

SEQUENCE LISTING

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<130> N406714W0

<150> GB1600380.8

<151> 2016-01-08

<150> GB1600381.6

<151> 2016-01-08

<150> GB1600382.4

<151> 2016-01-08

<160> 45

<170> PatentIn version 3.5

<210> 1

<211> 426

<212> DNA

<213> Mus musculus

<400> 1

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<212> DNA

<213> Mus musculus

<400> 2

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 <213> Homo sapiens

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<211> 141
<212> PRT
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<400> 5

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Val Glu Ala Ile Lys Glu Ala Leu Asn Leu Leu Asp Asp Met Pro Val
35 40 45

Thr Leu Asn Glu Glu Val Glu Val Val Ser Asn Glu Phe Ser Phe Lys
50 55 60

Lys Leu Thr Cys Val Gln Thr Arg Leu Lys Ile Phe Glu Gln Gly Leu
65 70 75 80

Arg Gly Asn Phe Thr Lys Leu Lys Gly Ala Leu Asn Met Thr Ala Ser
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Tyr Tyr Gln Thr Tyr Cys Pro Pro Thr Pro Glu Thr Asp Cys Glu Thr
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Gln Val Thr Thr Tyr Ala Asp Phe Ile Asp Ser Leu Lys Thr Phe Leu
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Thr Asp Ile Pro Phe Glu Cys Lys Lys Pro Val Gln Lys
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<210> 6
<211> 144
<212> PRT
<213> Homo sapiens

<400> 6

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 35 40 45

Thr Ala Ala Glu Met Asn Glu Thr Val Glu Val Ile Ser Glu Met Phe
 50 55 60

Asp Leu Gln Glu Pro Thr Cys Leu Gln Thr Arg Leu Glu Leu Tyr Lys
 65 70 75 80

Gln Gly Leu Arg Gly Ser Leu Thr Lys Leu Lys Gly Pro Leu Thr Met
 85 90 95

Met Ala Ser His Tyr Lys Gln His Cys Pro Pro Thr Pro Glu Thr Ser
 100 105 110

Cys Ala Thr Gln Ile Ile Thr Phe Glu Ser Phe Lys Glu Asn Leu Lys
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 130 135 140

<210> 7
 <211> 2010
 <212> DNA
 <213> Gibbon leukemia virus

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2010

<210> 8

<211> 2013

<212> DNA

<213> Gibbon leukemia virus

<400> 8

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ggaggcacct ccctgcagaa caagaatcct caccagccaa tgaccctgac atggcaggtg 180

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gcctacaagc agatcacctg gggcgccatc ggctgtagct atcctcgggc ccgcacaagg 420

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atcaatagct ccggcgacca ccagtacctg ctgccaagca accactcctg gtgggcatgc 1320

Trp Asp Ile Pro Gly Thr Asp Val Ser Ser Ser Lys Arg Val Arg Pro
100 105 110

Pro Asp Ser Asp Tyr Thr Ala Ala Tyr Lys Gln Ile Thr Trp Gly Ala
115 120 125

Ile Gly Cys Ser Tyr Pro Arg Ala Arg Thr Arg Met Ala Ser Ser Thr
130 135 140

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

Cys Gly Gly Leu Glu Ser Leu Tyr Cys Lys Glu Trp Asp Cys Glu Thr
165 170 175

Thr Gly Thr Gly Tyr Trp Leu Ser Lys Ser Ser Lys Asp Leu Ile Thr
180 185 190

Val Lys Trp Asp Gln Asn Ser Glu Trp Thr Gln Lys Phe Gln Gln Cys
195 200 205

His Gln Thr Gly Trp Cys Asn Pro Leu Lys Ile Asp Phe Thr Asp Lys
210 215 220

Gly Lys Leu Ser Lys Asp Trp Ile Thr Gly Lys Thr Trp Gly Leu Arg
225 230 235 240

Phe Tyr Val Ser Gly His Pro Gly Val Gln Phe Thr Ile Arg Leu Lys
245 250 255

Ile Thr Asn Met Pro Ala Val Ala Val Gly Pro Asp Leu Val Leu Val
260 265 270

Glu Gln Gly Pro Pro Arg Thr Ser Leu Ala Leu Pro Pro Pro Leu Pro
275 280 285

Pro Arg Glu Ala Pro Pro Pro Ser Leu Pro Asp Ser Asn Ser Thr Ala
290 295 300

Leu Ala Thr Ser Ala Gln Thr Pro Thr Val Arg Lys Thr Ile Val Thr
305 310 315 320

Leu Asn Thr Pro Pro Pro Thr Thr Gly Asp Arg Leu Phe Asp Leu Val
325 330 335

Gln Gly Ala Phe Leu Thr Leu Asn Ala Thr Asn Pro Gly Ala Thr Glu
340 345 350

Ser Cys Trp Leu Cys Leu Ala Met Gly Pro Pro Tyr Tyr Glu Ala Ile
355 360 365

Ala Ser Ser Gly Glu Val Ala Tyr Ser Thr Asp Leu Asp Arg Cys Arg
370 375 380

Trp Gly Thr Gln Gly Lys Leu Thr Leu Thr Glu Val Ser Gly His Gly
385 390 395 400

Leu Cys Ile Gly Lys Val Pro Phe Thr His Gln His Leu Cys Asn Gln
405 410 415

Thr Leu Ser Ile Asn Ser Ser Gly Asp His Gln Tyr Leu Leu Pro Ser
420 425 430

Asn His Ser Trp Trp Ala Cys Ser Thr Gly Leu Thr Pro Cys Leu Ser
435 440 445

Thr Ser Val Phe Asn Gln Thr Arg Asp Phe Cys Ile Gln Val Gln Leu
450 455 460

Ile Pro Arg Ile Tyr Tyr Tyr Pro Glu Glu Val Leu Leu Gln Ala Tyr
465 470 475 480

Asp Asn Ser His Pro Arg Thr Lys Arg Glu Ala Val Ser Leu Thr Leu
485 490 495

Ala Val Leu Leu Gly Leu Gly Ile Thr Ala Gly Ile Gly Thr Gly Ser
500 505 510

Thr Ala Leu Ile Lys Gly Pro Ile Asp Leu Gln Gln Gly Leu Thr Ser
515 520 525

Leu Gln Ile Ala Ile Asp Ala Asp Leu Arg Ala Leu Gln Asp Ser Val
530 535 540

Ser Lys Leu Glu Asp Ser Leu Thr Ser Leu Ser Glu Val Val Leu Gln
545 550 555 560

Asn Arg Arg Gly Leu Asp Leu Leu Phe Leu Lys Glu Gly Gly Leu Cys
565 570 575

Ala Ala Leu Lys Glu Glu Cys Cys Phe Tyr Ile Asp His Ser Gly Ala
580 585 590

Val Arg Asp Ser Met Lys Lys Leu Lys Glu Lys Leu Asp Lys Arg Gln
595 600 605

Leu Glu Arg Gln Lys Ser Gln Asn Trp Tyr Glu Gly Trp Phe Asn Asn
610 615 620

Ser Pro Trp Phe Thr Thr Leu Leu Ser Thr Ile Ala Gly Pro Leu Leu
625 630 635 640

Leu Leu Leu Leu Leu Leu Ile Leu Gly Pro Cys Ile Ile Asn Lys Leu
645 650 655

Val Gln Phe Ile Asn Asp Arg Ile Ser Ala Val Lys Ile
660 665

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<211> 759
<212> DNA
<213> Artificial Sequence

<220>
<223> Homo sapiens

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<210> 11
 <211> 252
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Homo sapiens

<400> 11

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Ile Thr Gln Met Ile Gly Ser Ala Leu Phe Ala Val Tyr Leu His Arg
 35 40 45

Arg Leu Asp Lys Ile Glu Asp Glu Arg Asn Leu His Glu Asp Phe Val
 50 55 60

Phe Met Lys Thr Ile Gln Arg Cys Asn Thr Gly Glu Arg Ser Leu Ser
 65 70 75 80

Leu Leu Asn Cys Glu Glu Ile Lys Ser Gln Phe Glu Gly Phe Val Lys
 85 90 95

Asp Ile Met Leu Asn Lys Glu Glu Thr Lys Lys Asp Glu Asp Pro Gln
100 105 110

Ile Ala Ala His Val Val Ser Glu Ala Asn Ser Asn Ala Ala Ser Val
115 120 125

Leu Gln Trp Ala Lys Lys Gly Tyr Tyr Thr Met Lys Ser Asn Leu Val
130 135 140

Thr Leu Glu Asn Gly Lys Gln Leu Thr Val Lys Arg Gln Gly Leu Tyr
145 150 155 160

Tyr Ile Tyr Ala Gln Val Thr Phe Cys Ser Asn Arg Glu Ala Ser Ser
165 170 175

Gln Ala Pro Phe Ile Val Gly Leu Trp Leu Lys Pro Ser Ser Gly Ser
180 185 190

Glu Arg Ile Leu Leu Lys Ala Ala Asn Thr His Ser Ser Ser Gln Leu
195 200 205

Cys Glu Gln Gln Ser Ile His Leu Gly Gly Val Phe Glu Leu Gln Pro
210 215 220

Gly Ala Ser Val Phe Val Asn Val Thr Asp Pro Ser Gln Val Ser His
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Gly Thr Gly Phe Thr Ser Phe Gly Leu Leu Lys Leu
245 250

<210> 12
<211> 1416
<212> DNA
<213> Homo sapiens

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<210> 13
 <211> 471
 <212> PRT
 <213> Homo sapiens

<400> 13

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Thr Cys Thr Leu Val Met Cys Ser Pro Thr Glu Asn Gly Leu Pro Gly
35 40 45

Arg Asp Gly Arg Asp Gly Arg Glu Gly Pro Arg Gly Glu Lys Gly Asp
50 55 60

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

Gly Pro Val Gly Pro Lys Gly Glu Asn Gly Ser Ala Gly Glu Pro Gly
85 90 95

Pro Lys Gly Glu Arg Gly Leu Ser Gly Pro Pro Gly Leu Pro Gly Ile
100 105 110

Pro Gly Pro Ala Gly Lys Glu Gly Pro Ser Gly Lys Gln Gly Asn Ile
115 120 125

Gly Pro Gln Gly Lys Pro Gly Pro Lys Gly Glu Ala Gly Pro Lys Gly
130 135 140

Glu Val Gly Ala Pro Gly Met Gln Gly Ser Thr Gly Ala Lys Gly Ser
145 150 155 160

Thr Gly Pro Lys Gly Glu Arg Gly Ala Pro Gly Val Gln Gly Ala Pro
165 170 175

Gly Asn Ala Gly Ala Ala Gly Pro Ala Gly Pro Ala Gly Pro Gln Gly
180 185 190

Ala Pro Gly Ser Arg Gly Pro Pro Gly Leu Lys Gly Asp Arg Gly Val
195 200 205

Pro Gly Asp Arg Gly Ile Lys Gly Glu Ser Gly Leu Pro Asp Ser Ala
210 215 220

Ala Leu Arg Gln Gln Met Glu Ala Leu Lys Gly Lys Leu Gln Arg Leu
225 230 235 240

Glu Val Ala Phe Ser His Tyr Gln Lys Ala Ala Leu Phe Pro Asp Gly
245 250 255

His Arg Arg Leu Asp Lys Ile Glu Asp Glu Arg Asn Leu His Glu Asp
260 265 270

Phe Val Phe Met Lys Thr Ile Gln Arg Cys Asn Thr Gly Glu Arg Ser
275 280 285

Leu Ser Leu Leu Asn Cys Glu Glu Ile Lys Ser Gln Phe Glu Gly Phe
290 295 300

Val Lys Asp Ile Met Leu Asn Lys Glu Glu Thr Lys Lys Glu Asn Ser
305 310 315 320

Phe Glu Met Gln Lys Gly Asp Gln Asn Pro Gln Ile Ala Ala His Val
325 330 335

Ile Ser Glu Ala Ser Ser Lys Thr Thr Ser Val Leu Gln Trp Ala Glu
340 345 350

Lys Gly Tyr Tyr Thr Met Ser Asn Asn Leu Val Thr Leu Glu Asn Gly
355 360 365

Lys Gln Leu Thr Val Lys Arg Gln Gly Leu Tyr Tyr Ile Tyr Ala Gln
370 375 380

Val Thr Phe Cys Ser Asn Arg Glu Ala Ser Ser Gln Ala Pro Phe Ile
385 390 395 400

Ala Ser Leu Cys Leu Lys Ser Pro Gly Arg Phe Glu Arg Ile Leu Leu
405 410 415

Arg Ala Ala Asn Thr His Ser Ser Ala Lys Pro Cys Gly Gln Gln Ser
420 425 430

Ile His Leu Gly Gly Val Phe Glu Leu Gln Pro Gly Ala Ser Val Phe
435 440 445

Val Asn Val Thr Asp Pro Ser Gln Val Ser His Gly Thr Gly Phe Thr
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Ser Phe Gly Leu Leu Lys Leu
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<210> 14
<211> 1412
<212> DNA
<213> Mus musculus

<400> 14
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 <211> 470
 <212> PRT
 <213> Mus musculus

<400> 15

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			20					25					30		

Thr	Cys	Thr	Leu	Val	Met	Cys	Ser	Pro	Thr	Glu	Asn	Gly	Leu	Pro	Gly
		35					40					45			

Arg	Asp	Gly	Arg	Asp	Gly	Arg	Glu	Gly	Pro	Arg	Gly	Glu	Lys	Gly	Asp
	50					55					60				

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

Gly	Pro	Val	Gly	Pro	Lys	Gly	Glu	Asn	Gly	Ser	Ala	Gly	Glu	Pro	Gly
				85					90					95	

Pro	Lys	Gly	Glu	Arg	Gly	Leu	Ser	Gly	Pro	Pro	Gly	Leu	Pro	Gly	Ile
			100					105					110		

Pro	Gly	Pro	Ala	Gly	Lys	Glu	Gly	Pro	Ser	Gly	Lys	Gln	Gly	Asn	Ile
		115					120					125			

Gly	Pro	Gln	Gly	Lys	Pro	Gly	Pro	Lys	Gly	Glu	Ala	Gly	Pro	Lys	Gly
	130					135					140				

Glu	Val	Gly	Ala	Pro	Gly	Met	Gln	Gly	Ser	Thr	Gly	Ala	Lys	Gly	Ser
145					150					155					160

Thr Gly Pro Lys Gly Glu Arg Gly Ala Pro Gly Val Gln Gly Ala Pro
165 170 175

Gly Asn Ala Gly Ala Ala Gly Pro Ala Gly Pro Ala Gly Pro Gln Gly
180 185 190

Ala Pro Gly Ser Arg Gly Pro Pro Gly Leu Lys Gly Asp Arg Gly Val
195 200 205

Pro Gly Asp Arg Gly Ile Lys Gly Glu Ser Gly Leu Pro Asp Ser Ala
210 215 220

Ala Leu Arg Gln Gln Met Glu Ala Leu Lys Gly Lys Leu Gln Arg Leu
225 230 235 240

Glu Val Ala Phe Ser His Tyr Gln Lys Ala Ala Leu Phe Pro Asp Gly
245 250 255

His Arg Arg Leu Asp Lys Val Glu Glu Glu Val Asn Leu His Glu Asp
260 265 270

Phe Val Phe Ile Lys Lys Leu Lys Arg Cys Asn Lys Gly Glu Gly Ser
275 280 285

Leu Ser Leu Leu Asn Cys Glu Glu Met Arg Arg Gln Phe Glu Asp Leu
290 295 300

Val Lys Asp Ile Thr Leu Asn Lys Glu Glu Lys Lys Glu Asn Ser Phe
305 310 315 320

Glu Met Gln Arg Gly Asp Glu Asp Pro Gln Ile Ala Ala His Val Val
325 330 335

Ser Glu Ala Asn Ser Asn Ala Ala Ser Val Leu Gln Trp Ala Lys Lys
340 345 350

Gly Tyr Tyr Thr Met Lys Ser Asn Leu Val Met Leu Glu Asn Gly Lys
355 360 365

Gln Leu Thr Val Lys Arg Glu Gly Leu Tyr Tyr Val Tyr Thr Gln Val
370 375 380

Thr Phe Cys Ser Asn Arg Glu Pro Ser Ser Gln Arg Pro Phe Ile Val
 385 390 395 400

Gly Leu Trp Leu Lys Pro Ser Ile Gly Ser Glu Arg Ile Leu Leu Lys
 405 410 415

Ala Ala Asn Thr His Ser Ser Ser Gln Leu Cys Glu Gln Gln Ser Val
 420 425 430

His Leu Gly Gly Val Phe Glu Leu Gln Ala Gly Ala Ser Val Phe Val
 435 440 445

Asn Val Thr Glu Ala Ser Gln Val Ile His Arg Val Gly Phe Ser Ser
 450 455 460

Phe Gly Leu Leu Lys Leu
 465 470

<210> 16
 <211> 786
 <212> DNA
 <213> Homo sapiens

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 ctttttgctg tgtatcttca tagaagggtg gacaagatag aagatgaaag gaatcttcat 180
 gaagattttg tattcatgaa aacgatacag agatgcaaca caggagaaag atccttatcc 240
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caatccattc acttgggagg agtatttgaa ttgcaaccag gtgcttcggt gtttgtcaat 720
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<210> 17
 <211> 261
 <212> PRT
 <213> Homo sapiens

<400> 17

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Ile Thr Gln Met Ile Gly Ser Ala Leu Phe Ala Val Tyr Leu His Arg
 35 40 45

Arg Leu Asp Lys Ile Glu Asp Glu Arg Asn Leu His Glu Asp Phe Val
 50 55 60

Phe Met Lys Thr Ile Gln Arg Cys Asn Thr Gly Glu Arg Ser Leu Ser
 65 70 75 80

Leu Leu Asn Cys Glu Glu Ile Lys Ser Gln Phe Glu Gly Phe Val Lys
 85 90 95

Asp Ile Met Leu Asn Lys Glu Glu Thr Lys Lys Glu Asn Ser Phe Glu
 100 105 110

Met Gln Lys Gly Asp Gln Asn Pro Gln Ile Ala Ala His Val Ile Ser
 115 120 125

Glu Ala Ser Ser Lys Thr Thr Ser Val Leu Gln Trp Ala Glu Lys Gly
 130 135 140

Tyr Tyr Thr Met Ser Asn Asn Leu Val Thr Leu Glu Asn Gly Lys Gln
 145 150 155 160

Leu Thr Val Lys Arg Gln Gly Leu Tyr Tyr Ile Tyr Ala Gln Val Thr
165 170 175

Phe Cys Ser Asn Arg Glu Ala Ser Ser Gln Ala Pro Phe Ile Ala Ser
180 185 190

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

Ala Asn Thr His Ser Ser Ala Lys Pro Cys Gly Gln Gln Ser Ile His
210 215 220

Leu Gly Gly Val Phe Glu Leu Gln Pro Gly Ala Ser Val Phe Val Asn
225 230 235 240

Val Thr Asp Pro Ser Gln Val Ser His Gly Thr Gly Phe Thr Ser Phe
245 250 255

Gly Leu Leu Lys Leu
260

<210> 18
<211> 783
<212> DNA
<213> Mus musculus

<400> 18
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ctttttgctg tgtatcttca tagaagattg gataaggtcg aagaggaagt aaaccttcat 180
gaagattttg tattcataaa aaagctaaag agatgcaaca aaggagaagg atctttatcc 240
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aacaaagaag agaaaaaaga aaacagcttt gaaatgcaaa gaggtgatga ggatcctcaa 360
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gagagaatct tactcaaggc ggcaaatacc cacagttcct cccagctttg cgagcagcag	660
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tga	783

<210> 19
 <211> 260
 <212> PRT
 <213> Mus musculus

<400> 19

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			20					25					30		

Ile	Thr	Gln	Met	Ile	Gly	Ser	Val	Leu	Phe	Ala	Val	Tyr	Leu	His	Arg
		35					40					45			

Arg	Leu	Asp	Lys	Val	Glu	Glu	Glu	Val	Asn	Leu	His	Glu	Asp	Phe	Val
	50					55					60				

Phe	Ile	Lys	Lys	Leu	Lys	Arg	Cys	Asn	Lys	Gly	Glu	Gly	Ser	Leu	Ser
65					70					75					80

Leu	Leu	Asn	Cys	Glu	Glu	Met	Arg	Arg	Gln	Phe	Glu	Asp	Leu	Val	Lys
				85					90					95	

Asp	Ile	Thr	Leu	Asn	Lys	Glu	Glu	Lys	Lys	Glu	Asn	Ser	Phe	Glu	Met
			100					105					110		

Gln	Arg	Gly	Asp	Glu	Asp	Pro	Gln	Ile	Ala	Ala	His	Val	Val	Ser	Glu
		115					120					125			

Ala	Asn	Ser	Asn	Ala	Ala	Ser	Val	Leu	Gln	Trp	Ala	Lys	Lys	Gly	Tyr
	130					135					140				

Tyr	Thr	Met	Lys	Ser	Asn	Leu	Val	Met	Leu	Glu	Asn	Gly	Lys	Gln	Leu
145					150					155					160

Thr Val Lys Arg Glu Gly Leu Tyr Tyr Val Tyr Thr Gln Val Thr Phe
165 170 175

Cys Ser Asn Arg Glu Pro Ser Ser Gln Arg Pro Phe Ile Val Gly Leu
180 185 190

Trp Leu Lys Pro Ser Ser Gly Ser Glu Arg Ile Leu Leu Lys Ala Ala
195 200 205

Asn Thr His Ser Ser Ser Gln Leu Cys Glu Gln Gln Ser Val His Leu
210 215 220

Gly Gly Val Phe Glu Leu Gln Ala Gly Ala Ser Val Phe Val Asn Val
225 230 235 240

Thr Glu Ala Ser Gln Val Ile His Arg Val Gly Phe Ser Ser Phe Gly
245 250 255

Leu Leu Lys Leu
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<210> 20
<211> 930
<212> DNA
<213> Mus musculus

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<210> 21
 <211> 765
 <212> DNA
 <213> Homo sapiens

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 <212> DNA
 <213> Mus musculus

<400> 22

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tgggagtga	1389

<210> 23
 <211> 1389
 <212> DNA
 <213> Homo sapiens

<400> 23

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<210> 24
 <211> 522
 <212> DNA
 <213> Mus musculus

<400> 24
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<210> 25
<211> 600
<212> DNA
<213> Homo sapiens

<400> 25
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<210> 26
<211> 1164
<212> DNA
<213> Mus musculus

<400> 26

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ccagatctgc cttttatcag ctga	1164

<210> 27
 <211> 1152
 <212> DNA
 <213> Homo sapiens

<400> 27	
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agcagcgtgg agagcggcct gccaggaagg gacggaagag atggaaggga gggaccacga	180
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 <211> 552
 <212> DNA
 <213> Homo sapiens

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 <212> DNA
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 <212> DNA
 <213> Homo sapiens

<400> 33	
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gaagggtccc ggttcgacct gaacgatgtc tacgtgtatt ggacagacctc tgagagtaag	180
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 <213> Mus musculus

<400> 34

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<211> 1484
<212> DNA
<213> Homo sapiens

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acacagaagt ctctgagtct gtcacccggc aaatgaggcg cgcc 1484

<210> 36
<211> 632
<212> DNA
<213> Artificial Sequence

<220>
<223> CMV promoter

<400> 36
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ccaacgaccc ccgcccattg acgtcaataa tgacgtatgt tcccatagta acgccaatag 180
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<210> 37
<211> 394
<212> DNA
<213> Artificial Sequence

<220>
<223> RSV promoter

<400> 37
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<210> 38
 <211> 188
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> BGH polyA

<400> 38	
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tgagtaggtg tcattctatt ctgggggggtg ggggtggggca ggacagcaag ggggaggatt	180
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<210> 39
 <211> 249
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> SV40 late polyA

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ctttatttgt aaccattata agctgcaata aacaagttaa caacaacaat tgcattcatt	180
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aatgtggta	249

<210> 40
 <211> 345
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> SV40 enhancer promoter

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<210> 41
 <211> 99
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Rabbit beta-globin polyA

<400> 41	
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<210> 42
 <211> 723
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> GFP

<400> 42	
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<210> 43
 <211> 454
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> MoMuLV LTR

<400> 43	
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<210> 44
 <211> 1349
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> EF1alpha promoter

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<220>
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