tyr Asp Val Pro Asp Tyr Ala Thr Val
130 135 140

Lys Glu Gly Thr Met Asp Asp Ile Lys Ile Ser Thr Ser Gly Pro Cys
145 150 155 160

Arg Arg Leu Ser Tyr Lys Gly Tyr Phe Leu Leu Ala Lys Cys Pro Pro
165 170 175

Gly Asp Ser Val Thr Val Ser Ile Val Ser Ser Asn Ser Ala Thr Ser
180 185 190

Cys Thr Leu Ala Arg Lys Ile Lys Pro Lys Phe Val Gly Arg Glu Lys
195 200 205

50189A_SeqListing

Tyr Asp Leu Pro Pro Val His Gly Lys Lys Ile Pro Cys Thr Val Tyr
210 215 220

Asp Arg Leu Ala Ala Thr Thr Ala Gly Tyr Ile Thr Met His Arg Pro
225 230 235 240

Arg Pro His Ala Tyr Thr Ser Tyr Leu Glu Glu Ser Ser Gly Lys Val
245 250 255

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

Gly Asp Tyr Lys Thr Gly Thr Val Ser Thr Arg Thr Glu Ile Thr Gly
275 280 285

Cys Thr Ala Ile Lys Gln Cys Val Ala Tyr Lys Ser Asp Gln Thr Lys
290 295 300

Trp Val Phe Asn Ser Pro Asp Leu Ile Arg His Asp Asp His Thr Ala
305 310 315 320

Gln Gly Lys Leu His Leu Pro Phe Lys Leu Ile Pro Ser Thr Cys Met
325 330 335

Val Pro Val Ala His Ala Pro Asn Val Ile His Gly Phe Lys His Ile
340 345 350

Ser Leu Gln Leu Asp Thr Asp His Leu Thr Leu Leu Thr Thr Arg Arg
355 360 365

Leu Gly Ala Asn Pro Glu Pro Thr Thr Glu Trp Ile Val Gly Lys Thr
370 375 380

Val Arg Asn Phe Thr Val Asp Arg Asp Gly Leu Glu Tyr Ile Trp Gly
385 390 395 400

50189A_SeqListing

Asn His Glu Pro Val Arg Val Tyr Ala Gln Glu Ser Ala Pro Gly Asp
405 410 415

Pro His Gly Trp Pro His Glu Ile Val Gln His Tyr Tyr His Arg His
420 425 430

Pro Val Tyr Thr Ile Leu Ala Val Ala Ser Ala Thr Val Ala Met Met
435 440 445

Ile Gly Val Thr Val Ala Val Leu Cys Ala Cys Lys Ala Arg Arg Glu
450 455 460

Cys Leu Thr Pro Tyr Ala Leu Ala Pro Asn Ala Val Ile Pro Thr Ser
465 470 475 480

Leu Ala Leu Leu Cys Cys Val Arg Ser Ala Asn Ala Glu Thr Phe Thr
485 490 495

Glu Thr Met Ser Tyr Leu Trp Ser Asn Ser Gln Pro Phe Phe Trp Val
500 505 510

Gln Leu Cys Ile Pro Leu Ala Ala Phe Ile Val Leu Met Arg Cys Cys
515 520 525

Ser Cys Cys Leu Pro Phe Leu Val Val Ala Gly Ala Tyr Leu Ala Lys
530 535 540

Val Asp Ala Tyr Glu His Ala Thr Thr Val Pro Asn Val Pro Gln Ile
545 550 555 560

Pro Tyr Lys Ala Leu Val Glu Arg Ala Gly Tyr Ala Pro Leu Asn Leu
565 570 575

Glu Ile Thr Val Met Ser Ser Glu Val Leu Pro Ser Thr Asn Gln Glu
580 585 590

50189A_SeqListing

Tyr Ile Thr Cys Lys Phe Thr Thr Val Val Pro Ser Pro Lys Ile Lys
595 600 605

Cys Cys Gly Ser Leu Glu Cys Gln Pro Ala Ala His Ala Gly Tyr Thr
610 615 620

Cys Lys Val Phe Gly Gly Val Tyr Pro Phe Met Trp Gly Gly Ala Gln
625 630 635 640

Cys Phe Cys Asp Ser Glu Asn Ser Gln Met Ser Glu Ala Tyr Val Glu
645 650 655

Leu Ser Ala Asp Cys Ala Ser Asp His Ala Gln Ala Ile Lys Val His
660 665 670

Thr Ala Ala Met Lys Val Gly Leu Arg Ile Val Tyr Gly Asn Thr Thr
675 680 685

Ser Phe Leu Asp Val Tyr Val Asn Gly Val Thr Pro Gly Thr Ser Lys
690 695 700

Asp Leu Lys Val Ile Ala Gly Pro Ile Ser Ala Ser Phe Thr Pro Phe
705 710 715 720

Asp His Lys Val Val Ile His Arg Gly Leu Val Tyr Asn Tyr Asp Phe
725 730 735

Pro Glu Tyr Gly Ala Met Lys Pro Gly Ala Phe Gly Asp Ile Gln Ala
740 745 750

Thr Ser Leu Thr Ser Lys Asp Leu Ile Ala Ser Thr Asp Ile Arg Leu
755 760 765

Leu Lys Pro Ser Ala Lys Asn Val His Val Pro Tyr Thr Gln Ala Ser
770 775 780

50189A_SeqListing

Ser Gly Phe Glu Met Trp Lys Asn Asn Ser Gly Arg Pro Leu Gln Glu
785 790 795 800

Thr Ala Pro Phe Gly Cys Lys Ile Ala Val Asn Pro Leu Arg Ala Val
805 810 815

Asp Cys Ser Tyr Gly Asn Ile Pro Ile Ser Ile Asp Ile Pro Asn Ala
820 825 830

Ala Phe Ile Arg Thr Ser Asp Ala Pro Leu Val Ser Thr Val Lys Cys
835 840 845

Glu Val Ser Glu Cys Thr Tyr Ser Ala Asp Phe Gly Gly Met Ala Thr
850 855 860

Leu Gln Tyr Val Ser Asp Arg Glu Gly Gln Cys Pro Val His Ser His
865 870 875 880

Ser Ser Thr Ala Thr Leu Gln Glu Ser Thr Val His Val Leu Glu Lys
885 890 895

Gly Ala Val Thr Val His Phe Ser Thr Ala Ser Pro Gln Ala Asn Phe
900 905 910

Ile Val Ser Leu Cys Gly Lys Lys Thr Thr Cys Asn Ala Glu Cys Lys
915 920 925

Pro Pro Ala Asp His Ile Val Ser Thr Pro His Lys Asn Asp Gln Glu
930 935 940

Phe Gln Ala Ala Ile Ser Lys Thr Ser Trp Ser Trp Leu Phe Ala Leu
945 950 955 960

Phe Gly Gly Ala Ser Ser Leu Leu Ile Ile Gly Leu Met Ile Phe Ala
965 970 975

50189A_SeqListing

Cys Ser Met Met Leu Thr Ser Thr Arg Arg
980 985

<210> 3
<211> 982
<212> PRT
<213> Sindbis virus

<400> 3

Met Ser Ala Ala Pro Leu Val Thr Ala Met Cys Leu Leu Gly Asn Val
1 5 10 15

Ser Phe Pro Cys Asp Arg Pro Pro Thr Cys Tyr Thr Arg Glu Pro Ser
20 25 30

Arg Ala Leu Asp Ile Leu Glu Glu Asn Val Asn His Glu Ala Tyr Asp
35 40 45

Thr Leu Leu Asn Ala Ile Leu Arg Cys Gly Ser Ser Gly Arg Ser Lys
50 55 60

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

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

Glu Gln Val Trp Asp Glu Ala Asp Asp Asn Thr Ile Arg Ile Gln Thr
100 105 110

Ser Ala Gln Phe Gly Tyr Asp Gln Ser Gly Ala Ala Ser Ala Asn Lys
115 120 125

Tyr Arg Tyr Met Ser Leu Glu Gln Asp His Thr Val Lys Glu Gly Thr
130 135 140

50189A_SeqListing

Met Asp Asp Ile Lys Ile Ser Thr Ser Gly Pro Cys Arg Arg Leu Ser
145 150 155 160

Tyr Lys Gly Tyr Phe Leu Leu Ala Lys Cys Pro Pro Gly Asp Ser Val
165 170 175

Thr Val Ser Ile Val Ser Ser Asn Ser Ala Thr Ser Cys Thr Leu Ala
180 185 190

Arg Lys Ile Lys Pro Lys Phe Val Gly Arg Glu Lys Tyr Asp Leu Pro
195 200 205

Pro Val His Gly Lys Lys Ile Pro Cys Thr Val Tyr Asp Arg Leu Glu
210 215 220

Gly Thr Thr Ala Gly Tyr Ile Thr Met His Arg Pro Arg Pro His Ala
225 230 235 240

Tyr Thr Ser Tyr Leu Glu Glu Ser Ser Gly Lys Val Tyr Ala Lys Pro
245 250 255

Pro Ser Gly Lys Asn Ile Thr Tyr Glu Cys Lys Cys Gly Asp Tyr Lys
260 265 270

Thr Gly Thr Val Ser Thr Arg Thr Glu Ile Thr Gly Cys Thr Ala Ile
275 280 285

Lys Gln Cys Val Ala Tyr Lys Ser Asp Gln Thr Lys Trp Val Phe Asn
290 295 300

Ser Pro Asp Leu Ile Arg His Asp Asp His Thr Ala Gln Gly Lys Leu
305 310 315 320

His Leu Pro Phe Lys Leu Ile Pro Ser Thr Cys Met Val Pro Val Ala
325 330 335

50189A_SeqListing

His Ala Pro Asn Val Ile His Gly Phe Lys His Ile Ser Leu Gln Leu
 340 345 350

Asp Thr Asp His Leu Thr Leu Leu Thr Thr Arg Arg Leu Gly Ala Asn
 355 360 365

Pro Glu Pro Thr Thr Glu Trp Ile Val Gly Lys Thr Val Arg Asn Phe
 370 375 380

Thr Val Asp Arg Asp Gly Leu Glu Tyr Ile Trp Gly Asn His Glu Pro
 385 390 395 400

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

Pro His Glu Ile Val Gln His Tyr Tyr His Arg His Pro Val Tyr Thr
 420 425 430

Ile Leu Ala Val Ala Ser Ala Thr Val Ala Met Met Ile Gly Val Thr
 435 440 445

Val Ala Val Leu Cys Ala Cys Lys Ala Arg Arg Glu Cys Leu Thr Pro
 450 455 460

Tyr Ala Leu Ala Pro Asn Ala Val Ile Pro Thr Ser Leu Ala Leu Leu
 465 470 475 480

Cys Cys Val Arg Ser Ala Asn Ala Glu Thr Phe Thr Glu Thr Met Ser
 485 490 495

Tyr Leu Trp Ser Asn Ser Gln Pro Phe Phe Trp Val Gln Leu Cys Ile
 500 505 510

Pro Leu Ala Ala Phe Ile Val Leu Met Arg Cys Cys Ser Cys Cys Leu
 515 520 525

50189A_SeqListing

Pro Phe Leu Val Val Ala Gly Ala Tyr Leu Ala Lys Val Asp Ala Tyr
530 535 540

Glu His Ala Thr Thr Val Pro Asn Val Pro Gln Ile Pro Tyr Lys Ala
545 550 555 560

Leu Val Glu Arg Ala Gly Tyr Ala Pro Leu Asn Leu Glu Ile Thr Val
565 570 575

Met Ser Ser Glu Val Leu Pro Ser Thr Asn Gln Glu Tyr Ile Thr Cys
580 585 590

Lys Phe Thr Thr Val Val Pro Ser Pro Lys Ile Lys Cys Cys Gly Ser
595 600 605

Leu Glu Cys Gln Pro Ala Ala His Ala Asp Tyr Thr Cys Lys Val Phe
610 615 620

Gly Gly Val Tyr Pro Phe Met Trp Gly Gly Ala Gln Cys Phe Cys Asp
625 630 635 640

Ser Glu Asn Ser Gln Met Ser Glu Ala Tyr Val Glu Leu Ser Ala Asp
645 650 655

Cys Ala Ser Asp His Ala Gln Ala Ile Lys Val His Thr Ala Ala Met
660 665 670

Lys Val Gly Leu Arg Ile Val Tyr Gly Asn Thr Thr Ser Phe Leu Asp
675 680 685

Val Tyr Val Asn Gly Val Thr Pro Gly Thr Ser Lys Asp Leu Lys Val
690 695 700

Ile Ala Gly Pro Ile Ser Ala Ser Phe Thr Pro Phe Asp His Lys Val
705 710 715 720

50189A_SeqListing

Val Ile His Arg Gly Leu Val Tyr Asn Tyr Asp Phe Pro Glu Tyr Gly
725 730 735

Ala Met Lys Pro Gly Ala Phe Gly Asp Ile Gln Ala Thr Ser Leu Thr
740 745 750

Ser Lys Asp Leu Ile Ala Ser Thr Asp Ile Arg Leu Leu Lys Pro Ser
755 760 765

Ala Lys Asn Val His Val Pro Tyr Thr Gln Ala Ser Ser Gly Phe Glu
770 775 780

Met Trp Lys Asn Asn Ser Gly Arg Pro Leu Gln Glu Thr Ala Pro Phe
785 790 795 800

Gly Cys Lys Ile Ala Val Asn Pro Leu Arg Ala Val Asp Cys Ser Tyr
805 810 815

Gly Asn Ile Pro Ile Ser Ile Asp Ile Pro Asn Ala Ala Phe Ile Arg
820 825 830

Thr Ser Asp Ala Pro Leu Val Ser Thr Val Lys Cys Glu Val Ser Glu
835 840 845

Cys Thr Tyr Ser Ala Asp Phe Gly Gly Met Ala Thr Leu Gln Tyr Val
850 855 860

Ser Asp Arg Glu Gly Gln Cys Pro Val His Ser His Ser Ser Thr Ala
865 870 875 880

Thr Leu Gln Glu Ser Thr Val His Val Leu Glu Lys Gly Ala Val Thr
885 890 895

Val His Phe Ser Thr Ala Ser Pro Gln Ala Asn Phe Ile Val Ser Leu
900 905 910

50189A_SeqListing

Cys Gly Lys Lys Thr Thr Cys Asn Ala Glu Cys Lys Pro Pro Ala Asp
 915 920 925

His Ile Val Ser Thr Pro His Lys Asn Asp Gln Glu Phe Gln Ala Ala
 930 935 940

Ile Ser Lys Thr Ser Trp Ser Trp Leu Phe Ala Leu Phe Gly Gly Ala
 945 950 955 960

Ser Ser Leu Leu Ile Ile Gly Leu Met Ile Phe Ala Cys Ser Met Met
 965 970 975

Leu Thr Ser Thr Arg Arg
 980

<210> 4
 <211> 982
 <212> PRT
 <213> Sindbis virus

<400> 4

Met Ser Ala Ala Pro Leu Val Thr Ala Met Cys Leu Leu Gly Asn Val
 1 5 10 15

Ser Phe Pro Cys Asp Arg Pro Pro Thr Cys Tyr Thr Arg Glu Pro Ser
 20 25 30

Arg Ala Leu Asp Ile Leu Glu Glu Asn Val Asn His Glu Ala Tyr Asp
 35 40 45

Thr Leu Leu Asn Ala Ile Leu Arg Cys Gly Ser Ser Gly Arg Ser Lys
 50 55 60

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

Cys Ser Tyr Cys His His Thr Val Pro Cys Phe Ser Pro Val Lys Ile

50189A_SeqListing

85

90

95

Glu Gln Val Trp Asp Glu Ala Asp Asp Asn Thr Ile Arg Ile Gln Thr
 100 105 110

Ser Ala Gln Phe Gly Tyr Asp Gln Ser Gly Ala Ala Ser Ala Asn Lys
 115 120 125

Tyr Arg Tyr Met Ser Leu Glu Gln Asp His Thr Val Glu Glu Gly Thr
 130 135 140

Met Asp Asp Ile Lys Ile Ser Thr Ser Gly Pro Cys Arg Arg Leu Ser
 145 150 155 160

Tyr Lys Gly Tyr Phe Leu Leu Ala Lys Cys Pro Pro Gly Asp Ser Val
 165 170 175

Thr Val Ser Ile Val Ser Ser Asn Ser Ala Thr Ser Cys Thr Leu Ala
 180 185 190

Arg Lys Ile Lys Pro Lys Phe Val Gly Arg Glu Lys Tyr Asp Leu Pro
 195 200 205

Pro Val His Gly Lys Lys Ile Pro Cys Thr Val Tyr Asp Arg Leu Glu
 210 215 220

Gly Thr Thr Ala Gly Tyr Ile Thr Met His Arg Pro Arg Pro His Ala
 225 230 235 240

Tyr Thr Ser Tyr Leu Glu Glu Ser Ser Gly Lys Val Tyr Ala Lys Pro
 245 250 255

Pro Ser Gly Lys Asn Ile Thr Tyr Glu Cys Lys Cys Gly Asp Tyr Lys
 260 265 270

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

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275

280

285

Lys Gln Cys Val Ala Tyr Lys Ser Asp Gln Thr Lys Trp Val Phe Asn
 290 295 300

Ser Pro Asp Leu Ile Arg His Asp Asp His Thr Ala Gln Gly Lys Leu
 305 310 315 320

His Leu Pro Phe Lys Leu Ile Pro Ser Thr Cys Met Val Pro Val Ala
 325 330 335

His Ala Pro Asn Val Ile His Gly Phe Lys His Ile Ser Leu Gln Leu
 340 345 350

Asp Thr Asp His Leu Thr Leu Leu Thr Thr Arg Arg Leu Gly Ala Asn
 355 360 365

Pro Glu Pro Thr Thr Glu Trp Ile Val Gly Lys Thr Val Arg Asn Phe
 370 375 380

Thr Val Asp Arg Asp Gly Leu Glu Tyr Ile Trp Gly Asn His Glu Pro
 385 390 395 400

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

Pro His Glu Ile Val Gln His Tyr Tyr His Arg His Pro Val Tyr Thr
 420 425 430

Ile Leu Ala Val Ala Ser Ala Thr Val Ala Met Met Ile Gly Val Thr
 435 440 445

Val Ala Val Leu Cys Ala Cys Lys Ala Arg Arg Glu Cys Leu Thr Pro
 450 455 460

Tyr Ala Leu Ala Pro Asn Ala Val Ile Pro Thr Ser Leu Ala Leu Leu

50189A_SeqListing

465		470		475		480									
Cys	Cys	Val	Arg	Ser	Ala	Asn	Ala	Glu	Thr	Phe	Thr	Glu	Thr	Met	Ser
				485					490					495	
Tyr	Leu	Trp	Ser	Asn	Ser	Gln	Pro	Phe	Phe	Trp	Val	Gln	Leu	Cys	Ile
			500					505					510		
Pro	Leu	Ala	Ala	Phe	Ile	Val	Leu	Met	Arg	Cys	Cys	Ser	Cys	Cys	Leu
		515					520					525			
Pro	Phe	Leu	Val	Val	Ala	Gly	Ala	Tyr	Leu	Ala	Lys	Val	Asp	Ala	Tyr
	530					535					540				
Glu	His	Ala	Thr	Thr	Val	Pro	Asn	Val	Pro	Gln	Ile	Pro	Tyr	Lys	Ala
545					550					555					560
Leu	Val	Glu	Arg	Ala	Gly	Tyr	Ala	Pro	Leu	Asn	Leu	Glu	Ile	Thr	Val
				565					570					575	
Met	Ser	Ser	Glu	Val	Leu	Pro	Ser	Thr	Asn	Gln	Glu	Tyr	Ile	Thr	Cys
			580					585					590		
Lys	Phe	Thr	Thr	Val	Val	Pro	Ser	Pro	Lys	Ile	Lys	Cys	Cys	Gly	Ser
		595					600					605			
Leu	Glu	Cys	Gln	Pro	Ala	Ala	His	Ala	Asp	Tyr	Thr	Cys	Lys	Val	Phe
	610					615					620				
Gly	Gly	Val	Tyr	Pro	Phe	Met	Trp	Gly	Gly	Ala	Gln	Cys	Phe	Cys	Asp
625					630					635					640
Ser	Glu	Asn	Ser	Gln	Met	Ser	Glu	Ala	Tyr	Val	Glu	Leu	Ser	Ala	Asp
				645					650					655	
Cys	Ala	Ser	Asp	His	Ala	Gln	Ala	Ile	Lys	Val	His	Thr	Ala	Ala	Met

50189A_SeqListing

660

665

670

Lys Val Gly Leu Arg Ile Val Tyr Gly Asn Thr Thr Ser Phe Leu Asp
 675 680 685

Val Tyr Val Asn Gly Val Thr Pro Gly Thr Ser Lys Asp Leu Lys Val
 690 695 700

Ile Ala Gly Pro Ile Ser Ala Ser Phe Thr Pro Phe Asp His Lys Val
 705 710 715 720

Val Ile His Arg Gly Leu Val Tyr Asn Tyr Asp Phe Pro Glu Tyr Gly
 725 730 735

Ala Met Lys Pro Gly Ala Phe Gly Asp Ile Gln Ala Thr Ser Leu Thr
 740 745 750

Ser Lys Asp Leu Ile Ala Ser Thr Asp Ile Arg Leu Leu Lys Pro Ser
 755 760 765

Ala Lys Asn Val His Val Pro Tyr Thr Gln Ala Ser Ser Gly Phe Glu
 770 775 780

Met Trp Lys Asn Asn Ser Gly Arg Pro Leu Gln Glu Thr Ala Pro Phe
 785 790 795 800

Gly Cys Lys Ile Ala Val Asn Pro Leu Arg Ala Val Asp Cys Ser Tyr
 805 810 815

Gly Asn Ile Pro Ile Ser Ile Asp Ile Pro Asn Ala Ala Phe Ile Arg
 820 825 830

Thr Ser Asp Ala Pro Leu Val Ser Thr Val Lys Cys Glu Val Ser Glu
 835 840 845

Cys Thr Tyr Ser Ala Asp Phe Gly Gly Met Ala Thr Leu Gln Tyr Val

50189A_SeqListing

850

855

860

Ser Asp Arg Glu Gly Gln Cys Pro Val His Ser His Ser Ser Thr Ala
865 870 875 880

Thr Leu Gln Glu Ser Thr Val His Val Leu Glu Lys Gly Ala Val Thr
885 890 895

Val His Phe Ser Thr Ala Ser Pro Gln Ala Asn Phe Ile Val Ser Leu
900 905 910

Cys Gly Lys Lys Thr Thr Cys Asn Ala Glu Cys Lys Pro Pro Ala Asp
915 920 925

His Ile Val Ser Thr Pro His Lys Asn Asp Gln Glu Phe Gln Ala Ala
930 935 940

Ile Ser Lys Thr Ser Trp Ser Trp Leu Phe Ala Leu Phe Gly Gly Ala
945 950 955 960

Ser Ser Leu Leu Ile Ile Gly Leu Met Ile Phe Ala Cys Ser Met Met
965 970 975

Leu Thr Ser Thr Arg Arg
980

<210> 5
<211> 980
<212> PRT
<213> Sindbis virus

<400> 5

Met Ser Ala Ala Pro Leu Val Thr Ala Met Cys Leu Leu Gly Asn Val
1 5 10 15

Ser Phe Pro Cys Asp Arg Pro Pro Thr Cys Tyr Thr Arg Glu Pro Ser
20 25 30

50189A_SeqListing

Arg Ala Leu Asp Ile Leu Glu Glu Asn Val Asn His Glu Ala Tyr Asp
35 40 45

Thr Leu Leu Asn Ala Ile Leu Arg Cys Gly Ser Ser Gly Arg Ser Lys
50 55 60

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

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

Glu Gln Val Trp Asp Glu Ala Asp Asp Asn Thr Ile Arg Ile Gln Thr
100 105 110

Ser Ala Gln Phe Gly Tyr Asp Gln Ser Gly Ala Ala Ser Ala Asn Lys
115 120 125

Tyr Arg Tyr Met Ser Leu Glu Gln Asp His Thr Val Lys Glu Gly Thr
130 135 140

Met Asp Asp Ile Lys Ile Ser Thr Ser Gly Pro Cys Arg Arg Leu Ser
145 150 155 160

Tyr Lys Gly Tyr Phe Leu Leu Ala Lys Cys Pro Pro Gly Asp Ser Val
165 170 175

Thr Val Ser Ile Val Ser Ser Asn Ser Ala Thr Ser Cys Thr Leu Ala
180 185 190

Arg Lys Ile Lys Pro Lys Phe Val Gly Arg Glu Lys Tyr Asp Leu Pro
195 200 205

Pro Val His Gly Lys Lys Ile Pro Cys Thr Val Tyr Asp Arg Leu Thr
210 215 220

50189A_SeqListing

Thr Ala Gly Tyr Ile Thr Met His Arg Pro Arg Pro His Ala Tyr Thr
225 230 235 240

Ser Tyr Leu Glu Glu Ser Ser Gly Lys Val Tyr Ala Lys Pro Pro Ser
245 250 255

Gly Lys Asn Ile Thr Tyr Glu Cys Lys Cys Gly Asp Tyr Lys Thr Gly
260 265 270

Thr Val Ser Thr Arg Thr Glu Ile Thr Gly Cys Thr Ala Ile Lys Gln
275 280 285

Cys Val Ala Tyr Lys Ser Asp Gln Thr Lys Trp Val Phe Asn Ser Pro
290 295 300

Asp Leu Ile Arg His Asp Asp His Thr Ala Gln Gly Lys Leu His Leu
305 310 315 320

Pro Phe Lys Leu Ile Pro Ser Thr Cys Met Val Pro Val Ala His Ala
325 330 335

Pro Asn Val Ile His Gly Phe Lys His Ile Ser Leu Gln Leu Asp Thr
340 345 350

Asp His Leu Thr Leu Leu Thr Thr Arg Arg Leu Gly Ala Asn Pro Glu
355 360 365

Pro Thr Thr Glu Trp Ile Val Gly Lys Thr Val Arg Asn Phe Thr Val
370 375 380

Asp Arg Asp Gly Leu Glu Tyr Ile Trp Gly Asn His Glu Pro Val Arg
385 390 395 400

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

50189A_SeqListing

Glu Ile Val Gln His Tyr Tyr His Arg His Pro Val Tyr Thr Ile Leu
420 425 430

Ala Val Ala Ser Ala Thr Val Ala Met Met Ile Gly Val Thr Val Ala
435 440 445

Val Leu Cys Ala Cys Lys Ala Arg Arg Glu Cys Leu Thr Pro Tyr Ala
450 455 460

Leu Ala Pro Asn Ala Val Ile Pro Thr Ser Leu Ala Leu Leu Cys Cys
465 470 475 480

Val Arg Ser Ala Asn Ala Glu Thr Phe Thr Glu Thr Met Ser Tyr Leu
485 490 495

Trp Ser Asn Ser Gln Pro Phe Phe Trp Val Gln Leu Cys Ile Pro Leu
500 505 510

Ala Ala Phe Ile Val Leu Met Arg Cys Cys Ser Cys Cys Leu Pro Phe
515 520 525

Leu Val Val Ala Gly Ala Tyr Leu Ala Lys Val Asp Ala Tyr Glu His
530 535 540

Ala Thr Thr Val Pro Asn Val Pro Gln Ile Pro Tyr Lys Ala Leu Val
545 550 555 560

Glu Arg Ala Gly Tyr Ala Pro Leu Asn Leu Glu Ile Thr Val Met Ser
565 570 575

Ser Glu Val Leu Pro Ser Thr Asn Gln Glu Tyr Ile Thr Cys Lys Phe
580 585 590

Thr Thr Val Val Pro Ser Pro Lys Ile Lys Cys Cys Gly Ser Leu Glu
595 600 605

50189A_SeqListing

Cys Gln Pro Ala Ala His Ala Asp Tyr Thr Cys Lys Val Phe Gly Gly
610 615 620

Val Tyr Pro Phe Met Trp Gly Gly Ala Gln Cys Phe Cys Asp Ser Glu
625 630 635 640

Asn Ser Gln Met Ser Glu Ala Tyr Val Glu Leu Ser Ala Asp Cys Ala
645 650 655

Ser Asp His Ala Gln Ala Ile Lys Val His Thr Ala Ala Met Lys Val
660 665 670

Gly Leu Arg Ile Val Tyr Gly Asn Thr Thr Ser Phe Leu Asp Val Tyr
675 680 685

Val Asn Gly Val Thr Pro Gly Thr Ser Lys Asp Leu Lys Val Ile Ala
690 695 700

Gly Pro Ile Ser Ala Ser Phe Thr Pro Phe Asp His Lys Val Val Ile
705 710 715 720

His Arg Gly Leu Val Tyr Asn Tyr Asp Phe Pro Glu Tyr Gly Ala Met
725 730 735

Lys Pro Gly Ala Phe Gly Asp Ile Gln Ala Thr Ser Leu Thr Ser Lys
740 745 750

Asp Leu Ile Ala Ser Thr Asp Ile Arg Leu Leu Lys Pro Ser Ala Lys
755 760 765

Asn Val His Val Pro Tyr Thr Gln Ala Ser Ser Gly Phe Glu Met Trp
770 775 780

Lys Asn Asn Ser Gly Arg Pro Leu Gln Glu Thr Ala Pro Phe Gly Cys
785 790 795 800

50189A_SeqListing

Lys Ile Ala Val Asn Pro Leu Arg Ala Val Asp Cys Ser Tyr Gly Asn
805 810 815

Ile Pro Ile Ser Ile Asp Ile Pro Asn Ala Ala Phe Ile Arg Thr Ser
820 825 830

Asp Ala Pro Leu Val Ser Thr Val Lys Cys Glu Val Ser Glu Cys Thr
835 840 845

Tyr Ser Ala Asp Phe Gly Gly Met Ala Thr Leu Gln Tyr Val Ser Asp
850 855 860

Arg Glu Gly Gln Cys Pro Val His Ser His Ser Ser Thr Ala Thr Leu
865 870 875 880

Gln Glu Ser Thr Val His Val Leu Glu Lys Gly Ala Val Thr Val His
885 890 895

Phe Ser Thr Ala Ser Pro Gln Ala Asn Phe Ile Val Ser Leu Cys Gly
900 905 910

Lys Lys Thr Thr Cys Asn Ala Glu Cys Lys Pro Pro Ala Asp His Ile
915 920 925

Val Ser Thr Pro His Lys Asn Asp Gln Glu Phe Gln Ala Ala Ile Ser
930 935 940

Lys Thr Ser Trp Ser Trp Leu Phe Ala Leu Phe Gly Gly Ala Ser Ser
945 950 955 960

Leu Leu Ile Ile Gly Leu Met Ile Phe Ala Cys Ser Met Met Leu Thr
965 970 975

Ser Thr Arg Arg
980

50189A_SeqListing

<210> 6
 <211> 981
 <212> PRT
 <213> Sindbis virus

<400> 6

Met Ser Ala Ala Pro Leu Val Thr Ala Met Cys Leu Leu Gly Asn Val
 1 5 10 15

Ser Phe Pro Cys Asp Arg Pro Pro Thr Cys Tyr Thr Arg Glu Pro Ser
 20 25 30

Arg Ala Leu Asp Ile Leu Glu Glu Asn Val Asn His Glu Ala Tyr Asp
 35 40 45

Thr Leu Leu Asn Ala Ile Leu Arg Cys Gly Ser Ser Gly Arg Ser Lys
 50 55 60

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

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

Glu Gln Val Trp Asp Glu Ala Asp Asp Asn Thr Ile Arg Ile Gln Thr
 100 105 110

Ser Ala Gln Phe Gly Tyr Asp Gln Ser Gly Ala Ala Ser Ala Asn Lys
 115 120 125

Tyr Arg Tyr Met Ser Leu Glu Gln Asp His Thr Val Glu Glu Gly Thr
 130 135 140

Met Asp Asp Ile Lys Ile Ser Thr Ser Gly Pro Cys Arg Arg Leu Ser
 145 150 155 160

50189A_SeqListing

Tyr Lys Gly Tyr Phe Leu Leu Ala Lys Cys Pro Pro Gly Asp Ser Val
165 170 175

Thr Val Ser Ile Val Ser Ser Asn Ser Ala Thr Ser Cys Thr Leu Ala
180 185 190

Arg Lys Ile Lys Pro Lys Phe Val Gly Arg Glu Lys Tyr Asp Leu Pro
195 200 205

Pro Val His Gly Lys Lys Ile Pro Cys Thr Val Tyr Asp Arg Leu Glu
210 215 220

Thr Thr Ala Gly Tyr Ile Thr Met His Arg Pro Arg Pro His Ala Tyr
225 230 235 240

Thr Ser Tyr Leu Glu Glu Ser Ser Gly Lys Val Tyr Ala Lys Pro Pro
245 250 255

Ser Gly Lys Asn Ile Thr Tyr Glu Cys Lys Cys Gly Asp Tyr Lys Thr
260 265 270

Gly Thr Val Ser Thr Arg Thr Glu Ile Thr Gly Cys Thr Ala Ile Lys
275 280 285

Gln Cys Val Ala Tyr Lys Ser Asp Gln Thr Lys Trp Val Phe Asn Ser
290 295 300

Pro Asp Leu Ile Arg His Asp Asp His Thr Ala Gln Gly Lys Leu His
305 310 315 320

Leu Pro Phe Lys Leu Ile Pro Ser Thr Cys Met Val Pro Val Ala His
325 330 335

Ala Pro Asn Val Ile His Gly Phe Lys His Ile Ser Leu Gln Leu Asp
340 345 350

50189A_SeqListing

Thr Asp His Leu Thr Leu Leu Thr Thr Arg Arg Leu Gly Ala Asn Pro
355 360 365

Glu Pro Thr Thr Glu Trp Ile Val Gly Lys Thr Val Arg Asn Phe Thr
370 375 380

Val Asp Arg Asp Gly Leu Glu Tyr Ile Trp Gly Asn His Glu Pro Val
385 390 395 400

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

His Glu Ile Val Gln His Tyr Tyr His Arg His Pro Val Tyr Thr Ile
420 425 430

Leu Ala Val Ala Ser Ala Thr Val Ala Met Met Ile Gly Val Thr Val
435 440 445

Ala Val Leu Cys Ala Cys Lys Ala Arg Arg Glu Cys Leu Thr Pro Tyr
450 455 460

Ala Leu Ala Pro Asn Ala Val Ile Pro Thr Ser Leu Ala Leu Leu Cys
465 470 475 480

Cys Val Arg Ser Ala Asn Ala Glu Thr Phe Thr Glu Thr Met Ser Tyr
485 490 495

Leu Trp Ser Asn Ser Gln Pro Phe Phe Trp Val Gln Leu Cys Ile Pro
500 505 510

Leu Ala Ala Phe Ile Val Leu Met Arg Cys Cys Ser Cys Cys Leu Pro
515 520 525

Phe Leu Val Val Ala Gly Ala Tyr Leu Ala Lys Val Asp Ala Tyr Glu
530 535 540

50189A_SeqListing

His Ala Thr Thr Val Pro Asn Val Pro Gln Ile Pro Tyr Lys Ala Leu
545 550 555 560

Val Glu Arg Ala Gly Tyr Ala Pro Leu Asn Leu Glu Ile Thr Val Met
565 570 575

Ser Ser Glu Val Leu Pro Ser Thr Asn Gln Glu Tyr Ile Thr Cys Lys
580 585 590

Phe Thr Thr Val Val Pro Ser Pro Lys Ile Lys Cys Cys Gly Ser Leu
595 600 605

Glu Cys Gln Pro Ala Ala His Ala Asp Tyr Thr Cys Lys Val Phe Gly
610 615 620

Gly Val Tyr Pro Phe Met Trp Gly Gly Ala Gln Cys Phe Cys Asp Ser
625 630 635 640

Glu Asn Ser Gln Met Ser Glu Ala Tyr Val Glu Leu Ser Ala Asp Cys
645 650 655

Ala Ser Asp His Ala Gln Ala Ile Lys Val His Thr Ala Ala Met Lys
660 665 670

Val Gly Leu Arg Ile Val Tyr Gly Asn Thr Thr Ser Phe Leu Asp Val
675 680 685

Tyr Val Asn Gly Val Thr Pro Gly Thr Ser Lys Asp Leu Lys Val Ile
690 695 700

Ala Gly Pro Ile Ser Ala Ser Phe Thr Pro Phe Asp His Lys Val Val
705 710 715 720

Ile His Arg Gly Leu Val Tyr Asn Tyr Asp Phe Pro Glu Tyr Gly Ala
725 730 735

50189A_SeqListing

Met Lys Pro Gly Ala Phe Gly Asp Ile Gln Ala Thr Ser Leu Thr Ser
740 745 750

Lys Asp Leu Ile Ala Ser Thr Asp Ile Arg Leu Leu Lys Pro Ser Ala
755 760 765

Lys Asn Val His Val Pro Tyr Thr Gln Ala Ser Ser Gly Phe Glu Met
770 775 780

Trp Lys Asn Asn Ser Gly Arg Pro Leu Gln Glu Thr Ala Pro Phe Gly
785 790 795 800

Cys Lys Ile Ala Val Asn Pro Leu Arg Ala Val Asp Cys Ser Tyr Gly
805 810 815

Asn Ile Pro Ile Ser Ile Asp Ile Pro Asn Ala Ala Phe Ile Arg Thr
820 825 830

Ser Asp Ala Pro Leu Val Ser Thr Val Lys Cys Glu Val Ser Glu Cys
835 840 845

Thr Tyr Ser Ala Asp Phe Gly Gly Met Ala Thr Leu Gln Tyr Val Ser
850 855 860

Asp Arg Glu Gly Gln Cys Pro Val His Ser His Ser Ser Thr Ala Thr
865 870 875 880

Leu Gln Glu Ser Thr Val His Val Leu Glu Lys Gly Ala Val Thr Val
885 890 895

His Phe Ser Thr Ala Ser Pro Gln Ala Asn Phe Ile Val Ser Leu Cys
900 905 910

Gly Lys Lys Thr Thr Cys Asn Ala Glu Cys Lys Pro Pro Ala Asp His
915 920 925

50189A_SeqListing

Ile Val Ser Thr Pro His Lys Asn Asp Gln Glu Phe Gln Ala Ala Ile
930 935 940

Ser Lys Thr Ser Trp Ser Trp Leu Phe Ala Leu Phe Gly Gly Ala Ser
945 950 955 960

Ser Leu Leu Ile Ile Gly Leu Met Ile Phe Ala Cys Ser Met Met Leu
965 970 975

Thr Ser Thr Arg Arg
980

<210> 7
<211> 982
<212> PRT
<213> Sindbis virus

<400> 7

Met Ser Ala Ala Pro Leu Val Thr Ala Met Cys Leu Leu Gly Asn Val
1 5 10 15

Ser Phe Pro Cys Asp Arg Pro Pro Thr Cys Tyr Thr Arg Glu Pro Ser
20 25 30

Arg Ala Leu Asp Ile Leu Glu Glu Asn Val Asn His Glu Ala Tyr Asp
35 40 45

Thr Leu Leu Asn Ala Ile Leu Arg Cys Gly Ser Ser Gly Arg Ser Lys
50 55 60

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

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

50189A_SeqListing

Glu Gln Val Trp Asp Glu Ala Asp Asp Asn Thr Ile Arg Ile Gln Thr
100 105 110

Ser Ala Gln Phe Gly Tyr Asp Gln Ser Gly Ala Ala Ser Ala Asn Lys
115 120 125

Tyr Arg Tyr Met Ser Leu Lys Gln Asp His Thr Val Lys Glu Gly Thr
130 135 140

Met Asp Asp Ile Lys Ile Ser Thr Ser Gly Pro Cys Arg Arg Leu Ser
145 150 155 160

Tyr Lys Gly Tyr Phe Leu Leu Ala Lys Cys Pro Pro Gly Asp Ser Val
165 170 175

Thr Val Ser Ile Val Ser Ser Asn Ser Ala Thr Ser Cys Thr Leu Ala
180 185 190

Arg Lys Ile Lys Pro Lys Phe Val Gly Arg Glu Lys Tyr Asp Leu Pro
195 200 205

Pro Val His Gly Lys Lys Ile Pro Cys Thr Val Tyr Asp Arg Leu Glu
210 215 220

Gly Thr Thr Ala Gly Tyr Ile Thr Met His Arg Pro Arg Pro His Ala
225 230 235 240

Tyr Thr Ser Tyr Leu Glu Glu Ser Ser Gly Lys Val Tyr Ala Lys Pro
245 250 255

Pro Ser Gly Lys Asn Ile Thr Tyr Glu Cys Lys Cys Gly Asp Tyr Lys
260 265 270

Thr Gly Thr Val Ser Thr Arg Thr Glu Ile Thr Gly Cys Thr Ala Ile
275 280 285

50189A_SeqListing

Lys Gln Cys Val Ala Tyr Lys Ser Asp Gln Thr Lys Trp Val Phe Asn
 290 295 300

Ser Pro Asp Leu Ile Arg His Asp Asp His Thr Ala Gln Gly Lys Leu
 305 310 315 320

His Leu Pro Phe Lys Leu Ile Pro Ser Thr Cys Met Val Pro Val Ala
 325 330 335

His Ala Pro Asn Val Ile His Gly Phe Lys His Ile Ser Leu Gln Leu
 340 345 350

Asp Thr Asp His Leu Thr Leu Leu Thr Thr Arg Arg Leu Gly Ala Asn
 355 360 365

Pro Glu Pro Thr Thr Glu Trp Ile Val Gly Lys Thr Val Arg Asn Phe
 370 375 380

Thr Val Asp Arg Asp Gly Leu Glu Tyr Ile Trp Gly Asn His Glu Pro
 385 390 395 400

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

Pro His Glu Ile Val Gln His Tyr Tyr His Arg His Pro Val Tyr Thr
 420 425 430

Ile Leu Ala Val Ala Ser Ala Thr Val Ala Met Met Ile Gly Val Thr
 435 440 445

Val Ala Val Leu Cys Ala Cys Lys Ala Arg Arg Glu Cys Leu Thr Pro
 450 455 460

Tyr Ala Leu Ala Pro Asn Ala Val Ile Pro Thr Ser Leu Ala Leu Leu
 465 470 475 480

50189A_SeqListing

Cys Cys Val Arg Ser Ala Asn Ala Glu Thr Phe Thr Glu Thr Met Ser
485 490 495

Tyr Leu Trp Ser Asn Ser Gln Pro Phe Phe Trp Val Gln Leu Cys Ile
500 505 510

Pro Leu Ala Ala Phe Ile Val Leu Met Arg Cys Cys Ser Cys Cys Leu
515 520 525

Pro Phe Leu Val Val Ala Gly Ala Tyr Leu Ala Lys Val Asp Ala Tyr
530 535 540

Glu His Ala Thr Thr Val Pro Asn Val Pro Gln Ile Pro Tyr Lys Ala
545 550 555 560

Leu Val Glu Arg Ala Gly Tyr Ala Pro Leu Asn Leu Glu Ile Thr Val
565 570 575

Met Ser Ser Glu Val Leu Pro Ser Thr Asn Gln Glu Tyr Ile Thr Cys
580 585 590

Lys Phe Thr Thr Val Val Pro Ser Pro Lys Ile Lys Cys Cys Gly Ser
595 600 605

Leu Glu Cys Gln Pro Ala Ala His Ala Asp Tyr Thr Cys Lys Val Phe
610 615 620

Gly Gly Val Tyr Pro Phe Met Trp Gly Gly Ala Gln Cys Phe Cys Asp
625 630 635 640

Ser Glu Asn Ser Gln Met Ser Glu Ala Tyr Val Glu Leu Ser Ala Asp
645 650 655

Cys Ala Ser Asp His Ala Gln Ala Ile Lys Val His Thr Ala Ala Met
660 665 670

50189A_SeqListing

Lys Val Gly Leu Arg Ile Val Tyr Gly Asn Thr Thr Ser Phe Leu Asp
675 680 685

Val Tyr Val Asn Gly Val Thr Pro Gly Thr Ser Lys Asp Leu Lys Val
690 695 700

Ile Ala Gly Pro Ile Ser Ala Ser Phe Thr Pro Phe Asp His Lys Val
705 710 715 720

Val Ile His Arg Gly Leu Val Tyr Asn Tyr Asp Phe Pro Glu Tyr Gly
725 730 735

Ala Met Lys Pro Gly Ala Phe Gly Asp Ile Gln Ala Thr Ser Leu Thr
740 745 750

Ser Lys Asp Leu Ile Ala Ser Thr Asp Ile Arg Leu Leu Lys Pro Ser
755 760 765

Ala Lys Asn Val His Val Pro Tyr Thr Gln Ala Ser Ser Gly Phe Glu
770 775 780

Met Trp Lys Asn Asn Ser Gly Arg Pro Leu Gln Glu Thr Ala Pro Phe
785 790 795 800

Gly Cys Lys Ile Ala Val Asn Pro Leu Arg Ala Val Asp Cys Ser Tyr
805 810 815

Gly Asn Ile Pro Ile Ser Ile Asp Ile Pro Asn Ala Ala Phe Ile Arg
820 825 830

Thr Ser Asp Ala Pro Leu Val Ser Thr Val Lys Cys Glu Val Ser Glu
835 840 845

Cys Thr Tyr Ser Ala Asp Phe Gly Gly Met Ala Thr Leu Gln Tyr Val
850 855 860

50189A_SeqListing

Ser Asp Arg Glu Gly Gln Cys Pro Val His Ser His Ser Ser Thr Ala
865 870 875 880

Thr Leu Gln Glu Ser Thr Val His Val Leu Glu Lys Gly Ala Val Thr
885 890 895

Val His Phe Ser Thr Ala Ser Pro Gln Ala Asn Phe Ile Val Ser Leu
900 905 910

Cys Gly Lys Lys Thr Thr Cys Asn Ala Glu Cys Lys Pro Pro Ala Asp
915 920 925

His Ile Val Ser Thr Pro His Lys Asn Asp Gln Glu Phe Gln Ala Ala
930 935 940

Ile Ser Lys Thr Ser Trp Ser Trp Leu Phe Ala Leu Phe Gly Gly Ala
945 950 955 960

Ser Ser Leu Leu Ile Ile Gly Leu Met Ile Phe Ala Cys Ser Met Met
965 970 975

Leu Thr Ser Thr Arg Arg
980

<210> 8
<211> 981
<212> PRT
<213> Sindbis virus

<400> 8

Met Ser Ala Ala Pro Leu Val Thr Ala Met Cys Leu Leu Gly Asn Val
1 5 10 15

Ser Phe Pro Cys Asp Arg Pro Pro Thr Cys Tyr Thr Arg Glu Pro Ser
20 25 30

Arg Ala Leu Asp Ile Leu Glu Glu Asn Val Asn His Glu Ala Tyr Asp

50189A_SeqListing

35

40

45

Thr Leu Leu Asn Ala Ile Leu Arg Cys Gly Ser Ser Gly Arg Ser Lys
 50 55 60

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

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

Glu Gln Val Trp Asp Glu Ala Asp Asp Asn Thr Ile Arg Ile Gln Thr
 100 105 110

Ser Ala Gln Phe Gly Tyr Asp Gln Ser Gly Ala Ala Ser Ala Asn Lys
 115 120 125

Tyr Arg Tyr Met Ser Leu Lys Gln Asp His Thr Val Lys Glu Gly Thr
 130 135 140

Met Asp Asp Ile Lys Ile Ser Thr Ser Gly Pro Cys Arg Arg Leu Ser
 145 150 155 160

Tyr Lys Gly Tyr Phe Leu Leu Ala Lys Cys Pro Pro Gly Asp Ser Val
 165 170 175

Thr Val Ser Ile Val Ser Ser Asn Ser Ala Thr Ser Cys Thr Leu Ala
 180 185 190

Arg Lys Ile Lys Pro Lys Phe Val Gly Arg Glu Lys Tyr Asp Leu Pro
 195 200 205

Pro Val His Gly Lys Lys Ile Pro Cys Thr Val Tyr Asp Arg Leu Glu
 210 215 220

Thr Thr Ala Gly Tyr Ile Thr Met His Arg Pro Arg Pro His Ala Tyr

50189A_SeqListing

225		230		235		240									
Thr	Ser	Tyr	Leu	Glu	Glu	Ser	Ser	Gly	Lys	Val	Tyr	Ala	Lys	Pro	Pro
			245						250					255	
Ser	Gly	Lys	Asn	Ile	Thr	Tyr	Glu	Cys	Lys	Cys	Gly	Asp	Tyr	Lys	Thr
			260					265					270		
Gly	Thr	Val	Ser	Thr	Arg	Thr	Glu	Ile	Thr	Gly	Cys	Thr	Ala	Ile	Lys
		275					280					285			
Gln	Cys	Val	Ala	Tyr	Lys	Ser	Asp	Gln	Thr	Lys	Trp	Val	Phe	Asn	Ser
	290					295					300				
Pro	Asp	Leu	Ile	Arg	His	Asp	Asp	His	Thr	Ala	Gln	Gly	Lys	Leu	His
305					310					315					320
Leu	Pro	Phe	Lys	Leu	Ile	Pro	Ser	Thr	Cys	Met	Val	Pro	Val	Ala	His
				325					330					335	
Ala	Pro	Asn	Val	Ile	His	Gly	Phe	Lys	His	Ile	Ser	Leu	Gln	Leu	Asp
			340					345					350		
Thr	Asp	His	Leu	Thr	Leu	Leu	Thr	Thr	Arg	Arg	Leu	Gly	Ala	Asn	Pro
		355					360					365			
Glu	Pro	Thr	Thr	Glu	Trp	Ile	Val	Gly	Lys	Thr	Val	Arg	Asn	Phe	Thr
	370					375					380				
Val	Asp	Arg	Asp	Gly	Leu	Glu	Tyr	Ile	Trp	Gly	Asn	His	Glu	Pro	Val
385					390					395					400
Arg	Val	Tyr	Ala	Gln	Glu	Ser	Ala	Pro	Gly	Asp	Pro	His	Gly	Trp	Pro
				405					410					415	
His	Glu	Ile	Val	Gln	His	Tyr	Tyr	His	Arg	His	Pro	Val	Tyr	Thr	Ile

50189A_SeqListing

420

425

430

Leu Ala Val Ala Ser Ala Thr Val Ala Met Met Ile Gly Val Thr Val
435 440 445

Ala Val Leu Cys Ala Cys Lys Ala Arg Arg Glu Cys Leu Thr Pro Tyr
450 455 460

Ala Leu Ala Pro Asn Ala Val Ile Pro Thr Ser Leu Ala Leu Leu Cys
465 470 475 480

Cys Val Arg Ser Ala Asn Ala Glu Thr Phe Thr Glu Thr Met Ser Tyr
485 490 495

Leu Trp Ser Asn Ser Gln Pro Phe Phe Trp Val Gln Leu Cys Ile Pro
500 505 510

Leu Ala Ala Phe Ile Val Leu Met Arg Cys Cys Ser Cys Cys Leu Pro
515 520 525

Phe Leu Val Val Ala Gly Ala Tyr Leu Ala Lys Val Asp Ala Tyr Glu
530 535 540

His Ala Thr Thr Val Pro Asn Val Pro Gln Ile Pro Tyr Lys Ala Leu
545 550 555 560

Val Glu Arg Ala Gly Tyr Ala Pro Leu Asn Leu Glu Ile Thr Val Met
565 570 575

Ser Ser Glu Val Leu Pro Ser Thr Asn Gln Glu Tyr Ile Thr Cys Lys
580 585 590

Phe Thr Thr Val Val Pro Ser Pro Lys Ile Lys Cys Cys Gly Ser Leu
595 600 605

Glu Cys Gln Pro Ala Ala His Ala Asp Tyr Thr Cys Lys Val Phe Gly

50189A_SeqListing

610

615

620

Gly Val Tyr Pro Phe Met Trp Gly Gly Ala Gln Cys Phe Cys Asp Ser
 625 630 635 640

Glu Asn Ser Gln Met Ser Glu Ala Tyr Val Glu Leu Ser Ala Asp Cys
 645 650 655

Ala Ser Asp His Ala Gln Ala Ile Lys Val His Thr Ala Ala Met Lys
 660 665 670

Val Gly Leu Arg Ile Val Tyr Gly Asn Thr Thr Ser Phe Leu Asp Val
 675 680 685

Tyr Val Asn Gly Val Thr Pro Gly Thr Ser Lys Asp Leu Lys Val Ile
 690 695 700

Ala Gly Pro Ile Ser Ala Ser Phe Thr Pro Phe Asp His Lys Val Val
 705 710 715 720

Ile His Arg Gly Leu Val Tyr Asn Tyr Asp Phe Pro Glu Tyr Gly Ala
 725 730 735

Met Lys Pro Gly Ala Phe Gly Asp Ile Gln Ala Thr Ser Leu Thr Ser
 740 745 750

Lys Asp Leu Ile Ala Ser Thr Asp Ile Arg Leu Leu Lys Pro Ser Ala
 755 760 765

Lys Asn Val His Val Pro Tyr Thr Gln Ala Ser Ser Gly Phe Glu Met
 770 775 780

Trp Lys Asn Asn Ser Gly Arg Pro Leu Gln Glu Thr Ala Pro Phe Gly
 785 790 795 800

Cys Lys Ile Ala Val Asn Pro Leu Arg Ala Val Asp Cys Ser Tyr Gly

50189A_SeqListing

805

810

815

Asn Ile Pro Ile Ser Ile Asp Ile Pro Asn Ala Ala Phe Ile Arg Thr
 820 825 830

Ser Asp Ala Pro Leu Val Ser Thr Val Lys Cys Glu Val Ser Glu Cys
 835 840 845

Thr Tyr Ser Ala Asp Phe Gly Gly Met Ala Thr Leu Gln Tyr Val Ser
 850 855 860

Asp Arg Glu Gly Gln Cys Pro Val His Ser His Ser Ser Thr Ala Thr
 865 870 875 880

Leu Gln Glu Ser Thr Val His Val Leu Glu Lys Gly Ala Val Thr Val
 885 890 895

His Phe Ser Thr Ala Ser Pro Gln Ala Asn Phe Ile Val Ser Leu Cys
 900 905 910

Gly Lys Lys Thr Thr Cys Asn Ala Glu Cys Lys Pro Pro Ala Asp His
 915 920 925

Ile Val Ser Thr Pro His Lys Asn Asp Gln Glu Phe Gln Ala Ala Ile
 930 935 940

Ser Lys Thr Ser Trp Ser Trp Leu Phe Ala Leu Phe Gly Gly Ala Ser
 945 950 955 960

Ser Leu Leu Ile Ile Gly Leu Met Ile Phe Ala Cys Ser Met Met Leu
 965 970 975

Thr Ser Thr Arg Arg
 980

<210> 9

50189A_SeqListing

<211> 982

<212> PRT

<213> Sindbis virus

<400> 9

Met Ser Ala Ala Pro Leu Val Thr Ala Met Cys Leu Leu Gly Asn Val
1 5 10 15

Ser Phe Pro Cys Asp Arg Pro Pro Thr Cys Tyr Thr Arg Glu Pro Ser
20 25 30

Arg Ala Leu Asp Ile Leu Glu Glu Asn Val Asn His Glu Ala Tyr Asp
35 40 45

Thr Leu Leu Asn Ala Ile Leu Arg Cys Gly Ser Ser Gly Arg Ser Lys
50 55 60

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

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

Glu Gln Val Trp Asp Glu Ala Asp Asp Asn Thr Ile Arg Ile Gln Thr
100 105 110

Ser Ala Gln Phe Gly Tyr Asp Gln Ser Gly Ala Ala Ser Ala Asn Lys
115 120 125

Tyr Arg Tyr Met Ser Leu Lys Gln Asp His Thr Val Glu Glu Gly Thr
130 135 140

Met Asp Asp Ile Lys Ile Ser Thr Ser Gly Pro Cys Arg Arg Leu Ser
145 150 155 160

Tyr Lys Gly Tyr Phe Leu Leu Ala Lys Cys Pro Pro Gly Asp Ser Val
165 170 175

50189A_SeqListing

Thr Val Ser Ile Val Ser Ser Asn Ser Ala Thr Ser Cys Thr Leu Ala
180 185 190

Arg Lys Ile Lys Pro Lys Phe Val Gly Arg Glu Lys Tyr Asp Leu Pro
195 200 205

Pro Val His Gly Lys Lys Ile Pro Cys Thr Val Tyr Asp Arg Leu Glu
210 215 220

Gly Thr Thr Ala Gly Tyr Ile Thr Met His Arg Pro Arg Pro His Ala
225 230 235 240

Tyr Thr Ser Tyr Leu Glu Glu Ser Ser Gly Lys Val Tyr Ala Lys Pro
245 250 255

Pro Ser Gly Lys Asn Ile Thr Tyr Glu Cys Lys Cys Gly Asp Tyr Lys
260 265 270

Thr Gly Thr Val Ser Thr Arg Thr Glu Ile Thr Gly Cys Thr Ala Ile
275 280 285

Lys Gln Cys Val Ala Tyr Lys Ser Asp Gln Thr Lys Trp Val Phe Asn
290 295 300

Ser Pro Asp Leu Ile Arg His Asp Asp His Thr Ala Gln Gly Lys Leu
305 310 315 320

His Leu Pro Phe Lys Leu Ile Pro Ser Thr Cys Met Val Pro Val Ala
325 330 335

His Ala Pro Asn Val Ile His Gly Phe Lys His Ile Ser Leu Gln Leu
340 345 350

Asp Thr Asp His Leu Thr Leu Leu Thr Thr Arg Arg Leu Gly Ala Asn
355 360 365

50189A_SeqListing

Pro Glu Pro Thr Thr Glu Trp Ile Val Gly Lys Thr Val Arg Asn Phe
370 375 380

Thr Val Asp Arg Asp Gly Leu Glu Tyr Ile Trp Gly Asn His Glu Pro
385 390 395 400

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

Pro His Glu Ile Val Gln His Tyr Tyr His Arg His Pro Val Tyr Thr
420 425 430

Ile Leu Ala Val Ala Ser Ala Thr Val Ala Met Met Ile Gly Val Thr
435 440 445

Val Ala Val Leu Cys Ala Cys Lys Ala Arg Arg Glu Cys Leu Thr Pro
450 455 460

Tyr Ala Leu Ala Pro Asn Ala Val Ile Pro Thr Ser Leu Ala Leu Leu
465 470 475 480

Cys Cys Val Arg Ser Ala Asn Ala Glu Thr Phe Thr Glu Thr Met Ser
485 490 495

Tyr Leu Trp Ser Asn Ser Gln Pro Phe Phe Trp Val Gln Leu Cys Ile
500 505 510

Pro Leu Ala Ala Phe Ile Val Leu Met Arg Cys Cys Ser Cys Cys Leu
515 520 525

Pro Phe Leu Val Val Ala Gly Ala Tyr Leu Ala Lys Val Asp Ala Tyr
530 535 540

Glu His Ala Thr Thr Val Pro Asn Val Pro Gln Ile Pro Tyr Lys Ala
545 550 555 560

50189A_SeqListing

Leu Val Glu Arg Ala Gly Tyr Ala Pro Leu Asn Leu Glu Ile Thr Val
565 570 575

Met Ser Ser Glu Val Leu Pro Ser Thr Asn Gln Glu Tyr Ile Thr Cys
580 585 590

Lys Phe Thr Thr Val Val Pro Ser Pro Lys Ile Lys Cys Cys Gly Ser
595 600 605

Leu Glu Cys Gln Pro Ala Ala His Ala Asp Tyr Thr Cys Lys Val Phe
610 615 620

Gly Gly Val Tyr Pro Phe Met Trp Gly Gly Ala Gln Cys Phe Cys Asp
625 630 635 640

Ser Glu Asn Ser Gln Met Ser Glu Ala Tyr Val Glu Leu Ser Ala Asp
645 650 655

Cys Ala Ser Asp His Ala Gln Ala Ile Lys Val His Thr Ala Ala Met
660 665 670

Lys Val Gly Leu Arg Ile Val Tyr Gly Asn Thr Thr Ser Phe Leu Asp
675 680 685

Val Tyr Val Asn Gly Val Thr Pro Gly Thr Ser Lys Asp Leu Lys Val
690 695 700

Ile Ala Gly Pro Ile Ser Ala Ser Phe Thr Pro Phe Asp His Lys Val
705 710 715 720

Val Ile His Arg Gly Leu Val Tyr Asn Tyr Asp Phe Pro Glu Tyr Gly
725 730 735

Ala Met Lys Pro Gly Ala Phe Gly Asp Ile Gln Ala Thr Ser Leu Thr
740 745 750

50189A_SeqListing

Ser Lys Asp Leu Ile Ala Ser Thr Asp Ile Arg Leu Leu Lys Pro Ser
755 760 765

Ala Lys Asn Val His Val Pro Tyr Thr Gln Ala Ser Ser Gly Phe Glu
770 775 780

Met Trp Lys Asn Asn Ser Gly Arg Pro Leu Gln Glu Thr Ala Pro Phe
785 790 795 800

Gly Cys Lys Ile Ala Val Asn Pro Leu Arg Ala Val Asp Cys Ser Tyr
805 810 815

Gly Asn Ile Pro Ile Ser Ile Asp Ile Pro Asn Ala Ala Phe Ile Arg
820 825 830

Thr Ser Asp Ala Pro Leu Val Ser Thr Val Lys Cys Glu Val Ser Glu
835 840 845

Cys Thr Tyr Ser Ala Asp Phe Gly Gly Met Ala Thr Leu Gln Tyr Val
850 855 860

Ser Asp Arg Glu Gly Gln Cys Pro Val His Ser His Ser Ser Thr Ala
865 870 875 880

Thr Leu Gln Glu Ser Thr Val His Val Leu Glu Lys Gly Ala Val Thr
885 890 895

Val His Phe Ser Thr Ala Ser Pro Gln Ala Asn Phe Ile Val Ser Leu
900 905 910

Cys Gly Lys Lys Thr Thr Cys Asn Ala Glu Cys Lys Pro Pro Ala Asp
915 920 925

His Ile Val Ser Thr Pro His Lys Asn Asp Gln Glu Phe Gln Ala Ala
930 935 940

50189A_SeqListing

Ile Ser Lys Thr Ser Trp Ser Trp Leu Phe Ala Leu Phe Gly Gly Ala
945 950 955 960

Ser Ser Leu Leu Ile Ile Gly Leu Met Ile Phe Ala Cys Ser Met Met
965 970 975

Leu Thr Ser Thr Arg Arg
980

<210> 10
<211> 981
<212> PRT
<213> Sindbis virus

<400> 10

Met Ser Ala Ala Pro Leu Val Thr Ala Met Cys Leu Leu Gly Asn Val
1 5 10 15

Ser Phe Pro Cys Asp Arg Pro Pro Thr Cys Tyr Thr Arg Glu Pro Ser
20 25 30

Arg Ala Leu Asp Ile Leu Glu Glu Asn Val Asn His Glu Ala Tyr Asp
35 40 45

Thr Leu Leu Asn Ala Ile Leu Arg Cys Gly Ser Ser Gly Arg Ser Lys
50 55 60

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

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

Glu Gln Val Trp Asp Glu Ala Asp Asp Asn Thr Ile Arg Ile Gln Thr
100 105 110

50189A_SeqListing

Ser Ala Gln Phe Gly Tyr Asp Gln Ser Gly Ala Ala Ser Ala Asn Lys
115 120 125

Tyr Arg Tyr Met Ser Leu Lys Gln Asp His Thr Val Glu Glu Gly Thr
130 135 140

Met Asp Asp Ile Lys Ile Ser Thr Ser Gly Pro Cys Arg Arg Leu Ser
145 150 155 160

Tyr Lys Gly Tyr Phe Leu Leu Ala Lys Cys Pro Pro Gly Asp Ser Val
165 170 175

Thr Val Ser Ile Val Ser Ser Asn Ser Ala Thr Ser Cys Thr Leu Ala
180 185 190

Arg Lys Ile Lys Pro Lys Phe Val Gly Arg Glu Lys Tyr Asp Leu Pro
195 200 205

Pro Val His Gly Lys Lys Ile Pro Cys Thr Val Tyr Asp Arg Leu Glu
210 215 220

Thr Thr Ala Gly Tyr Ile Thr Met His Arg Pro Arg Pro His Ala Tyr
225 230 235 240

Thr Ser Tyr Leu Glu Glu Ser Ser Gly Lys Val Tyr Ala Lys Pro Pro
245 250 255

Ser Gly Lys Asn Ile Thr Tyr Glu Cys Lys Cys Gly Asp Tyr Lys Thr
260 265 270

Gly Thr Val Ser Thr Arg Thr Glu Ile Thr Gly Cys Thr Ala Ile Lys
275 280 285

Gln Cys Val Ala Tyr Lys Ser Asp Gln Thr Lys Trp Val Phe Asn Ser
290 295 300

50189A_SeqListing

Pro Asp Leu Ile Arg His Asp Asp His Thr Ala Gln Gly Lys Leu His
305 310 315 320

Leu Pro Phe Lys Leu Ile Pro Ser Thr Cys Met Val Pro Val Ala His
325 330 335

Ala Pro Asn Val Ile His Gly Phe Lys His Ile Ser Leu Gln Leu Asp
340 345 350

Thr Asp His Leu Thr Leu Leu Thr Thr Arg Arg Leu Gly Ala Asn Pro
355 360 365

Glu Pro Thr Thr Glu Trp Ile Val Gly Lys Thr Val Arg Asn Phe Thr
370 375 380

Val Asp Arg Asp Gly Leu Glu Tyr Ile Trp Gly Asn His Glu Pro Val
385 390 395 400

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

His Glu Ile Val Gln His Tyr Tyr His Arg His Pro Val Tyr Thr Ile
420 425 430

Leu Ala Val Ala Ser Ala Thr Val Ala Met Met Ile Gly Val Thr Val
435 440 445

Ala Val Leu Cys Ala Cys Lys Ala Arg Arg Glu Cys Leu Thr Pro Tyr
450 455 460

Ala Leu Ala Pro Asn Ala Val Ile Pro Thr Ser Leu Ala Leu Leu Cys
465 470 475 480

Cys Val Arg Ser Ala Asn Ala Glu Thr Phe Thr Glu Thr Met Ser Tyr
485 490 495

50189A_SeqListing

Leu Trp Ser Asn Ser Gln Pro Phe Phe Trp Val Gln Leu Cys Ile Pro
500 505 510

Leu Ala Ala Phe Ile Val Leu Met Arg Cys Cys Ser Cys Cys Leu Pro
515 520 525

Phe Leu Val Val Ala Gly Ala Tyr Leu Ala Lys Val Asp Ala Tyr Glu
530 535 540

His Ala Thr Thr Val Pro Asn Val Pro Gln Ile Pro Tyr Lys Ala Leu
545 550 555 560

Val Glu Arg Ala Gly Tyr Ala Pro Leu Asn Leu Glu Ile Thr Val Met
565 570 575

Ser Ser Glu Val Leu Pro Ser Thr Asn Gln Glu Tyr Ile Thr Cys Lys
580 585 590

Phe Thr Thr Val Val Pro Ser Pro Lys Ile Lys Cys Cys Gly Ser Leu
595 600 605

Glu Cys Gln Pro Ala Ala His Ala Asp Tyr Thr Cys Lys Val Phe Gly
610 615 620

Gly Val Tyr Pro Phe Met Trp Gly Gly Ala Gln Cys Phe Cys Asp Ser
625 630 635 640

Glu Asn Ser Gln Met Ser Glu Ala Tyr Val Glu Leu Ser Ala Asp Cys
645 650 655

Ala Ser Asp His Ala Gln Ala Ile Lys Val His Thr Ala Ala Met Lys
660 665 670

Val Gly Leu Arg Ile Val Tyr Gly Asn Thr Thr Ser Phe Leu Asp Val
675 680 685

50189A_SeqListing

Tyr Val Asn Gly Val Thr Pro Gly Thr Ser Lys Asp Leu Lys Val Ile
690 695 700

Ala Gly Pro Ile Ser Ala Ser Phe Thr Pro Phe Asp His Lys Val Val
705 710 715 720

Ile His Arg Gly Leu Val Tyr Asn Tyr Asp Phe Pro Glu Tyr Gly Ala
725 730 735

Met Lys Pro Gly Ala Phe Gly Asp Ile Gln Ala Thr Ser Leu Thr Ser
740 745 750

Lys Asp Leu Ile Ala Ser Thr Asp Ile Arg Leu Leu Lys Pro Ser Ala
755 760 765

Lys Asn Val His Val Pro Tyr Thr Gln Ala Ser Ser Gly Phe Glu Met
770 775 780

Trp Lys Asn Asn Ser Gly Arg Pro Leu Gln Glu Thr Ala Pro Phe Gly
785 790 795 800

Cys Lys Ile Ala Val Asn Pro Leu Arg Ala Val Asp Cys Ser Tyr Gly
805 810 815

Asn Ile Pro Ile Ser Ile Asp Ile Pro Asn Ala Ala Phe Ile Arg Thr
820 825 830

Ser Asp Ala Pro Leu Val Ser Thr Val Lys Cys Glu Val Ser Glu Cys
835 840 845

Thr Tyr Ser Ala Asp Phe Gly Gly Met Ala Thr Leu Gln Tyr Val Ser
850 855 860

Asp Arg Glu Gly Gln Cys Pro Val His Ser His Ser Ser Thr Ala Thr
865 870 875 880

50189A_SeqListing

Leu Gln Glu Ser Thr Val His Val Leu Glu Lys Gly Ala Val Thr Val
885 890 895

His Phe Ser Thr Ala Ser Pro Gln Ala Asn Phe Ile Val Ser Leu Cys
900 905 910

Gly Lys Lys Thr Thr Cys Asn Ala Glu Cys Lys Pro Pro Ala Asp His
915 920 925

Ile Val Ser Thr Pro His Lys Asn Asp Gln Glu Phe Gln Ala Ala Ile
930 935 940

Ser Lys Thr Ser Trp Ser Trp Leu Phe Ala Leu Phe Gly Gly Ala Ser
945 950 955 960

Ser Leu Leu Ile Ile Gly Leu Met Ile Phe Ala Cys Ser Met Met Leu
965 970 975

Thr Ser Thr Arg Arg
980

<210> 11
<211> 982
<212> PRT
<213> Sindbis virus

<400> 11

Met Ser Ala Ala Pro Leu Val Thr Ala Met Cys Leu Leu Gly Asn Val
1 5 10 15

Ser Phe Pro Cys Asp Arg Pro Pro Thr Cys Tyr Thr Arg Glu Pro Ser
20 25 30

Arg Ala Leu Asp Ile Leu Glu Glu Asn Val Asn His Glu Ala Tyr Asp
35 40 45

50189A_SeqListing

Thr Leu Leu Asn Ala Ile Leu Arg Cys Gly Ser Ser Gly Arg Ser Lys
50 55 60

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

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

Glu Gln Val Trp Asp Glu Ala Asp Asp Asn Thr Ile Arg Ile Gln Thr
100 105 110

Ser Ala Gln Phe Gly Tyr Asp Gln Ser Gly Ala Ala Ser Ala Asn Lys
115 120 125

Tyr Arg Tyr Met Ser Leu Lys Gln Asp His Thr Val Glu Glu Gly Thr
130 135 140

Met Asp Asp Ile Lys Ile Ser Thr Ser Gly Pro Cys Arg Arg Leu Ser
145 150 155 160

Tyr Lys Gly Tyr Phe Leu Leu Ala Lys Cys Pro Pro Gly Asp Ser Val
165 170 175

Thr Val Ser Ile Val Ser Ser Asn Ser Ala Thr Ser Cys Thr Leu Ala
180 185 190

Arg Lys Ile Lys Pro Lys Phe Val Gly Arg Glu Lys Tyr Asp Leu Pro
195 200 205

Pro Val His Gly Lys Lys Ile Pro Cys Thr Val Tyr Asp Arg Leu Lys
210 215 220

Gly Thr Thr Ala Gly Tyr Ile Thr Met His Arg Pro Arg Pro His Ala
225 230 235 240

50189A_SeqListing

Tyr Thr Ser Tyr Leu Glu Glu Ser Ser Gly Lys Val Tyr Ala Lys Pro
 245 250 255

Pro Ser Gly Lys Asn Ile Thr Tyr Glu Cys Lys Cys Gly Asp Tyr Lys
 260 265 270

Thr Gly Thr Val Ser Thr Arg Thr Glu Ile Thr Gly Cys Thr Ala Ile
 275 280 285

Lys Gln Cys Val Ala Tyr Lys Ser Asp Gln Thr Lys Trp Val Phe Asn
 290 295 300

Ser Pro Asp Leu Ile Arg His Asp Asp His Thr Ala Gln Gly Lys Leu
 305 310 315 320

His Leu Pro Phe Lys Leu Ile Pro Ser Thr Cys Met Val Pro Val Ala
 325 330 335

His Ala Pro Asn Val Ile His Gly Phe Lys His Ile Ser Leu Gln Leu
 340 345 350

Asp Thr Asp His Leu Thr Leu Leu Thr Thr Arg Arg Leu Gly Ala Asn
 355 360 365

Pro Glu Pro Thr Thr Glu Trp Ile Val Gly Lys Thr Val Arg Asn Phe
 370 375 380

Thr Val Asp Arg Asp Gly Leu Glu Tyr Ile Trp Gly Asn His Glu Pro
 385 390 395 400

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

Pro His Glu Ile Val Gln His Tyr Tyr His Arg His Pro Val Tyr Thr
 420 425 430

50189A_SeqListing

Ile Leu Ala Val Ala Ser Ala Thr Val Ala Met Met Ile Gly Val Thr
435 440 445

Val Ala Val Leu Cys Ala Cys Lys Ala Arg Arg Glu Cys Leu Thr Pro
450 455 460

Tyr Ala Leu Ala Pro Asn Ala Val Ile Pro Thr Ser Leu Ala Leu Leu
465 470 475 480

Cys Cys Val Arg Ser Ala Asn Ala Glu Thr Phe Thr Glu Thr Met Ser
485 490 495

Tyr Leu Trp Ser Asn Ser Gln Pro Phe Phe Trp Val Gln Leu Cys Ile
500 505 510

Pro Leu Ala Ala Phe Ile Val Leu Met Arg Cys Cys Ser Cys Cys Leu
515 520 525

Pro Phe Leu Val Val Ala Gly Ala Tyr Leu Ala Lys Val Asp Ala Tyr
530 535 540

Glu His Ala Thr Thr Val Pro Asn Val Pro Gln Ile Pro Tyr Lys Ala
545 550 555 560

Leu Val Glu Arg Ala Gly Tyr Ala Pro Leu Asn Leu Glu Ile Thr Val
565 570 575

Met Ser Ser Glu Val Leu Pro Ser Thr Asn Gln Glu Tyr Ile Thr Cys
580 585 590

Lys Phe Thr Thr Val Val Pro Ser Pro Lys Ile Lys Cys Cys Gly Ser
595 600 605

Leu Glu Cys Gln Pro Ala Ala His Ala Asp Tyr Thr Cys Lys Val Phe
610 615 620

50189A_SeqListing

Gly Gly Val Tyr Pro Phe Met Trp Gly Gly Ala Gln Cys Phe Cys Asp
625 630 635 640

Ser Glu Asn Ser Gln Met Ser Glu Ala Tyr Val Glu Leu Ser Ala Asp
645 650 655

Cys Ala Ser Asp His Ala Gln Ala Ile Lys Val His Thr Ala Ala Met
660 665 670

Lys Val Gly Leu Arg Ile Val Tyr Gly Asn Thr Thr Ser Phe Leu Asp
675 680 685

Val Tyr Val Asn Gly Val Thr Pro Gly Thr Ser Lys Asp Leu Lys Val
690 695 700

Ile Ala Gly Pro Ile Ser Ala Ser Phe Thr Pro Phe Asp His Lys Val
705 710 715 720

Val Ile His Arg Gly Leu Val Tyr Asn Tyr Asp Phe Pro Glu Tyr Gly
725 730 735

Ala Met Lys Pro Gly Ala Phe Gly Asp Ile Gln Ala Thr Ser Leu Thr
740 745 750

Ser Lys Asp Leu Ile Ala Ser Thr Asp Ile Arg Leu Leu Lys Pro Ser
755 760 765

Ala Lys Asn Val His Val Pro Tyr Thr Gln Ala Ser Ser Gly Phe Glu
770 775 780

Met Trp Lys Asn Asn Ser Gly Arg Pro Leu Gln Glu Thr Ala Pro Phe
785 790 795 800

Gly Cys Lys Ile Ala Val Asn Pro Leu Arg Ala Val Asp Cys Ser Tyr
805 810 815

50189A_SeqListing

Gly Asn Ile Pro Ile Ser Ile Asp Ile Pro Asn Ala Ala Phe Ile Arg
820 825 830

Thr Ser Asp Ala Pro Leu Val Ser Thr Val Lys Cys Glu Val Ser Glu
835 840 845

Cys Thr Tyr Ser Ala Asp Phe Gly Gly Met Ala Thr Leu Gln Tyr Val
850 855 860

Ser Asp Arg Glu Gly Gln Cys Pro Val His Ser His Ser Ser Thr Ala
865 870 875 880

Thr Leu Gln Glu Ser Thr Val His Val Leu Glu Lys Gly Ala Val Thr
885 890 895

Val His Phe Ser Thr Ala Ser Pro Gln Ala Asn Phe Ile Val Ser Leu
900 905 910

Cys Gly Lys Lys Thr Thr Cys Asn Ala Glu Cys Lys Pro Pro Ala Asp
915 920 925

His Ile Val Ser Thr Pro His Lys Asn Asp Gln Glu Phe Gln Ala Ala
930 935 940

Ile Ser Lys Thr Ser Trp Ser Trp Leu Phe Ala Leu Phe Gly Gly Ala
945 950 955 960

Ser Ser Leu Leu Ile Ile Gly Leu Met Ile Phe Ala Cys Ser Met Met
965 970 975

Leu Thr Ser Thr Arg Arg
980

<210> 12
<211> 981
<212> PRT
<213> Sindbis virus

50189A_SeqListing

<400> 12

Met Ser Ala Ala Pro Leu Val Thr Ala Met Cys Leu Leu Gly Asn Val
1 5 10 15

Ser Phe Pro Cys Asp Arg Pro Pro Thr Cys Tyr Thr Arg Glu Pro Ser
20 25 30

Arg Ala Leu Asp Ile Leu Glu Glu Asn Val Asn His Glu Ala Tyr Asp
35 40 45

Thr Leu Leu Asn Ala Ile Leu Arg Cys Gly Ser Ser Gly Arg Ser Lys
50 55 60

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

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

Glu Gln Val Trp Asp Glu Ala Asp Asp Asn Thr Ile Arg Ile Gln Thr
100 105 110

Ser Ala Gln Phe Gly Tyr Asp Gln Ser Gly Ala Ala Ser Ala Asn Lys
115 120 125

Tyr Arg Tyr Met Ser Leu Lys Gln Asp His Thr Val Glu Glu Gly Thr
130 135 140

Met Asp Asp Ile Lys Ile Ser Thr Ser Gly Pro Cys Arg Arg Leu Ser
145 150 155 160

Tyr Lys Gly Tyr Phe Leu Leu Ala Lys Cys Pro Pro Gly Asp Ser Val
165 170 175

Thr Val Ser Ile Val Ser Ser Asn Ser Ala Thr Ser Cys Thr Leu Ala

50189A_SeqListing

180

185

190

Arg Lys Ile Lys Pro Lys Phe Val Gly Arg Glu Lys Tyr Asp Leu Pro
195 200 205

Pro Val His Gly Lys Lys Ile Pro Cys Thr Val Tyr Asp Arg Leu Lys
210 215 220

Thr Thr Ala Gly Tyr Ile Thr Met His Arg Pro Arg Pro His Ala Tyr
225 230 235 240

Thr Ser Tyr Leu Glu Glu Ser Ser Gly Lys Val Tyr Ala Lys Pro Pro
245 250 255

Ser Gly Lys Asn Ile Thr Tyr Glu Cys Lys Cys Gly Asp Tyr Lys Thr
260 265 270

Gly Thr Val Ser Thr Arg Thr Glu Ile Thr Gly Cys Thr Ala Ile Lys
275 280 285

Gln Cys Val Ala Tyr Lys Ser Asp Gln Thr Lys Trp Val Phe Asn Ser
290 295 300

Pro Asp Leu Ile Arg His Asp Asp His Thr Ala Gln Gly Lys Leu His
305 310 315 320

Leu Pro Phe Lys Leu Ile Pro Ser Thr Cys Met Val Pro Val Ala His
325 330 335

Ala Pro Asn Val Ile His Gly Phe Lys His Ile Ser Leu Gln Leu Asp
340 345 350

Thr Asp His Leu Thr Leu Leu Thr Thr Arg Arg Leu Gly Ala Asn Pro
355 360 365

Glu Pro Thr Thr Glu Trp Ile Val Gly Lys Thr Val Arg Asn Phe Thr

50189A_SeqListing

370

375

380

Val Asp Arg Asp Gly Leu Glu Tyr Ile Trp Gly Asn His Glu Pro Val
 385 390 395 400

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

His Glu Ile Val Gln His Tyr Tyr His Arg His Pro Val Tyr Thr Ile
 420 425 430

Leu Ala Val Ala Ser Ala Thr Val Ala Met Met Ile Gly Val Thr Val
 435 440 445

Ala Val Leu Cys Ala Cys Lys Ala Arg Arg Glu Cys Leu Thr Pro Tyr
 450 455 460

Ala Leu Ala Pro Asn Ala Val Ile Pro Thr Ser Leu Ala Leu Leu Cys
 465 470 475 480

Cys Val Arg Ser Ala Asn Ala Glu Thr Phe Thr Glu Thr Met Ser Tyr
 485 490 495

Leu Trp Ser Asn Ser Gln Pro Phe Phe Trp Val Gln Leu Cys Ile Pro
 500 505 510

Leu Ala Ala Phe Ile Val Leu Met Arg Cys Cys Ser Cys Cys Leu Pro
 515 520 525

Phe Leu Val Val Ala Gly Ala Tyr Leu Ala Lys Val Asp Ala Tyr Glu
 530 535 540

His Ala Thr Thr Val Pro Asn Val Pro Gln Ile Pro Tyr Lys Ala Leu
 545 550 555 560

Val Glu Arg Ala Gly Tyr Ala Pro Leu Asn Leu Glu Ile Thr Val Met

50189A_SeqListing

565

570

575

Ser Ser Glu Val Leu Pro Ser Thr Asn Gln Glu Tyr Ile Thr Cys Lys
 580 585 590

Phe Thr Thr Val Val Pro Ser Pro Lys Ile Lys Cys Cys Gly Ser Leu
 595 600 605

Glu Cys Gln Pro Ala Ala His Ala Asp Tyr Thr Cys Lys Val Phe Gly
 610 615 620

Gly Val Tyr Pro Phe Met Trp Gly Gly Ala Gln Cys Phe Cys Asp Ser
 625 630 635 640

Glu Asn Ser Gln Met Ser Glu Ala Tyr Val Glu Leu Ser Ala Asp Cys
 645 650 655

Ala Ser Asp His Ala Gln Ala Ile Lys Val His Thr Ala Ala Met Lys
 660 665 670

Val Gly Leu Arg Ile Val Tyr Gly Asn Thr Thr Ser Phe Leu Asp Val
 675 680 685

Tyr Val Asn Gly Val Thr Pro Gly Thr Ser Lys Asp Leu Lys Val Ile
 690 695 700

Ala Gly Pro Ile Ser Ala Ser Phe Thr Pro Phe Asp His Lys Val Val
 705 710 715 720

Ile His Arg Gly Leu Val Tyr Asn Tyr Asp Phe Pro Glu Tyr Gly Ala
 725 730 735

Met Lys Pro Gly Ala Phe Gly Asp Ile Gln Ala Thr Ser Leu Thr Ser
 740 745 750

Lys Asp Leu Ile Ala Ser Thr Asp Ile Arg Leu Leu Lys Pro Ser Ala

50189A_SeqListing

755

760

765

Lys Asn Val His Val Pro Tyr Thr Gln Ala Ser Ser Gly Phe Glu Met
 770 775 780

Trp Lys Asn Asn Ser Gly Arg Pro Leu Gln Glu Thr Ala Pro Phe Gly
 785 790 795 800

Cys Lys Ile Ala Val Asn Pro Leu Arg Ala Val Asp Cys Ser Tyr Gly
 805 810 815

Asn Ile Pro Ile Ser Ile Asp Ile Pro Asn Ala Ala Phe Ile Arg Thr
 820 825 830

Ser Asp Ala Pro Leu Val Ser Thr Val Lys Cys Glu Val Ser Glu Cys
 835 840 845

Thr Tyr Ser Ala Asp Phe Gly Gly Met Ala Thr Leu Gln Tyr Val Ser
 850 855 860

Asp Arg Glu Gly Gln Cys Pro Val His Ser His Ser Ser Thr Ala Thr
 865 870 875 880

Leu Gln Glu Ser Thr Val His Val Leu Glu Lys Gly Ala Val Thr Val
 885 890 895

His Phe Ser Thr Ala Ser Pro Gln Ala Asn Phe Ile Val Ser Leu Cys
 900 905 910

Gly Lys Lys Thr Thr Cys Asn Ala Glu Cys Lys Pro Pro Ala Asp His
 915 920 925

Ile Val Ser Thr Pro His Lys Asn Asp Gln Glu Phe Gln Ala Ala Ile
 930 935 940

Ser Lys Thr Ser Trp Ser Trp Leu Phe Ala Leu Phe Gly Gly Ala Ser

945 950 50189A_SeqListing 955 960

Thr Ser Thr Arg Arg
980

<400> 13

Ser Phe Pro Cys Asp Arg Pro Pro Thr Cys Tyr Thr Arg Glu Pro Ser
20 25 30

Thr Leu Leu Asn Ala Ile Leu Arg Cys Gly Ser Ser Gly Arg Ser Lys
50 55 60

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

Ser Ala Gln Phe Gly Tyr Asp Gln Ser Gly Ala Ala Ser Ala Asn Lys
115 120 125

50189A_SeqListing

Tyr Arg Tyr Met Ser Leu Glu Gln Asp His Thr Val Lys Glu Gly Thr
130 135 140

Met Asp Asp Ile Lys Ile Ser Thr Ser Gly Pro Cys Arg Arg Leu Ser
145 150 155 160

Tyr Lys Gly Tyr Phe Leu Leu Ala Lys Cys Pro Pro Gly Asp Ser Val
165 170 175

Thr Val Ser Ile Val Ser Ser Asn Ser Ala Thr Ser Cys Thr Leu Ala
180 185 190

Arg Lys Ile Lys Pro Lys Phe Val Gly Arg Glu Lys Tyr Asp Leu Pro
195 200 205

Pro Val His Gly Lys Lys Ile Pro Cys Thr Val Tyr Asp Arg Leu Lys
210 215 220

Gly Thr Thr Ala Gly Tyr Ile Thr Met His Arg Pro Arg Pro His Ala
225 230 235 240

Tyr Thr Ser Tyr Leu Glu Glu Ser Ser Gly Lys Val Tyr Ala Lys Pro
245 250 255

Pro Ser Gly Lys Asn Ile Thr Tyr Glu Cys Lys Cys Gly Asp Tyr Lys
260 265 270

Thr Gly Thr Val Ser Thr Arg Thr Glu Ile Thr Gly Cys Thr Ala Ile
275 280 285

Lys Gln Cys Val Ala Tyr Lys Ser Asp Gln Thr Lys Trp Val Phe Asn
290 295 300

Ser Pro Asp Leu Ile Arg His Asp Asp His Thr Ala Gln Gly Lys Leu
305 310 315 320

50189A_SeqListing

His Leu Pro Phe Lys Leu Ile Pro Ser Thr Cys Met Val Pro Val Ala
325 330 335

His Ala Pro Asn Val Ile His Gly Phe Lys His Ile Ser Leu Gln Leu
340 345 350

Asp Thr Asp His Leu Thr Leu Leu Thr Thr Arg Arg Leu Gly Ala Asn
355 360 365

Pro Glu Pro Thr Thr Glu Trp Ile Val Gly Lys Thr Val Arg Asn Phe
370 375 380

Thr Val Asp Arg Asp Gly Leu Glu Tyr Ile Trp Gly Asn His Glu Pro
385 390 395 400

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

Pro His Glu Ile Val Gln His Tyr Tyr His Arg His Pro Val Tyr Thr
420 425 430

Ile Leu Ala Val Ala Ser Ala Thr Val Ala Met Met Ile Gly Val Thr
435 440 445

Val Ala Val Leu Cys Ala Cys Lys Ala Arg Arg Glu Cys Leu Thr Pro
450 455 460

Tyr Ala Leu Ala Pro Asn Ala Val Ile Pro Thr Ser Leu Ala Leu Leu
465 470 475 480

Cys Cys Val Arg Ser Ala Asn Ala Glu Thr Phe Thr Glu Thr Met Ser
485 490 495

Tyr Leu Trp Ser Asn Ser Gln Pro Phe Phe Trp Val Gln Leu Cys Ile
500 505 510

50189A_SeqListing

Pro Leu Ala Ala Phe Ile Val Leu Met Arg Cys Cys Ser Cys Cys Leu
515 520 525

Pro Phe Leu Val Val Ala Gly Ala Tyr Leu Ala Lys Val Asp Ala Tyr
530 535 540

Glu His Ala Thr Thr Val Pro Asn Val Pro Gln Ile Pro Tyr Lys Ala
545 550 555 560

Leu Val Glu Arg Ala Gly Tyr Ala Pro Leu Asn Leu Glu Ile Thr Val
565 570 575

Met Ser Ser Glu Val Leu Pro Ser Thr Asn Gln Glu Tyr Ile Thr Cys
580 585 590

Lys Phe Thr Thr Val Val Pro Ser Pro Lys Ile Lys Cys Cys Gly Ser
595 600 605

Leu Glu Cys Gln Pro Ala Ala His Ala Asp Tyr Thr Cys Lys Val Phe
610 615 620

Gly Gly Val Tyr Pro Phe Met Trp Gly Gly Ala Gln Cys Phe Cys Asp
625 630 635 640

Ser Glu Asn Ser Gln Met Ser Glu Ala Tyr Val Glu Leu Ser Ala Asp
645 650 655

Cys Ala Ser Asp His Ala Gln Ala Ile Lys Val His Thr Ala Ala Met
660 665 670

Lys Val Gly Leu Arg Ile Val Tyr Gly Asn Thr Thr Ser Phe Leu Asp
675 680 685

Val Tyr Val Asn Gly Val Thr Pro Gly Thr Ser Lys Asp Leu Lys Val
690 695 700

50189A_SeqListing

Ile Ala Gly Pro Ile Ser Ala Ser Phe Thr Pro Phe Asp His Lys Val
705 710 715 720

Val Ile His Arg Gly Leu Val Tyr Asn Tyr Asp Phe Pro Glu Tyr Gly
725 730 735

Ala Met Lys Pro Gly Ala Phe Gly Asp Ile Gln Ala Thr Ser Leu Thr
740 745 750

Ser Lys Asp Leu Ile Ala Ser Thr Asp Ile Arg Leu Leu Lys Pro Ser
755 760 765

Ala Lys Asn Val His Val Pro Tyr Thr Gln Ala Ser Ser Gly Phe Glu
770 775 780

Met Trp Lys Asn Asn Ser Gly Arg Pro Leu Gln Glu Thr Ala Pro Phe
785 790 795 800

Gly Cys Lys Ile Ala Val Asn Pro Leu Arg Ala Val Asp Cys Ser Tyr
805 810 815

Gly Asn Ile Pro Ile Ser Ile Asp Ile Pro Asn Ala Ala Phe Ile Arg
820 825 830

Thr Ser Asp Ala Pro Leu Val Ser Thr Val Lys Cys Glu Val Ser Glu
835 840 845

Cys Thr Tyr Ser Ala Asp Phe Gly Gly Met Ala Thr Leu Gln Tyr Val
850 855 860

Ser Asp Arg Glu Gly Gln Cys Pro Val His Ser His Ser Ser Thr Ala
865 870 875 880

Thr Leu Gln Glu Ser Thr Val His Val Leu Glu Lys Gly Ala Val Thr
885 890 895

50189A_SeqListing

Val His Phe Ser Thr Ala Ser Pro Gln Ala Asn Phe Ile Val Ser Leu
900 905 910

Cys Gly Lys Lys Thr Thr Cys Asn Ala Glu Cys Lys Pro Pro Ala Asp
915 920 925

His Ile Val Ser Thr Pro His Lys Asn Asp Gln Glu Phe Gln Ala Ala
930 935 940

Ile Ser Lys Thr Ser Trp Ser Trp Leu Phe Ala Leu Phe Gly Gly Ala
945 950 955 960

Ser Ser Leu Leu Ile Ile Gly Leu Met Ile Phe Ala Cys Ser Met Met
965 970 975

Leu Thr Ser Thr Arg Arg
980

<210> 14
<211> 981
<212> PRT
<213> Sindbis virus

<400> 14

Met Ser Ala Ala Pro Leu Val Thr Ala Met Cys Leu Leu Gly Asn Val
1 5 10 15

Ser Phe Pro Cys Asp Arg Pro Pro Thr Cys Tyr Thr Arg Glu Pro Ser
20 25 30

Arg Ala Leu Asp Ile Leu Glu Glu Asn Val Asn His Glu Ala Tyr Asp
35 40 45

Thr Leu Leu Asn Ala Ile Leu Arg Cys Gly Ser Ser Gly Arg Ser Lys
50 55 60

50189A_SeqListing

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

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

Glu Gln Val Trp Asp Glu Ala Asp Asp Asn Thr Ile Arg Ile Gln Thr
100 105 110

Ser Ala Gln Phe Gly Tyr Asp Gln Ser Gly Ala Ala Ser Ala Asn Lys
115 120 125

Tyr Arg Tyr Met Ser Leu Glu Gln Asp His Thr Val Lys Glu Gly Thr
130 135 140

Met Asp Asp Ile Lys Ile Ser Thr Ser Gly Pro Cys Arg Arg Leu Ser
145 150 155 160

Tyr Lys Gly Tyr Phe Leu Leu Ala Lys Cys Pro Pro Gly Asp Ser Val
165 170 175

Thr Val Ser Ile Val Ser Ser Asn Ser Ala Thr Ser Cys Thr Leu Ala
180 185 190

Arg Lys Ile Lys Pro Lys Phe Val Gly Arg Glu Lys Tyr Asp Leu Pro
195 200 205

Pro Val His Gly Lys Lys Ile Pro Cys Thr Val Tyr Asp Arg Leu Lys
210 215 220

Thr Thr Ala Gly Tyr Ile Thr Met His Arg Pro Arg Pro His Ala Tyr
225 230 235 240

Thr Ser Tyr Leu Glu Glu Ser Ser Gly Lys Val Tyr Ala Lys Pro Pro
245 250 255

50189A_SeqListing

Ser Gly Lys Asn Ile Thr Tyr Glu Cys Lys Cys Gly Asp Tyr Lys Thr
260 265 270

Gly Thr Val Ser Thr Arg Thr Glu Ile Thr Gly Cys Thr Ala Ile Lys
275 280 285

Gln Cys Val Ala Tyr Lys Ser Asp Gln Thr Lys Trp Val Phe Asn Ser
290 295 300

Pro Asp Leu Ile Arg His Asp Asp His Thr Ala Gln Gly Lys Leu His
305 310 315 320

Leu Pro Phe Lys Leu Ile Pro Ser Thr Cys Met Val Pro Val Ala His
325 330 335

Ala Pro Asn Val Ile His Gly Phe Lys His Ile Ser Leu Gln Leu Asp
340 345 350

Thr Asp His Leu Thr Leu Leu Thr Thr Arg Arg Leu Gly Ala Asn Pro
355 360 365

Glu Pro Thr Thr Glu Trp Ile Val Gly Lys Thr Val Arg Asn Phe Thr
370 375 380

Val Asp Arg Asp Gly Leu Glu Tyr Ile Trp Gly Asn His Glu Pro Val
385 390 395 400

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

His Glu Ile Val Gln His Tyr Tyr His Arg His Pro Val Tyr Thr Ile
420 425 430

Leu Ala Val Ala Ser Ala Thr Val Ala Met Met Ile Gly Val Thr Val
435 440 445

50189A_SeqListing

Ala Val Leu Cys Ala Cys Lys Ala Arg Arg Glu Cys Leu Thr Pro Tyr
450 455 460

Ala Leu Ala Pro Asn Ala Val Ile Pro Thr Ser Leu Ala Leu Leu Cys
465 470 475 480

Cys Val Arg Ser Ala Asn Ala Glu Thr Phe Thr Glu Thr Met Ser Tyr
485 490 495

Leu Trp Ser Asn Ser Gln Pro Phe Phe Trp Val Gln Leu Cys Ile Pro
500 505 510

Leu Ala Ala Phe Ile Val Leu Met Arg Cys Cys Ser Cys Cys Leu Pro
515 520 525

Phe Leu Val Val Ala Gly Ala Tyr Leu Ala Lys Val Asp Ala Tyr Glu
530 535 540

His Ala Thr Thr Val Pro Asn Val Pro Gln Ile Pro Tyr Lys Ala Leu
545 550 555 560

Val Glu Arg Ala Gly Tyr Ala Pro Leu Asn Leu Glu Ile Thr Val Met
565 570 575

Ser Ser Glu Val Leu Pro Ser Thr Asn Gln Glu Tyr Ile Thr Cys Lys
580 585 590

Phe Thr Thr Val Val Pro Ser Pro Lys Ile Lys Cys Cys Gly Ser Leu
595 600 605

Glu Cys Gln Pro Ala Ala His Ala Asp Tyr Thr Cys Lys Val Phe Gly
610 615 620

Gly Val Tyr Pro Phe Met Trp Gly Gly Ala Gln Cys Phe Cys Asp Ser
625 630 635 640

50189A_SeqListing

Glu Asn Ser Gln Met Ser Glu Ala Tyr Val Glu Leu Ser Ala Asp Cys
645 650 655

Ala Ser Asp His Ala Gln Ala Ile Lys Val His Thr Ala Ala Met Lys
660 665 670

Val Gly Leu Arg Ile Val Tyr Gly Asn Thr Thr Ser Phe Leu Asp Val
675 680 685

Tyr Val Asn Gly Val Thr Pro Gly Thr Ser Lys Asp Leu Lys Val Ile
690 695 700

Ala Gly Pro Ile Ser Ala Ser Phe Thr Pro Phe Asp His Lys Val Val
705 710 715 720

Ile His Arg Gly Leu Val Tyr Asn Tyr Asp Phe Pro Glu Tyr Gly Ala
725 730 735

Met Lys Pro Gly Ala Phe Gly Asp Ile Gln Ala Thr Ser Leu Thr Ser
740 745 750

Lys Asp Leu Ile Ala Ser Thr Asp Ile Arg Leu Leu Lys Pro Ser Ala
755 760 765

Lys Asn Val His Val Pro Tyr Thr Gln Ala Ser Ser Gly Phe Glu Met
770 775 780

Trp Lys Asn Asn Ser Gly Arg Pro Leu Gln Glu Thr Ala Pro Phe Gly
785 790 795 800

Cys Lys Ile Ala Val Asn Pro Leu Arg Ala Val Asp Cys Ser Tyr Gly
805 810 815

Asn Ile Pro Ile Ser Ile Asp Ile Pro Asn Ala Ala Phe Ile Arg Thr
820 825 830

50189A_SeqListing

Ser Asp Ala Pro Leu Val Ser Thr Val Lys Cys Glu Val Ser Glu Cys
835 840 845

Thr Tyr Ser Ala Asp Phe Gly Gly Met Ala Thr Leu Gln Tyr Val Ser
850 855 860

Asp Arg Glu Gly Gln Cys Pro Val His Ser His Ser Ser Thr Ala Thr
865 870 875 880

Leu Gln Glu Ser Thr Val His Val Leu Glu Lys Gly Ala Val Thr Val
885 890 895

His Phe Ser Thr Ala Ser Pro Gln Ala Asn Phe Ile Val Ser Leu Cys
900 905 910

Gly Lys Lys Thr Thr Cys Asn Ala Glu Cys Lys Pro Pro Ala Asp His
915 920 925

Ile Val Ser Thr Pro His Lys Asn Asp Gln Glu Phe Gln Ala Ala Ile
930 935 940

Ser Lys Thr Ser Trp Ser Trp Leu Phe Ala Leu Phe Gly Gly Ala Ser
945 950 955 960

Ser Leu Leu Ile Ile Gly Leu Met Ile Phe Ala Cys Ser Met Met Leu
965 970 975

Thr Ser Thr Arg Arg
980

<210> 15
<211> 982
<212> PRT
<213> Sindbis virus

<400> 15

50189A_SeqListing

Met Ser Ala Ala Pro Leu Val Thr Ala Met Cys Leu Leu Gly Asn Val
1 5 10 15

Ser Phe Pro Cys Asp Arg Pro Pro Thr Cys Tyr Thr Arg Glu Pro Ser
20 25 30

Arg Ala Leu Asp Ile Leu Glu Glu Asn Val Asn His Glu Ala Tyr Asp
35 40 45

Thr Leu Leu Asn Ala Ile Leu Arg Cys Gly Ser Ser Gly Arg Ser Lys
50 55 60

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

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

Glu Gln Val Trp Asp Glu Ala Asp Asp Asn Thr Ile Arg Ile Gln Thr
100 105 110

Ser Ala Gln Phe Gly Tyr Asp Gln Ser Gly Ala Ala Ser Ala Asn Lys
115 120 125

Tyr Arg Tyr Met Ser Leu Glu Gln Asp His Thr Val Glu Glu Gly Thr
130 135 140

Met Asp Asp Ile Lys Ile Ser Thr Ser Gly Pro Cys Arg Arg Leu Ser
145 150 155 160

Tyr Lys Gly Tyr Phe Leu Leu Ala Lys Cys Pro Pro Gly Asp Ser Val
165 170 175

Thr Val Ser Ile Val Ser Ser Asn Ser Ala Thr Ser Cys Thr Leu Ala
180 185 190

50189A_SeqListing

Arg Lys Ile Lys Pro Lys Phe Val Gly Arg Glu Lys Tyr Asp Leu Pro
195 200 205

Pro Val His Gly Lys Lys Ile Pro Cys Thr Val Tyr Asp Arg Leu Lys
210 215 220

Gly Thr Thr Ala Gly Tyr Ile Thr Met His Arg Pro Arg Pro His Ala
225 230 235 240

Tyr Thr Ser Tyr Leu Glu Glu Ser Ser Gly Lys Val Tyr Ala Lys Pro
245 250 255

Pro Ser Gly Lys Asn Ile Thr Tyr Glu Cys Lys Cys Gly Asp Tyr Lys
260 265 270

Thr Gly Thr Val Ser Thr Arg Thr Glu Ile Thr Gly Cys Thr Ala Ile
275 280 285

Lys Gln Cys Val Ala Tyr Lys Ser Asp Gln Thr Lys Trp Val Phe Asn
290 295 300

Ser Pro Asp Leu Ile Arg His Asp Asp His Thr Ala Gln Gly Lys Leu
305 310 315 320

His Leu Pro Phe Lys Leu Ile Pro Ser Thr Cys Met Val Pro Val Ala
325 330 335

His Ala Pro Asn Val Ile His Gly Phe Lys His Ile Ser Leu Gln Leu
340 345 350

Asp Thr Asp His Leu Thr Leu Leu Thr Thr Arg Arg Leu Gly Ala Asn
355 360 365

Pro Glu Pro Thr Thr Glu Trp Ile Val Gly Lys Thr Val Arg Asn Phe
370 375 380

50189A_SeqListing

Thr Val Asp Arg Asp Gly Leu Glu Tyr Ile Trp Gly Asn His Glu Pro
385 390 395 400

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

Pro His Glu Ile Val Gln His Tyr Tyr His Arg His Pro Val Tyr Thr
420 425 430

Ile Leu Ala Val Ala Ser Ala Thr Val Ala Met Met Ile Gly Val Thr
435 440 445

Val Ala Val Leu Cys Ala Cys Lys Ala Arg Arg Glu Cys Leu Thr Pro
450 455 460

Tyr Ala Leu Ala Pro Asn Ala Val Ile Pro Thr Ser Leu Ala Leu Leu
465 470 475 480

Cys Cys Val Arg Ser Ala Asn Ala Glu Thr Phe Thr Glu Thr Met Ser
485 490 495

Tyr Leu Trp Ser Asn Ser Gln Pro Phe Phe Trp Val Gln Leu Cys Ile
500 505 510

Pro Leu Ala Ala Phe Ile Val Leu Met Arg Cys Cys Ser Cys Cys Leu
515 520 525

Pro Phe Leu Val Val Ala Gly Ala Tyr Leu Ala Lys Val Asp Ala Tyr
530 535 540

Glu His Ala Thr Thr Val Pro Asn Val Pro Gln Ile Pro Tyr Lys Ala
545 550 555 560

Leu Val Glu Arg Ala Gly Tyr Ala Pro Leu Asn Leu Glu Ile Thr Val
565 570 575

50189A_SeqListing

Met Ser Ser Glu Val Leu Pro Ser Thr Asn Gln Glu Tyr Ile Thr Cys
580 585 590

Lys Phe Thr Thr Val Val Pro Ser Pro Lys Ile Lys Cys Cys Gly Ser
595 600 605

Leu Glu Cys Gln Pro Ala Ala His Ala Asp Tyr Thr Cys Lys Val Phe
610 615 620

Gly Gly Val Tyr Pro Phe Met Trp Gly Gly Ala Gln Cys Phe Cys Asp
625 630 635 640

Ser Glu Asn Ser Gln Met Ser Glu Ala Tyr Val Glu Leu Ser Ala Asp
645 650 655

Cys Ala Ser Asp His Ala Gln Ala Ile Lys Val His Thr Ala Ala Met
660 665 670

Lys Val Gly Leu Arg Ile Val Tyr Gly Asn Thr Thr Ser Phe Leu Asp
675 680 685

Val Tyr Val Asn Gly Val Thr Pro Gly Thr Ser Lys Asp Leu Lys Val
690 695 700

Ile Ala Gly Pro Ile Ser Ala Ser Phe Thr Pro Phe Asp His Lys Val
705 710 715 720

Val Ile His Arg Gly Leu Val Tyr Asn Tyr Asp Phe Pro Glu Tyr Gly
725 730 735

Ala Met Lys Pro Gly Ala Phe Gly Asp Ile Gln Ala Thr Ser Leu Thr
740 745 750

Ser Lys Asp Leu Ile Ala Ser Thr Asp Ile Arg Leu Leu Lys Pro Ser
755 760 765

50189A_SeqListing

Ala Lys Asn Val His Val Pro Tyr Thr Gln Ala Ser Ser Gly Phe Glu
770 775 780

Met Trp Lys Asn Asn Ser Gly Arg Pro Leu Gln Glu Thr Ala Pro Phe
785 790 795 800

Gly Cys Lys Ile Ala Val Asn Pro Leu Arg Ala Val Asp Cys Ser Tyr
805 810 815

Gly Asn Ile Pro Ile Ser Ile Asp Ile Pro Asn Ala Ala Phe Ile Arg
820 825 830

Thr Ser Asp Ala Pro Leu Val Ser Thr Val Lys Cys Glu Val Ser Glu
835 840 845

Cys Thr Tyr Ser Ala Asp Phe Gly Gly Met Ala Thr Leu Gln Tyr Val
850 855 860

Ser Asp Arg Glu Gly Gln Cys Pro Val His Ser His Ser Ser Thr Ala
865 870 875 880

Thr Leu Gln Glu Ser Thr Val His Val Leu Glu Lys Gly Ala Val Thr
885 890 895

Val His Phe Ser Thr Ala Ser Pro Gln Ala Asn Phe Ile Val Ser Leu
900 905 910

Cys Gly Lys Lys Thr Thr Cys Asn Ala Glu Cys Lys Pro Pro Ala Asp
915 920 925

His Ile Val Ser Thr Pro His Lys Asn Asp Gln Glu Phe Gln Ala Ala
930 935 940

Ile Ser Lys Thr Ser Trp Ser Trp Leu Phe Ala Leu Phe Gly Gly Ala
945 950 955 960

50189A_SeqListing

Ser Ser Leu Leu Ile Ile Gly Leu Met Ile Phe Ala Cys Ser Met Met
 965 970 975

Leu Thr Ser Thr Arg Arg
 980

<210> 16
 <211> 981
 <212> PRT
 <213> Sindbis virus

<400> 16

Met Ser Ala Ala Pro Leu Val Thr Ala Met Cys Leu Leu Gly Asn Val
 1 5 10 15

Ser Phe Pro Cys Asp Arg Pro Pro Thr Cys Tyr Thr Arg Glu Pro Ser
 20 25 30

Arg Ala Leu Asp Ile Leu Glu Glu Asn Val Asn His Glu Ala Tyr Asp
 35 40 45

Thr Leu Leu Asn Ala Ile Leu Arg Cys Gly Ser Ser Gly Arg Ser Lys
 50 55 60

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

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

Glu Gln Val Trp Asp Glu Ala Asp Asp Asn Thr Ile Arg Ile Gln Thr
 100 105 110

Ser Ala Gln Phe Gly Tyr Asp Gln Ser Gly Ala Ala Ser Ala Asn Lys
 115 120 125

Tyr Arg Tyr Met Ser Leu Glu Gln Asp His Thr Val Glu Glu Gly Thr

50189A_SeqListing

130

135

140

Met Asp Asp Ile Lys Ile Ser Thr Ser Gly Pro Cys Arg Arg Leu Ser
145 150 155 160

Tyr Lys Gly Tyr Phe Leu Leu Ala Lys Cys Pro Pro Gly Asp Ser Val
165 170 175

Thr Val Ser Ile Val Ser Ser Asn Ser Ala Thr Ser Cys Thr Leu Ala
180 185 190

Arg Lys Ile Lys Pro Lys Phe Val Gly Arg Glu Lys Tyr Asp Leu Pro
195 200 205

Pro Val His Gly Lys Lys Ile Pro Cys Thr Val Tyr Asp Arg Leu Lys
210 215 220

Thr Thr Ala Gly Tyr Ile Thr Met His Arg Pro Arg Pro His Ala Tyr
225 230 235 240

Thr Ser Tyr Leu Glu Glu Ser Ser Gly Lys Val Tyr Ala Lys Pro Pro
245 250 255

Ser Gly Lys Asn Ile Thr Tyr Glu Cys Lys Cys Gly Asp Tyr Lys Thr
260 265 270

Gly Thr Val Ser Thr Arg Thr Glu Ile Thr Gly Cys Thr Ala Ile Lys
275 280 285

Gln Cys Val Ala Tyr Lys Ser Asp Gln Thr Lys Trp Val Phe Asn Ser
290 295 300

Pro Asp Leu Ile Arg His Asp Asp His Thr Ala Gln Gly Lys Leu His
305 310 315 320

Leu Pro Phe Lys Leu Ile Pro Ser Thr Cys Met Val Pro Val Ala His

50189A_SeqListing

325

330

335

Ala Pro Asn Val Ile His Gly Phe Lys His Ile Ser Leu Gln Leu Asp
 340 345 350

Thr Asp His Leu Thr Leu Leu Thr Thr Arg Arg Leu Gly Ala Asn Pro
 355 360 365

Glu Pro Thr Thr Glu Trp Ile Val Gly Lys Thr Val Arg Asn Phe Thr
 370 375 380

Val Asp Arg Asp Gly Leu Glu Tyr Ile Trp Gly Asn His Glu Pro Val
 385 390 395 400

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

His Glu Ile Val Gln His Tyr Tyr His Arg His Pro Val Tyr Thr Ile
 420 425 430

Leu Ala Val Ala Ser Ala Thr Val Ala Met Met Ile Gly Val Thr Val
 435 440 445

Ala Val Leu Cys Ala Cys Lys Ala Arg Arg Glu Cys Leu Thr Pro Tyr
 450 455 460

Ala Leu Ala Pro Asn Ala Val Ile Pro Thr Ser Leu Ala Leu Leu Cys
 465 470 475 480

Cys Val Arg Ser Ala Asn Ala Glu Thr Phe Thr Glu Thr Met Ser Tyr
 485 490 495

Leu Trp Ser Asn Ser Gln Pro Phe Phe Trp Val Gln Leu Cys Ile Pro
 500 505 510

Leu Ala Ala Phe Ile Val Leu Met Arg Cys Cys Ser Cys Cys Leu Pro

50189A_SeqListing

515

520

525

Phe Leu Val Val Ala Gly Ala Tyr Leu Ala Lys Val Asp Ala Tyr Glu
 530 535 540

His Ala Thr Thr Val Pro Asn Val Pro Gln Ile Pro Tyr Lys Ala Leu
 545 550 555 560

Val Glu Arg Ala Gly Tyr Ala Pro Leu Asn Leu Glu Ile Thr Val Met
 565 570 575

Ser Ser Glu Val Leu Pro Ser Thr Asn Gln Glu Tyr Ile Thr Cys Lys
 580 585 590

Phe Thr Thr Val Val Pro Ser Pro Lys Ile Lys Cys Cys Gly Ser Leu
 595 600 605

Glu Cys Gln Pro Ala Ala His Ala Asp Tyr Thr Cys Lys Val Phe Gly
 610 615 620

Gly Val Tyr Pro Phe Met Trp Gly Gly Ala Gln Cys Phe Cys Asp Ser
 625 630 635 640

Glu Asn Ser Gln Met Ser Glu Ala Tyr Val Glu Leu Ser Ala Asp Cys
 645 650 655

Ala Ser Asp His Ala Gln Ala Ile Lys Val His Thr Ala Ala Met Lys
 660 665 670

Val Gly Leu Arg Ile Val Tyr Gly Asn Thr Thr Ser Phe Leu Asp Val
 675 680 685

Tyr Val Asn Gly Val Thr Pro Gly Thr Ser Lys Asp Leu Lys Val Ile
 690 695 700

Ala Gly Pro Ile Ser Ala Ser Phe Thr Pro Phe Asp His Lys Val Val

50189A_SeqListing

705		710		715		720									
Ile	His	Arg	Gly	Leu	Val	Tyr	Asn	Tyr	Asp	Phe	Pro	Glu	Tyr	Gly	Ala
			725						730					735	
Met	Lys	Pro	Gly	Ala	Phe	Gly	Asp	Ile	Gln	Ala	Thr	Ser	Leu	Thr	Ser
			740					745					750		
Lys	Asp	Leu	Ile	Ala	Ser	Thr	Asp	Ile	Arg	Leu	Leu	Lys	Pro	Ser	Ala
		755					760					765			
Lys	Asn	Val	His	Val	Pro	Tyr	Thr	Gln	Ala	Ser	Ser	Gly	Phe	Glu	Met
	770					775					780				
Trp	Lys	Asn	Asn	Ser	Gly	Arg	Pro	Leu	Gln	Glu	Thr	Ala	Pro	Phe	Gly
785					790					795					800
Cys	Lys	Ile	Ala	Val	Asn	Pro	Leu	Arg	Ala	Val	Asp	Cys	Ser	Tyr	Gly
				805					810					815	
Asn	Ile	Pro	Ile	Ser	Ile	Asp	Ile	Pro	Asn	Ala	Ala	Phe	Ile	Arg	Thr
			820					825					830		
Ser	Asp	Ala	Pro	Leu	Val	Ser	Thr	Val	Lys	Cys	Glu	Val	Ser	Glu	Cys
		835					840					845			
Thr	Tyr	Ser	Ala	Asp	Phe	Gly	Gly	Met	Ala	Thr	Leu	Gln	Tyr	Val	Ser
	850					855					860				
Asp	Arg	Glu	Gly	Gln	Cys	Pro	Val	His	Ser	His	Ser	Ser	Thr	Ala	Thr
865					870					875					880
Leu	Gln	Glu	Ser	Thr	Val	His	Val	Leu	Glu	Lys	Gly	Ala	Val	Thr	Val
				885					890					895	
His	Phe	Ser	Thr	Ala	Ser	Pro	Gln	Ala	Asn	Phe	Ile	Val	Ser	Leu	Cys

50189A_SeqListing

900

905

910

Gly Lys Lys Thr Thr Cys Asn Ala Glu Cys Lys Pro Pro Ala Asp His
915 920 925

Ile Val Ser Thr Pro His Lys Asn Asp Gln Glu Phe Gln Ala Ala Ile
930 935 940

Ser Lys Thr Ser Trp Ser Trp Leu Phe Ala Leu Phe Gly Gly Ala Ser
945 950 955 960

Ser Leu Leu Ile Ile Gly Leu Met Ile Phe Ala Cys Ser Met Met Leu
965 970 975

Thr Ser Thr Arg Arg
980

<210> 17
<211> 982
<212> PRT
<213> Sindbis virus

<400> 17

Met Ser Ala Ala Pro Leu Val Thr Ala Met Cys Leu Leu Gly Asn Val
1 5 10 15

Ser Phe Pro Cys Asp Arg Pro Pro Thr Cys Tyr Thr Arg Glu Pro Ser
20 25 30

Arg Ala Leu Asp Ile Leu Glu Glu Asn Val Asn His Glu Ala Tyr Asp
35 40 45

Thr Leu Leu Asn Ala Ile Leu Arg Cys Gly Ser Ser Gly Arg Ser Lys
50 55 60

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

50189A_SeqListing

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

Glu Gln Val Trp Asp Glu Ala Asp Asp Asn Thr Ile Arg Ile Gln Thr
100 105 110

Ser Ala Gln Phe Gly Tyr Asp Gln Ser Gly Ala Ala Ser Ala Asn Lys
115 120 125

Tyr Arg Tyr Met Ser Leu Lys Gln Asp His Thr Val Lys Glu Gly Thr
130 135 140

Met Asp Asp Ile Lys Ile Ser Thr Ser Gly Pro Cys Arg Arg Leu Ser
145 150 155 160

Tyr Lys Gly Tyr Phe Leu Leu Ala Lys Cys Pro Pro Gly Asp Ser Val
165 170 175

Thr Val Ser Ile Val Ser Ser Asn Ser Ala Thr Ser Cys Thr Leu Ala
180 185 190

Arg Lys Ile Lys Pro Lys Phe Val Gly Arg Glu Lys Tyr Asp Leu Pro
195 200 205

Pro Val His Gly Lys Lys Ile Pro Cys Thr Val Tyr Asp Arg Leu Lys
210 215 220

Glu Thr Thr Ala Gly Tyr Ile Thr Met His Arg Pro Arg Pro His Ala
225 230 235 240

Tyr Thr Ser Tyr Leu Glu Glu Ser Ser Gly Lys Val Tyr Ala Lys Pro
245 250 255

Pro Ser Gly Lys Asn Ile Thr Tyr Glu Cys Lys Cys Gly Asp Tyr Lys
260 265 270

50189A_SeqListing

Thr Gly Thr Val Ser Thr Arg Thr Glu Ile Thr Gly Cys Thr Ala Ile
275 280 285

Lys Gln Cys Val Ala Tyr Lys Ser Asp Gln Thr Lys Trp Val Phe Asn
290 295 300

Ser Pro Asp Leu Ile Arg His Asp Asp His Thr Ala Gln Gly Lys Leu
305 310 315 320

His Leu Pro Phe Lys Leu Ile Pro Ser Thr Cys Met Val Pro Val Ala
325 330 335

His Ala Pro Asn Val Ile His Gly Phe Lys His Ile Ser Leu Gln Leu
340 345 350

Asp Thr Asp His Leu Thr Leu Leu Thr Thr Arg Arg Leu Gly Ala Asn
355 360 365

Pro Glu Pro Thr Thr Glu Trp Ile Val Gly Lys Thr Val Arg Asn Phe
370 375 380

Thr Val Asp Arg Asp Gly Leu Glu Tyr Ile Trp Gly Asn His Glu Pro
385 390 395 400

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

Pro His Glu Ile Val Gln His Tyr Tyr His Arg His Pro Val Tyr Thr
420 425 430

Ile Leu Ala Val Ala Ser Ala Thr Val Ala Met Met Ile Gly Val Thr
435 440 445

Val Ala Val Leu Cys Ala Cys Lys Ala Arg Arg Glu Cys Leu Thr Pro
450 455 460

50189A_SeqListing

Tyr Ala Leu Ala Pro Asn Ala Val Ile Pro Thr Ser Leu Ala Leu Leu
465 470 475 480

Cys Cys Val Arg Ser Ala Asn Ala Glu Thr Phe Thr Glu Thr Met Ser
485 490 495

Tyr Leu Trp Ser Asn Ser Gln Pro Phe Phe Trp Val Gln Leu Cys Ile
500 505 510

Pro Leu Ala Ala Phe Ile Val Leu Met Arg Cys Cys Ser Cys Cys Leu
515 520 525

Pro Phe Leu Val Val Ala Gly Ala Tyr Leu Ala Lys Val Asp Ala Tyr
530 535 540

Glu His Ala Thr Thr Val Pro Asn Val Pro Gln Ile Pro Tyr Lys Ala
545 550 555 560

Leu Val Glu Arg Ala Gly Tyr Ala Pro Leu Asn Leu Glu Ile Thr Val
565 570 575

Met Ser Ser Glu Val Leu Pro Ser Thr Asn Gln Glu Tyr Ile Thr Cys
580 585 590

Lys Phe Thr Thr Val Val Pro Ser Pro Lys Ile Lys Cys Cys Gly Ser
595 600 605

Leu Glu Cys Gln Pro Ala Ala His Ala Asp Tyr Thr Cys Lys Val Phe
610 615 620

Gly Gly Val Tyr Pro Phe Met Trp Gly Gly Ala Gln Cys Phe Cys Asp
625 630 635 640

Ser Glu Asn Ser Gln Met Ser Glu Ala Tyr Val Glu Leu Ser Ala Asp
645 650 655

50189A_SeqListing

Cys Ala Ser Asp His Ala Gln Ala Ile Lys Val His Thr Ala Ala Met
660 665 670

Lys Val Gly Leu Arg Ile Val Tyr Gly Asn Thr Thr Ser Phe Leu Asp
675 680 685

Val Tyr Val Asn Gly Val Thr Pro Gly Thr Ser Lys Asp Leu Lys Val
690 695 700

Ile Ala Gly Pro Ile Ser Ala Ser Phe Thr Pro Phe Asp His Lys Val
705 710 715 720

Val Ile His Arg Gly Leu Val Tyr Asn Tyr Asp Phe Pro Glu Tyr Gly
725 730 735

Ala Met Lys Pro Gly Ala Phe Gly Asp Ile Gln Ala Thr Ser Leu Thr
740 745 750

Ser Lys Asp Leu Ile Ala Ser Thr Asp Ile Arg Leu Leu Lys Pro Ser
755 760 765

Ala Lys Asn Val His Val Pro Tyr Thr Gln Ala Ser Ser Gly Phe Glu
770 775 780

Met Trp Lys Asn Asn Ser Gly Arg Pro Leu Gln Glu Thr Ala Pro Phe
785 790 795 800

Gly Cys Lys Ile Ala Val Asn Pro Leu Arg Ala Val Asp Cys Ser Tyr
805 810 815

Gly Asn Ile Pro Ile Ser Ile Asp Ile Pro Asn Ala Ala Phe Ile Arg
820 825 830

Thr Ser Asp Ala Pro Leu Val Ser Thr Val Lys Cys Glu Val Ser Glu
835 840 845

50189A_SeqListing

Cys Thr Tyr Ser Ala Asp Phe Gly Gly Met Ala Thr Leu Gln Tyr Val
850 855 860

Ser Asp Arg Glu Gly Gln Cys Pro Val His Ser His Ser Ser Thr Ala
865 870 875 880

Thr Leu Gln Glu Ser Thr Val His Val Leu Glu Lys Gly Ala Val Thr
885 890 895

Val His Phe Ser Thr Ala Ser Pro Gln Ala Asn Phe Ile Val Ser Leu
900 905 910

Cys Gly Lys Lys Thr Thr Cys Asn Ala Glu Cys Lys Pro Pro Ala Asp
915 920 925

His Ile Val Ser Thr Pro His Lys Asn Asp Gln Glu Phe Gln Ala Ala
930 935 940

Ile Ser Lys Thr Ser Trp Ser Trp Leu Phe Ala Leu Phe Gly Gly Ala
945 950 955 960

Ser Ser Leu Leu Ile Ile Gly Leu Met Ile Phe Ala Cys Ser Met Met
965 970 975

Leu Thr Ser Thr Arg Arg
980

<210> 18
<211> 423
<212> PRT
<213> Sindbis virus

<400> 18

Ser Val Ile Asp Asp Phe Thr Leu Thr Ser Pro Tyr Leu Gly Thr Cys
1 5 10 15

50189A_SeqListing

Ser Tyr Cys His His Thr Val Pro Cys Phe Ser Pro Val Lys Ile Glu
20 25 30

Gln Val Trp Asp Glu Ala Asp Asp Asn Thr Ile Arg Ile Gln Thr Ser
35 40 45

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

Arg Tyr Met Ser Leu Lys Gln Asp His Thr Val Lys Glu Gly Thr Met
65 70 75 80

Asp Asp Ile Lys Ile Ser Thr Ser Gly Pro Cys Arg Arg Leu Ser Tyr
85 90 95

Lys Gly Tyr Phe Leu Leu Ala Lys Cys Pro Pro Gly Asp Ser Val Thr
100 105 110

Val Ser Ile Val Ser Ser Asn Ser Ala Thr Ser Cys Thr Leu Ala Arg
115 120 125

Lys Ile Lys Pro Lys Phe Val Gly Arg Glu Lys Tyr Asp Leu Pro Pro
130 135 140

Val His Gly Lys Lys Ile Pro Cys Thr Val Tyr Asp Arg Leu Lys Glu
145 150 155 160

Thr Thr Ala Gly Tyr Ile Thr Met His Arg Pro Arg Pro His Ala Tyr
165 170 175

Thr Ser Tyr Leu Glu Glu Ser Ser Gly Lys Val Tyr Ala Lys Pro Pro
180 185 190

Ser Gly Lys Asn Ile Thr Tyr Glu Cys Lys Cys Gly Asp Tyr Lys Thr
195 200 205

50189A_SeqListing

Gly Thr Val Ser Thr Arg Thr Glu Ile Thr Gly Cys Thr Ala Ile Lys
 210 215 220

Gln Cys Val Ala Tyr Lys Ser Asp Gln Thr Lys Trp Val Phe Asn Ser
 225 230 235 240

Pro Asp Leu Ile Arg His Asp Asp His Thr Ala Gln Gly Lys Leu His
 245 250 255

Leu Pro Phe Lys Leu Ile Pro Ser Thr Cys Met Val Pro Val Ala His
 260 265 270

Ala Pro Asn Val Ile His Gly Phe Lys His Ile Ser Leu Gln Leu Asp
 275 280 285

Thr Asp His Leu Thr Leu Leu Thr Thr Arg Arg Leu Gly Ala Asn Pro
 290 295 300

Glu Pro Thr Thr Glu Trp Ile Val Gly Lys Thr Val Arg Asn Phe Thr
 305 310 315 320

Val Asp Arg Asp Gly Leu Glu Tyr Ile Trp Gly Asn His Glu Pro Val
 325 330 335

Arg Val Tyr Ala Gln Glu Ser Ala Pro Gly Asp Pro His Gly Trp Pro
 340 345 350

His Glu Ile Val Gln His Tyr Tyr His Arg His Pro Val Tyr Thr Ile
 355 360 365

Leu Ala Val Ala Ser Ala Thr Val Ala Met Met Ile Gly Val Thr Val
 370 375 380

Ala Val Leu Cys Ala Cys Lys Ala Arg Arg Glu Cys Leu Thr Pro Tyr
 385 390 395 400

50189A_SeqListing

Ala Leu Ala Pro Asn Ala Val Ile Pro Thr Ser Leu Ala Leu Leu Cys
 405 410 415

Cys Val Arg Ser Ala Asn Ala
 420

<210> 19
 <211> 65
 <212> PRT
 <213> Sindbis virus

<400> 19

Met Ser Ala Ala Pro Leu Val Thr Ala Met Cys Leu Leu Gly Asn Val
 1 5 10 15

Ser Phe Pro Cys Asp Arg Pro Pro Thr Cys Tyr Thr Arg Glu Pro Ser
 20 25 30

Arg Ala Leu Asp Ile Leu Glu Glu Asn Val Asn His Glu Ala Tyr Asp
 35 40 45

Thr Leu Leu Asn Ala Ile Leu Arg Cys Gly Ser Ser Gly Arg Ser Lys
 50 55 60

Arg
 65

<210> 20
 <211> 488
 <212> PRT
 <213> Sindbis virus

<400> 20

Met Ser Ala Ala Pro Leu Val Thr Ala Met Cys Leu Leu Gly Asn Val
 1 5 10 15

Ser Phe Pro Cys Asp Arg Pro Pro Thr Cys Tyr Thr Arg Glu Pro Ser

50189A_SeqListing

20

25

30

Arg Ala Leu Asp Ile Leu Glu Glu Asn Val Asn His Glu Ala Tyr Asp
 35 40 45

Thr Leu Leu Asn Ala Ile Leu Arg Cys Gly Ser Ser Gly Arg Ser Lys
 50 55 60

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

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

Glu Gln Val Trp Asp Glu Ala Asp Asp Asn Thr Ile Arg Ile Gln Thr
 100 105 110

Ser Ala Gln Phe Gly Tyr Asp Gln Ser Gly Ala Ala Ser Ala Asn Lys
 115 120 125

Tyr Arg Tyr Met Ser Leu Lys Gln Asp His Thr Val Lys Glu Gly Thr
 130 135 140

Met Asp Asp Ile Lys Ile Ser Thr Ser Gly Pro Cys Arg Arg Leu Ser
 145 150 155 160

Tyr Lys Gly Tyr Phe Leu Leu Ala Lys Cys Pro Pro Gly Asp Ser Val
 165 170 175

Thr Val Ser Ile Val Ser Ser Asn Ser Ala Thr Ser Cys Thr Leu Ala
 180 185 190

Arg Lys Ile Lys Pro Lys Phe Val Gly Arg Glu Lys Tyr Asp Leu Pro
 195 200 205

Pro Val His Gly Lys Lys Ile Pro Cys Thr Val Tyr Asp Arg Leu Lys

50189A_SeqListing

210

215

220

Glu Thr Thr Ala Gly Tyr Ile Thr Met His Arg Pro Arg Pro His Ala
 225 230 235 240

Tyr Thr Ser Tyr Leu Glu Glu Ser Ser Gly Lys Val Tyr Ala Lys Pro
 245 250 255

Pro Ser Gly Lys Asn Ile Thr Tyr Glu Cys Lys Cys Gly Asp Tyr Lys
 260 265 270

Thr Gly Thr Val Ser Thr Arg Thr Glu Ile Thr Gly Cys Thr Ala Ile
 275 280 285

Lys Gln Cys Val Ala Tyr Lys Ser Asp Gln Thr Lys Trp Val Phe Asn
 290 295 300

Ser Pro Asp Leu Ile Arg His Asp Asp His Thr Ala Gln Gly Lys Leu
 305 310 315 320

His Leu Pro Phe Lys Leu Ile Pro Ser Thr Cys Met Val Pro Val Ala
 325 330 335

His Ala Pro Asn Val Ile His Gly Phe Lys His Ile Ser Leu Gln Leu
 340 345 350

Asp Thr Asp His Leu Thr Leu Leu Thr Thr Arg Arg Leu Gly Ala Asn
 355 360 365

Pro Glu Pro Thr Thr Glu Trp Ile Val Gly Lys Thr Val Arg Asn Phe
 370 375 380

Thr Val Asp Arg Asp Gly Leu Glu Tyr Ile Trp Gly Asn His Glu Pro
 385 390 395 400

Val Arg Val Tyr Ala Gln Glu Ser Ala Pro Gly Asp Pro His Gly Trp

50189A_SeqListing

405

410

415

Pro His Glu Ile Val Gln His Tyr Tyr His Arg His Pro Val Tyr Thr
 420 425 430

Ile Leu Ala Val Ala Ser Ala Thr Val Ala Met Met Ile Gly Val Thr
 435 440 445

Val Ala Val Leu Cys Ala Cys Lys Ala Arg Arg Glu Cys Leu Thr Pro
 450 455 460

Tyr Ala Leu Ala Pro Asn Ala Val Ile Pro Thr Ser Leu Ala Leu Leu
 465 470 475 480

Cys Cys Val Arg Ser Ala Asn Ala
 485

<210> 21

<211> 683

<212> DNA

<213> Human immunodeficiency virus type 1

<400> 21

cctagaaaaa catggagcaa tcacaagtag caatacagca gctaccaatg ctgattgtgc 60
 ctggctagaa gcacaagagg aggaggaggt gggttttcca gtcacacctc aggtaccttt 120
 aagaccaatg acttacaagg cagctgtaga tcttagccac tttttaaaag aaaagggggg 180
 actggaaggg ctaattcact cccaacgaag acaagatatc cttgatctgt ggatctacca 240
 cacacaaggc tacttccctg attggcagaa ctacacacca gggccaggga tcagatatcc 300
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 caatgaagga gagaacaccc gcttggttaca ccctgtgagc ctgcatggga tggatgaccc 420
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