

SP102075SGM SEQUENCE LISTING

<110> YOO YOUNG PHARM CO.,LTD.
 <120> CHIMERIC ANTIGEN RECEPTOR AND T CELLS EXPRESSING THE CHIMERIC ANTIGEN RECEPTOR
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 <150> 10-2015-0110788
 <151> 2015-08-05
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 35 40 45
 Ser Leu Ile Asn Val Ser Gly Cys Ser Ala Ile Glu Lys Thr Gln Arg
 50 55 60
 Met Leu Ser Gly Phe Cys Pro His Lys Val Ser Ala Gly Gln Phe Ser
 65 70 75 80
 Ser Leu His Val Arg Asp Thr Lys Ile Glu Val Ala Gln Phe Val Lys
 85 90 95
 Asp Leu Leu Leu His Leu Lys Lys Leu Phe Arg Glu Gly Gln Phe Asn
 100 105 110

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 Pro His Phe Pro Thr Arg Tyr Tyr Arg Ile Thr Tyr Gly Glu Thr Gly
 20 25 30
 Gly Asn Ser Pro Val Gln Glu Phe Thr Val Leu Gln Pro Pro Ser Thr
 35 40 45
 Ala Thr Ile Ser Gly Leu Lys Pro Gly Val Asp Tyr Thr Ile Thr Val
 Page 1

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50 55 60

Tyr Ala Val Val Glu Arg Asn Gly Arg Glu Leu Asn Thr Pro Pro Ile
65 70 75 80

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

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

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

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

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

Gly Thr Met Val Thr Val Ser Ser Gly Ser Thr Ser Gly Ser Gly Lys
115 120 125

Pro Gly Ser Gly Glu Gly Ser Asp Ile Gln Met Thr Gln Ser Pro Ser
130 135 140

Ser Leu Ser Ala Ser Val Gly Asp Arg Val Thr Ile Thr Cys Met Thr
145 150 155 160

Ser Thr Asp Ile Asp Asp Asp Met Asn Trp Tyr Gln Gln Lys Pro Gly
165 170 175

Lys Thr Pro Lys Leu Leu Ile Tyr Glu Gly Asn Thr Leu Arg Pro Gly
180 185 190

Val Pro Ser Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Ile Phe
195 200 205

Thr Ile Ser Ser Leu Gln Pro Glu Asp Ile Ala Thr Tyr Tyr Cys Leu
210 215 220

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

Ile Lys Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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 Cys Pro His Tyr Glu Asp His Ser₄₀ Val Ala Asp Ala₄₅ Ala Met Glu Gln
 Tyr Ile₅₀ Leu Tyr Leu Val₅₅ Glu His Glu Glu Tyr Gln₆₀ Leu Cys Gln Pro
 Gln Ser Lys Asp Gln Val₇₀ Arg Trp Gln Cys Asn₇₅ Arg Pro Ser Ala Lys₈₀
 His Gly Pro Glu Lys₈₅ Leu Ser Glu Lys Phe₉₀ Gln Arg Phe Thr Ala₉₅ Phe
 Ala Leu Ala Lys₁₀₀ Glu Phe Lys Ala Gly₁₀₅ His Ser Tyr Tyr Tyr₁₁₀ Ile Ser
 Lys Pro Ile₁₁₅ His Gln His Glu Asp₁₂₀ Arg Cys Leu Arg Leu₁₂₅ Lys Val Thr
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 Arg Ile Thr Tyr Gly Glu Thr Gly₄₀ Gly Asn Ser Pro Val₄₅ Gln Glu Phe
 Thr Val₅₀ Pro Gly Ser Lys Ser₅₅ Thr Ala Thr Ile Ser₆₀ Gly Leu Lys Pro
 Gly Val Asp Tyr Thr Ile₇₀ Thr Val Tyr Ala Val₇₅ Thr Pro Arg Gly Asp₈₀
 Trp Asn Glu Gly Ser₈₅ Lys Pro Ile Ser Ile₉₀ Asn Tyr Arg Thr Glu Gln₉₅

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Glu Lys Cys Glu Lys Leu Val Ser Val Asn Phe Ile Asn Lys Glu Ser
35 40 45
Tyr Leu Gln Ile Pro Ser Ala Lys Val Arg Pro Gln Thr Asn Ile Thr
50 55 60
Leu Gln Ile Ala Thr Asp Glu Asp Ser Gly Ile Leu Leu Tyr Lys Gly
65 70 75 80
Asp Lys Asp His Ile Ala Val Glu Leu Tyr Arg Gly Arg Val Arg Ala
85 90 95
Ser Tyr Asp Thr Gly Ser His Pro Ala Ser Ala Ile Tyr Ser Val Glu
100 105 110
Thr Ile Asn Asp Gly Asn Phe His Ile Val Glu Leu Leu Ala Leu Asp
115 120 125
Gln Ser Leu Ser Leu Ser Val Asp Gly Gly Asn Pro Lys Ile Ile Thr
130 135 140
Asn Leu Ser Lys Gln Ser Thr Leu Asn Phe Asp Ser Pro Leu Tyr Val
145 150 155 160
Gly Gly Met Pro Gly Lys Ser Asn Val Ala Ser Leu Arg Gln Ala Pro
165 170 175
Gly Gln Asn Gly Thr Ser Phe His Gly Cys Ile Arg Asn Leu Tyr Ile
180 185 190
Asn Ser Glu Leu Gln Asp Phe Gln Lys Val Pro Met Gln Thr Gly Ile
195 200 205
Leu Pro Gly Cys Glu Pro Cys His Lys Lys Val Cys Ala His Gly Thr
210 215 220
Cys Gln Pro Ser Ser Gln Ala Gly Phe Thr Cys Glu Cys Gln Glu Gly
225 230 235 240
Trp Met Gly Pro Leu Cys Asp Gln Arg Thr Asn Asp Pro Cys Leu Gly
245 250 255
Asn Lys Cys Val His Gly Thr Cys Leu Pro Ile Asn Ala Phe Ser Tyr
260 265 270
Ser Cys Lys Cys Leu Glu Gly His Gly Gly Val Leu Cys Asp Glu Glu
275 280 285

SP102075SGM SEQUENCE LISTING

Glu Asp Leu Phe Asn Pro Cys Gln Ala Ile Lys Cys Lys His Gly Lys
 290 295 300
 Cys Arg Leu Ser Gly Leu Gly Gln Pro Tyr Cys Glu Cys Ser Ser Gly
 305 310 315 320
 Tyr Thr Gly Asp Ser Cys Asp Arg Glu Ile Ser Cys Arg Gly Glu Arg
 325 330 335
 Ile Arg Asp Tyr Tyr Gln Lys Gln Gln Gly Tyr Ala Ala Cys Gln Thr
 340 345 350
 Thr Lys Lys Val Ser Arg Leu Glu Cys Arg Gly Gly Cys Ala Gly Gly
 355 360 365
 Gln Cys Cys Gly Pro Leu Arg Ser Lys Arg Arg Lys Tyr Ser Phe Glu
 370 375 380
 Cys Thr Asp Gly Ser Ser Phe Val Asp Glu Val Glu Lys Val Val Lys
 385 390 395 400
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 405 410 415

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 Pro Glu Glu Glu Gly Gly Cys Glu Leu
 35 40

SP102075SGM SEQUENCE LISTING

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 35 40 45

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SP102075SGM SEQUENCE LISTING

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 35 40 45
 Pro Gln Arg Arg Lys Asn Pro Gln Glu Gly Leu Tyr Asn Glu Leu Gln
 50 55 60
 Lys Asp Lys Met Ala Glu Ala Tyr Ser Glu Ile Gly Met Lys Gly Glu
 65 70 75 80
 Arg Arg Arg Gly Lys Gly His Asp Gly Leu Tyr Gln Gly Leu Ser Thr
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 Ala Thr Lys Asp Thr Tyr Asp Ala Leu His Met Gln Ala Leu Pro Pro
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 Glu Glu Leu Val Asn Ile Thr Gln Asn Gln Lys Ala Pro Leu Cys Asn
 35 40 45
 Gly Ser Met Val Trp Ser Ile Asn Leu Thr Ala Gly Met Tyr Cys Ala
 50 55 60
 Ala Leu Glu Ser Leu Ile Asn Val Ser Gly Cys Ser Ala Ile Glu Lys
 65 70 75 80

SP102075SGM SEQUENCE LISTING

Thr Gln Asp Met Leu Asp Gly Phe Cys Pro His Lys Val Ser Ala Gly
85 90 95

Gln Phe Ser Ser Leu His Val Arg Asp Thr Lys Ile Glu Val Ala Gln
100 105 110

Phe Val Lys Asp Leu Leu Leu His Leu Lys Lys Leu Phe Lys Glu Gly
115 120 125

Gln Phe Asn Gly Gly Gly Pro Arg Lys Pro Thr Thr Thr Pro Ala Pro
130 135 140

Arg Pro Pro Thr Pro Ala Pro Thr Ile Ala Ser Gln Pro Leu Ser Leu
145 150 155 160

Arg Pro Glu Ala Cys Arg Pro Ala Ala Gly Gly Ala Val His Thr Arg
165 170 175

Gly Leu Asp Phe Ala Cys Asp Ile Tyr Ile Trp Ala Pro Leu Ala Gly
180 185 190

Thr Cys Gly Val Leu Leu Leu Ser Leu Val Ile Thr Lys Arg Gly Arg
195 200 205

Lys Lys Leu Leu Tyr Ile Phe Lys Gln Pro Phe Met Arg Pro Val Gln
210 215 220

Thr Thr Gln Glu Glu Asp Gly Cys Ser Cys Arg Phe Pro Glu Glu Glu
225 230 235 240

Glu Gly Gly Cys Glu Leu Arg Val Lys Phe Ser Arg Ser Ala Asp Ala
245 250 255

Pro Ala Tyr Gln Gln Gly Gln Asn Gln Leu Tyr Asn Glu Leu Asn Leu
260 265 270

Gly Arg Arg Glu Glu Tyr Asp Val Leu Asp Lys Arg Arg Gly Arg Asp
275 280 285

Pro Glu Met Gly Gly Lys Pro Gln Arg Arg Lys Asn Pro Gln Glu Gly
290 295 300

Leu Tyr Asn Glu Leu Gln Lys Asp Lys Met Ala Glu Ala Tyr Ser Glu
305 310 315 320

Ile Gly Met Lys Gly Glu Arg Arg Arg Gly Lys Gly His Asp Gly Leu
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20 25 30
Glu Glu Leu Val Asn Ile Thr Gln Asn Gln Lys Ala Pro Leu Cys Asn
35 40 45
Gly Ser Met Val Trp Ser Ile Asn Leu Thr Ala Gly Met Tyr Cys Ala
50 55 60
Ala Leu Glu Ser Leu Ile Asn Val Ser Gly Cys Ser Ala Ile Glu Lys
65 70 75 80
Thr Gln Asp Met Leu Asp Gly Phe Cys Pro His Lys Val Ser Ala Gly
85 90 95
Gln Phe Ser Ser Leu His Val Arg Asp Thr Lys Ile Glu Val Ala Gln
100 105 110
Phe Val Lys Asp Leu Leu Leu His Leu Lys Lys Leu Phe Lys Glu Gly
115 120 125
Gln Phe Asn Gly Gly Gly Pro Arg Lys Pro Thr Thr Thr Pro Ala Pro
130 135 140
Arg Pro Pro Thr Pro Ala Pro Thr Ile Ala Ser Gln Pro Leu Ser Leu
145 150 155 160
Arg Pro Glu Ala Cys Arg Pro Ala Ala Gly Gly Ala Val His Thr Arg
165 170 175
Gly Leu Asp Phe Ala Cys Asp Ile Tyr Ile Trp Ala Pro Leu Ala Gly
180 185 190
Thr Cys Gly Val Leu Leu Leu Ser Leu Val Ile Thr Arg Ser Lys Arg
195 200 205
Ser Arg Gly Gly His Ser Asp Tyr Met Asn Met Thr Pro Arg Arg Pro
210 215 220
Gly Pro Thr Arg Lys His Tyr Gln Pro Tyr Ala Pro Pro Arg Asp Phe
225 230 235 240
Ala Ala Tyr Arg Ser Lys Arg Gly Arg Lys Lys Leu Leu Tyr Ile Phe
245 250 255
Lys Gln Pro Phe Met Arg Pro Val Gln Thr Thr Gln Glu Glu Asp Gly
260 265 270
Cys Ser Cys Arg Phe Pro Glu Glu Glu Gly Gly Cys Glu Leu Arg
275 280 285
Val Lys Phe Ser Arg Ser Ala Asp Ala Pro Ala Tyr Gln Gln Gly Gln
290 295 300
Asn Gln Leu Tyr Asn Glu Leu Asn Leu Gly Arg Arg Glu Glu Tyr Asp
305 310 315 320
Val Leu Asp Lys Arg Arg Gly Arg Asp Pro Glu Met Gly Gly Lys Pro
325 330 335
Gln Arg Arg Lys Asn Pro Gln Glu Gly Leu Tyr Asn Glu Leu Gln Lys
340 345 350
Asp Lys Met Ala Glu Ala Tyr Ser Glu Ile Gly Met Lys Gly Glu Arg
355 360 365
Arg Arg Gly Lys Gly His Asp Gly Leu Tyr Gln Gly Leu Ser Thr Ala
370 375 380

SP102075SGM SEQUENCE LISTING

Thr Lys Asp Thr Tyr Asp Ala Leu His Met Gln Ala Leu Pro Pro Arg
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Glu Glu Leu Val Asn Ile Thr Gln Asn Gln Lys Ala Pro Leu Cys Asn
35 40 45
Gly Ser Met Val Trp Ser Ile Asn Leu Thr Ala Gly Met Tyr Cys Ala
50 55 60
Ala Leu Glu Ser Leu Ile Asn Val Ser Gly Cys Ser Ala Ile Glu Lys
65 70 75 80
Thr Gln Arg Met Leu Ser Gly Phe Cys Pro His Lys Val Ser Ala Gly
85 90 95
Gln Phe Ser Ser Leu His Val Arg Asp Thr Lys Ile Glu Val Ala Gln
100 105 110
Phe Val Lys Asp Leu Leu Leu His Leu Lys Lys Leu Phe Lys Glu Gly
115 120 125
Gln Phe Asn Gly Gly Gly Pro Arg Lys Pro Thr Thr Thr Pro Ala Pro
130 135 140
Arg Pro Pro Thr Pro Ala Pro Thr Ile Ala Ser Gln Pro Leu Ser Leu
145 150 155 160
Arg Pro Glu Ala Cys Arg Pro Ala Ala Gly Gly Ala Val His Thr Arg
165 170 175
Gly Leu Asp Phe Ala Cys Asp Ile Tyr Ile Trp Ala Pro Leu Ala Gly
180 185 190
Thr Cys Gly Val Leu Leu Leu Ser Leu Val Ile Thr Arg Ser Lys Arg
195 200 205
Ser Arg Gly Gly His Ser Asp Tyr Met Asn Met Thr Pro Arg Arg Pro
210 215 220
Gly Pro Thr Arg Lys His Tyr Gln Pro Tyr Ala Pro Pro Arg Asp Phe
225 230 235 240
Ala Ala Tyr Arg Ser Lys Arg Gly Arg Lys Lys Leu Leu Tyr Ile Phe
245 250 255
Lys Gln Pro Phe Met Arg Pro Val Gln Thr Thr Gln Glu Glu Asp Gly
260 265 270
Cys Ser Cys Arg Phe Pro Glu Glu Glu Gly Gly Cys Glu Leu Arg
275 280 285

SP102075SGM SEQUENCE LISTING

Val Lys Phe Ser Arg Ser Ala Asp Ala Pro Ala Tyr Gln Gln Gly Gln
 290 300
 Asn Gln Leu Tyr Asn Glu Leu Asn Leu Gly Arg Arg Glu Glu Tyr Asp
 305 310 315 320
 Val Leu Asp Lys Arg Arg Gly Arg Asp Pro Glu Met Gly Gly Lys Pro
 325 330 335
 Gln Arg Arg Lys Asn Pro Gln Glu Gly Leu Tyr Asn Glu Leu Gln Lys
 340 345 350
 Asp Lys Met Ala Glu Ala Tyr Ser Glu Ile Gly Met Lys Gly Glu Arg
 355 360 365
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 370 375 380
 Thr Lys Asp Thr Tyr Asp Ala Leu His Met Gln Ala Leu Pro Pro Arg
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 Trp Arg His Pro His Phe Pro Thr Arg Tyr Tyr Arg Ile Thr Tyr Gly
 35 40 45
 Glu Thr Gly Gly Asn Ser Pro Val Gln Glu Phe Thr Val Leu Gln Pro
 50 55 60
 Pro Ser Thr Ala Thr Ile Ser Gly Leu Lys Pro Gly Val Asp Tyr Thr
 65 70 75 80
 Ile Thr Val Tyr Ala Val Val Glu Arg Asn Gly Arg Glu Leu Asn Thr
 85 90 95
 Pro Pro Ile Ser Ile Asn Tyr Arg Thr His His His His His Gly
 100 105 110
 Gly Gly Pro Arg Lys Pro Thr Thr Thr Pro Ala Pro Arg Pro Pro Thr
 115 120 125
 Pro Ala Pro Thr Ile Ala Ser Gln Pro Leu Ser Leu Arg Pro Glu Ala
 130 135 140
 Cys Arg Pro Ala Ala Gly Gly Ala Val His Thr Arg Gly Leu Asp Phe
 145 150 155 160
 Ala Cys Asp Ile Tyr Ile Trp Ala Pro Leu Ala Gly Thr Cys Gly Val
 165 170 175
 Leu Leu Leu Ser Leu Val Ile Thr Lys Arg Gly Arg Lys Lys Leu Leu
 180 185 190

SP102075SGM SEQUENCE LISTING

Tyr Ile Phe Lys Gln Pro Phe Met Arg Pro Val Gln Thr Thr Gln Glu
195 200 205
Glu Asp Gly Cys Ser Cys Arg Phe Pro Glu Glu Glu Glu Gly Gly Cys
210 215 220
Glu Leu Arg Val Lys Phe Ser Arg Ser Ala Asp Ala Pro Ala Tyr Gln
225 230 235 240
Gln Gly Gln Asn Gln Leu Tyr Asn Glu Leu Asn Leu Gly Arg Arg Glu
245 250 255
Glu Tyr Asp Val Leu Asp Lys Arg Arg Gly Arg Asp Pro Glu Met Gly
260 265 270
Gly Lys Pro Gln Arg Arg Lys Asn Pro Gln Glu Gly Leu Tyr Asn Glu
275 280 285
Leu Gln Lys Asp Lys Met Ala Glu Ala Tyr Ser Glu Ile Gly Met Lys
290 295 300
Gly Glu Arg Arg Arg Gly Lys Gly His Asp Gly Leu Tyr Gln Gly Leu
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Trp Arg His Pro His Phe Pro Thr Arg Tyr Tyr Arg Ile Thr Tyr Gly
35 40 45
Glu Thr Gly Gly Asn Ser Pro Val Gln Glu Phe Thr Val Leu Gln Pro
50 55 60
Pro Ser Thr Ala Thr Ile Ser Gly Leu Lys Pro Gly Val Asp Tyr Thr
65 70 75 80
Ile Thr Val Tyr Ala Val Val Glu Arg Asn Gly Arg Glu Leu Asn Thr
85 90 95
Pro Pro Ile Ser Ile Asn Tyr Arg Thr His His His His His His Gly
100 105 110
Gly Gly Pro Arg Lys Pro Thr Thr Thr Pro Ala Pro Arg Pro Pro Thr
115 120 125
Pro Ala Pro Thr Ile Ala Ser Gln Pro Leu Ser Leu Arg Pro Glu Ala
130 135 140

SP102075SGM SEQUENCE LISTING

Cys Arg Pro Ala Ala Gly Gly Ala Val His Thr Arg Gly Leu Asp Phe
 145 150 155 160
 Ala Cys Asp Ile Tyr Ile Trp Ala Pro Leu Ala Gly Thr Cys Gly Val
 165 170 175
 Leu Leu Leu Ser Leu Val Ile Thr Arg Ser Lys Arg Ser Arg Gly Gly
 180 185 190
 His Ser Asp Tyr Met Asn Met Thr Pro Arg Arg Pro Gly Pro Thr Arg
 195 200 205
 Lys His Tyr Gln Pro Tyr Ala Pro Pro Arg Asp Phe Ala Ala Tyr Arg
 210 215 220
 Ser Lys Arg Gly Arg Lys Lys Leu Leu Tyr Ile Phe Lys Gln Pro Phe
 225 230 235 240
 Met Arg Pro Val Gln Thr Thr Gln Glu Glu Asp Gly Cys Ser Cys Arg
 245 250 255
 Phe Pro Glu Glu Glu Glu Gly Gly Cys Glu Leu Arg Val Lys Phe Ser
 260 265 270
 Arg Ser Ala Asp Ala Pro Ala Tyr Gln Gln Gly Gln Asn Gln Leu Tyr
 275 280 285
 Asn Glu Leu Asn Leu Gly Arg Arg Glu Glu Tyr Asp Val Leu Asp Lys
 290 295 300
 Arg Arg Gly Arg Asp Pro Glu Met Gly Gly Lys Pro Gln Arg Arg Lys
 305 310 315 320
 Asn Pro Gln Glu Gly Leu Tyr Asn Glu Leu Gln Lys Asp Lys Met Ala
 325 330 335
 Glu Ala Tyr Ser Glu Ile Gly Met Lys Gly Glu Arg Arg Arg Gly Lys
 340 345 350
 Gly His Asp Gly Leu Tyr Gln Gly Leu Ser Thr Ala Thr Lys Asp Thr
 355 360 365
 Tyr Asp Ala Leu His Met Gln Ala Leu Pro Pro Arg
 370 375 380

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<220>
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 20 25 30
 Pro Gly Gly Ser Leu Lys Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe
 35 40 45
 Ser Lys Phe Gly Met Ser Trp Val Arg Gln Thr Pro Asp Lys Arg Leu
 50 55 60

SP102075SGM SEQUENCE LISTING

Glu Trp Val Ala Ser Ile Ser Thr Gly Gly Tyr Asn Thr Phe Tyr Ser
 65 70 75 80
 Asp Asn Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn
 85 90 95
 Thr Leu Tyr Leu Gln Met Ser Ser Leu Lys Ser Glu Asp Thr Ala Met
 100 105 110
 Tyr Tyr Cys Ala Arg Gly Tyr Ser Ser Thr Ser Phe Ala Met Asp Tyr
 115 120 125
 Trp Gly Gln Gly Thr Met Val Thr Val Ser Ser Gly Ser Thr Ser Gly
 130 135 140
 Ser Gly Lys Pro Gly Ser Gly Glu Gly Ser Asp Ile Gln Met Thr Gln
 145 150 155 160
 Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp Arg Val Thr Ile Thr
 165 170 175
 Cys Met Thr Ser Thr Asp Ile Asp Asp Asp Met Asn Trp Tyr Gln Gln
 180 185 190
 Lys Pro Gly Lys Thr Pro Lys Leu Leu Ile Tyr Glu Gly Asn Thr Leu
 195 200 205
 Arg Pro Gly Val Pro Ser Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp
 210 215 220
 Phe Ile Phe Thr Ile Ser Ser Leu Gln Pro Glu Asp Ile Ala Thr Tyr
 225 230 235 240
 Tyr Cys Leu Gln Ser Phe Asn Val Pro Leu Thr Phe Gly Gly Gly Thr
 245 250 255
 Lys Val Glu Ile Lys Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Gly
 260 265 270
 Gly Gly Pro Arg Lys Pro Thr Thr Thr Pro Ala Pro Arg Pro Pro Thr
 275 280 285
 Pro Ala Pro Thr Ile Ala Ser Gln Pro Leu Ser Leu Arg Pro Glu Ala
 290 295 300
 Ala Arg Pro Ala Ala Gly Gly Ala Val His Thr Arg Gly Leu Asp Phe
 305 310 315 320
 Ala Leu Ala Tyr Leu Leu Asp Gly Ile Leu Phe Ile Tyr Gly Val Ile
 325 330 335
 Leu Thr Ala Leu Phe Leu Arg Val Lys Arg Gly Arg Lys Lys Leu Leu
 340 345 350
 Tyr Ile Phe Lys Gln Pro Phe Met Arg Pro Val Gln Thr Thr Gln Glu
 355 360 365
 Glu Asp Gly Cys Ser Cys Arg Phe Pro Glu Glu Glu Glu Gly Gly Cys
 370 375 380
 Glu Leu Lys Phe Ser Arg Ser Ala Asp Ala Pro Ala Tyr Gln Gln Gly
 385 390 395 400
 Gln Asn Gln Leu Tyr Asn Glu Leu Asn Leu Gly Arg Arg Glu Glu Tyr
 405 410 415
 Asp Val Leu Asp Lys Arg Arg Gly Arg Asp Pro Glu Met Gly Gly Lys

SP102075SGM SEQUENCE LISTING

420 425 430
 Pro Gln Arg Arg Lys Asn Pro Gln Glu Gly Leu Tyr Asn Glu Leu Gln
 435 440 445
 Lys Asp Lys Met Ala Glu Ala Tyr Ser Glu Ile Gly Met Lys Gly Glu
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Arg

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 35 40 45
 Ser Lys Phe Gly Met Ser Trp Val Arg Gln Thr Pro Asp Lys Arg Leu
 50 55 60
 Glu Trp Val Ala Ser Ile Ser Thr Gly Gly Tyr Asn Thr Phe Tyr Ser
 65 70 75 80
 Asp Asn Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn
 85 90 95
 Thr Leu Tyr Leu Gln Met Ser Ser Leu Lys Ser Glu Asp Thr Ala Met
 100 105 110
 Tyr Tyr Cys Ala Arg Gly Tyr Ser Ser Thr Ser Phe Ala Met Asp Tyr
 115 120 125
 Trp Gly Gln Gly Thr Met Val Thr Val Ser Ser Gly Ser Thr Ser Gly
 130 135 140
 Ser Gly Lys Pro Gly Ser Gly Glu Gly Ser Asp Ile Gln Met Thr Gln
 145 150 155 160
 Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp Arg Val Thr Ile Thr
 165 170 175
 Cys Met Thr Ser Thr Asp Ile Asp Asp Asp Met Asn Trp Tyr Gln Gln
 180 185 190
 Lys Pro Gly Lys Thr Pro Lys Leu Leu Ile Tyr Glu Gly Asn Thr Leu
 195 200 205
 Arg Pro Gly Val Pro Ser Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp

SP102075SGM SEQUENCE LISTING

210	215	220
Phe 225	Ile Phe Thr Ile Ser 230	Ser Leu Gln Pro Glu 235
Tyr 240	Cys Leu Gln Ser 245	Phe Asn Val Pro Leu 250
Lys 255	Val Glu Ile Lys 260	Glu Gln Lys Leu 265
Gly 270	Gly Pro Arg Lys 275	Pro Thr Thr Thr 280
Pro 285	Ala Pro Thr Ile Ala 290	Ser Gln Pro Leu Ser 295
Ala 300	Arg Pro Thr Ile Ala 305	Gly Gly Ala Val His 310
Phe 315	Ala Leu Ala Tyr Leu 320	Leu Asp Gly Ile Leu 325
Ile 330	Thr Ala Leu Phe Leu 335	Arg Val Arg Val Arg 340
Gly 345	His Ser Asp Tyr Met 350	Asn Met Thr Pro Arg 355
Arg 360	Lys His Tyr Gln Pro 365	Tyr Ala Pro Pro Arg 370
Arg 375	Ser Lys Arg Ser Arg 380	Thr Phe Lys Gln Pro 385
Phe 390	Met Arg Pro Val Gln 395	Thr Thr Gln Glu Glu 400
Arg 405	Phe Pro Glu Glu Glu 410	Gly Gly Cys Glu Leu 415
Ser 420	Ala Asp Ala Pro Ala 425	Tyr Gln Gln Gly Gln 430
Arg 435	Leu Asn Leu Gly Arg 440	Arg Glu Tyr Asp Val 445
Arg 450	Gly Arg Asp Pro Glu 455	Met Gly Gly Lys Pro 460
Pro 465	Gln Glu Gly Leu Tyr 470	Asn Glu Leu Gln Lys 475
Ala 480	Tyr Ser Glu Ile Gly 485	Met Lys Gly Glu Arg 490
Gly 495	Asp Gly Leu Tyr Gln 500	Gly Leu Ser Thr Ala 505
Thr 510	Thr Ser Glu Ile Gly 515	Met Lys Gly Glu Arg 520
Asp 525	Ala Leu His Met Gln 530	Ala Leu Pro Pro Arg 535

<210> 21
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 <213> Unknown

SP102075SGM SEQUENCE LISTING

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<223> HUMAN

<400> 21

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      20      25      30
Phe Arg Asn Glu Asp Tyr Thr Ile His Val Gln Leu Asn Asp Tyr Val
      35      40      45
Asp Ile Ile Cys Pro His Tyr Glu Asp His Ser Val Ala Asp Ala Ala
 50      55      60
Met Glu Gln Tyr Ile Leu Tyr Leu Val Glu His Glu Glu Tyr Gln Leu
 65      70      75      80
Cys Gln Pro Gln Ser Lys Asp Gln Val Arg Trp Gln Cys Asn Arg Pro
      85      90      95
Ser Ala Lys His Gly Pro Glu Lys Leu Ser Glu Lys Phe Gln Arg Phe
      100      105      110
Thr Ala Phe Ala Leu Ala Lys Glu Phe Lys Ala Gly His Ser Tyr Tyr
      115      120      125
Tyr Ile Ser Lys Pro Ile His Gln His Glu Asp Arg Cys Leu Arg Leu
      130      135      140
Lys Val Thr Val Ser Gly Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
      145      150      155      160
Gly Gly Gly Pro Arg Lys Pro Thr Thr Thr Pro Ala Pro Arg Pro Pro
      165      170      175
Thr Pro Ala Pro Thr Ile Ala Ser Gln Pro Leu Ser Leu Arg Pro Glu
      180      185      190
Ala Cys Arg Pro Ala Ala Gly Gly Ala Val His Thr Arg Gly Leu Asp
      195      200      205
Phe Ala Cys Asp Ile Tyr Ile Trp Ala Pro Leu Ala Gly Thr Cys Gly
      210      215      220
Val Leu Leu Leu Ser Leu Val Ile Thr Lys Arg Gly Arg Lys Lys Leu
      225      230      235      240
Leu Tyr Ile Phe Lys Gln Pro Phe Met Arg Pro Val Gln Thr Thr Gln
      245      250      255
Glu Glu Asp Gly Cys Ser Cys Arg Phe Pro Glu Glu Glu Glu Gly Gly
      260      265      270
Cys Glu Leu Arg Val Lys Phe Ser Arg Ser Ala Asp Ala Pro Ala Tyr
      275      280      285
Gln Gln Gly Gln Asn Gln Leu Tyr Asn Glu Leu Asn Leu Gly Arg Arg
      290      295      300
Glu Glu Tyr Asp Val Leu Asp Lys Arg Arg Gly Arg Asp Pro Glu Met
      305      310      315      320
Gly Gly Lys Pro Gln Arg Arg Lys Asn Pro Gln Glu Gly Leu Tyr Asn
      325      330      335

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SP102075SGM SEQUENCE LISTING

Glu Leu Gln Lys Asp Lys Met Ala Glu Ala Tyr Ser Glu Ile Gly Met
 340 345 350
 Lys Gly Glu Arg Arg Arg Gly Lys Gly His Asp Gly Leu Tyr Gln Gly
 355 360 365
 Leu Ser Thr Ala Thr Lys Asp Thr Tyr Asp Ala Leu His Met Gln Ala
 370 375 380
 Leu Pro Pro Arg
 385

<210> 22
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 <212> PRT
 <213> Unknown

<220>
 <223> HUMAN

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 35 40 45
 Asp Ile Ile Cys Pro His Tyr Glu Asp His Ser Val Ala Asp Ala Ala
 50 55 60
 Met Glu Gln Tyr Ile Leu Tyr Leu Val Glu His Glu Glu Tyr Gln Leu
 65 70 75 80
 Cys Gln Pro Gln Ser Lys Asp Gln Val Arg Trp Gln Cys Asn Arg Pro
 85 90 95
 Ser Ala Lys His Gly Pro Glu Lys Leu Ser Glu Lys Phe Gln Arg Phe
 100 105 110
 Thr Ala Phe Ala Leu Ala Lys Glu Phe Lys Ala Gly His Ser Tyr Tyr
 115 120 125
 Tyr Ile Ser Lys Pro Ile His Gln His Glu Asp Arg Cys Leu Arg Leu
 130 135 140
 Lys Val Thr Val Ser Gly Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
 145 150 155 160
 Gly Gly Gly Pro Arg Lys Pro Thr Thr Thr Pro Ala Pro Arg Pro Pro
 165 170 175
 Thr Pro Ala Pro Thr Ile Ala Ser Gln Pro Leu Ser Leu Arg Pro Glu
 180 185 190
 Ala Cys Arg Pro Ala Ala Gly Gly Ala Val His Thr Arg Gly Leu Asp
 195 200 205
 Phe Ala Cys Asp Ile Tyr Ile Trp Ala Pro Leu Ala Gly Thr Cys Gly
 210 215 220
 Val Leu Leu Leu Ser Leu Val Ile Thr Arg Ser Lys Arg Ser Arg Gly
 225 230 235 240

SP102075SGM SEQUENCE LISTING

Gly His Ser Asp Tyr Met Asn Met Thr Pro Arg Arg Pro Gly Pro Thr
245 250 255
Arg Lys His Tyr Gln Pro Tyr Ala Pro Pro Arg Asp Phe Ala Ala Tyr
260 265 270
Arg Ser Lys Arg Gly Arg Lys Lys Leu Leu Tyr Ile Phe Lys Gln Pro
275 280 285
Phe Met Arg Pro Val Gln Thr Thr Gln Glu Glu Asp Gly Cys Ser Cys
290 295 300
Arg Phe Pro Glu Glu Glu Glu Gly Gly Cys Glu Leu Arg Val Lys Phe
305 310 315 320
Ser Arg Ser Ala Asp Ala Pro Ala Tyr Gln Gln Gly Gln Asn Gln Leu
325 330 335
Tyr Asn Glu Leu Asn Leu Gly Arg Arg Glu Glu Tyr Asp Val Leu Asp
340 345 350
Lys Arg Arg Gly Arg Asp Pro Glu Met Gly Gly Lys Pro Gln Arg Arg
355 360 365
Lys Asn Pro Gln Glu Gly Leu Tyr Asn Glu Leu Gln Lys Asp Lys Met
370 375 380
Ala Glu Ala Tyr Ser Glu Ile Gly Met Lys Gly Glu Arg Arg Arg Gly
385 390 395 400
Lys Gly His Asp Gly Leu Tyr Gln Gly Leu Ser Thr Ala Thr Lys Asp
405 410 415
Thr Tyr Asp Ala Leu His Met Gln Ala Leu Pro Pro Arg
420 425

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<212> PRT
<213> Unknown

<220>
<223> HUMAN

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

SP102075SGM SEQUENCE LISTING

Thr Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Gly Gly Gly Pro Arg
115 120 125
Lys Pro Thr Thr Thr Pro Ala Pro Arg Pro Pro Thr Pro Ala Pro Thr
130 135 140
Ile Ala Ser Gln Pro Leu Ser Leu Arg Pro Glu Ala Cys Arg Pro Ala
145 150 155 160
Ala Gly Gly Ala Val His Thr Arg Gly Leu Asp Phe Ala Cys Asp Ile
165 170 175
Tyr Ile Trp Ala Pro Leu Ala Gly Thr Cys Gly Val Leu Leu Leu Ser
180 185 190
Leu Val Ile Thr Lys Arg Gly Arg Lys Lys Leu Leu Tyr Ile Phe Lys
195 200 205
Gln Pro Phe Met Arg Pro Val Gln Thr Thr Gln Glu Glu Asp Gly Cys
210 215 220
Ser Cys Arg Phe Pro Glu Glu Glu Glu Gly Gly Cys Glu Leu Arg Val
225 230 235 240
Lys Phe Ser Arg Ser Ala Asp Ala Pro Ala Tyr Gln Gln Gly Gln Asn
245 250 255
Gln Leu Tyr Asn Glu Leu Asn Leu Gly Arg Arg Glu Glu Tyr Asp Val
260 265 270
Leu Asp Lys Arg Arg Gly Arg Asp Pro Glu Met Gly Gly Lys Pro Gln
275 280 285
Arg Arg Lys Asn Pro Gln Glu Gly Leu Tyr Asn Glu Leu Gln Lys Asp
290 295 300
Lys Met Ala Glu Ala Tyr Ser Glu Ile Gly Met Lys Gly Glu Arg Arg
305 310 315 320
Arg Gly Lys Gly His Asp Gly Leu Tyr Gln Gly Leu Ser Thr Ala Thr
325 330 335
Lys Asp Thr Tyr Asp Ala Leu His Met Gln Ala Leu Pro Pro Arg
340 345 350

<210> 24
<211> 392
<212> PRT
<213> Unknown

<220>
<223> HUMAN

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Met Gly Trp Ser Cys Ile Ile Leu Phe Leu Val Ala Thr Ala Thr Gly
1 5 10 15
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20 25 30
Thr Pro Thr Ser Leu Leu Ile Ser Trp Asp Ala Pro Ala Val Thr Val
35 40 45
Arg Tyr Tyr Arg Ile Thr Tyr Gly Glu Thr Gly Gly Asn Ser Pro Val
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SP102075SGM SEQUENCE LISTING

50		55		60
Gln 65	Glu Phe Thr Val	Pro 70	Gly Ser Lys Ser Thr	Ala Thr Ile Ser Gly 80
Leu	Lys Pro Gly Val 85	Asp Tyr Thr Ile Thr	Val Tyr Ala Val Thr 95	Pro
Arg	Gly Asp Trp 100	Asn Glu Gly Ser Lys 105	Pro Ile Ser Ile Asn Tyr Arg 110	
Thr	Glu Gln Lys Leu Ile Ser	Glu 120	Glu Asp Leu Gly Gly Gly Pro Arg 125	
Lys	Pro Thr Thr Thr Pro	Ala 135	Pro Arg Pro Pro Thr Pro Ala Pro Thr 140	
Ile 145	Ala Ser Gln Pro Leu Ser Leu Arg Pro	Glu 155	Ala Cys Arg Pro Ala 160	
Ala	Gly Gly Ala Val 165	His Thr Arg Gly Leu Asp Phe Ala Cys Asp Ile 175		
Tyr Ile Trp	Ala 180	Pro Leu Ala Gly Thr Cys Gly Val Leu Leu Leu Ser 190		
Leu Val	Ile Thr Arg Ser Lys Arg Ser Arg Gly Gly His Ser Asp Tyr 205			
Met	Asn Met Thr Pro Arg Arg Pro Gly Pro Thr Arg Lys His Tyr Gln 210			
Pro 225	Tyr Ala Pro Pro Arg Asp Phe Ala Ala Tyr Arg Ser Lys Arg Gly 230			
Arg	Lys Lys Leu Leu Tyr Ile Phe Lys Gln Pro Phe Met Arg Pro Val 245			
Gln Thr Thr	Gln 260	Glu Glu Asp Gly Cys Ser Cys Arg Phe Pro Glu Glu 270		
Glu Glu	Gly Gly Cys Glu Leu Arg Val Lys Phe Ser Arg Ser Ala Asp 275			
Ala	Pro Ala Tyr Gln Gln Gly Gln Asn Gln Leu Tyr Asn Glu Leu Asn 290			
Leu 305	Gly Arg Arg Glu Glu Tyr Asp Val Leu Asp Lys Arg Arg Gly Arg 310			
Asp	Pro Glu Met Gly Gly Lys Pro Gln Arg Arg Lys Asn Pro Gln Glu 325			
Gly Leu Tyr	Asn 340	Glu Leu Gln Lys Asp Lys Met Ala Glu Ala Tyr Ser 345		
Glu Ile	Gly Met Lys Gly Glu Arg Arg Arg Gly Lys Gly His Asp Gly 355			
Leu Tyr	Gln Gly Leu Ser Thr Ala Thr Lys Asp Thr Tyr Asp Ala Leu 370			
His 385	Met Gln Ala Leu Pro Pro Arg 390			

<210> 25
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SP102075SGM SEQUENCE LISTING

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<220>
<223> HUMAN

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35 40 45
Tyr Gln Gly Glu Lys Cys Glu Lys Leu Val Ser Val Asn Phe Ile Asn
50 55 60
Lys Glu Ser Tyr Leu Gln Ile Pro Ser Ala Lys Val Arg Pro Gln Thr
65 70 75 80
Asn Ile Thr Leu Gln Ile Ala Thr Asp Glu Asp Ser Gly Ile Leu Leu
85 90 95
Tyr Lys Gly Asp Lys Asp His Ile Ala Val Glu Leu Tyr Arg Gly Arg
100 105 110
Val Arg Ala Ser Tyr Asp Thr Gly Ser His Pro Ala Ser Ala Ile Tyr
115 120 125
Ser Val Glu Thr Ile Asn Asp Gly Asn Phe His Ile Val Glu Leu Leu
130 135 140
Ala Leu Asp Gln Ser Leu Ser Leu Ser Val Asp Gly Gly Asn Pro Lys
145 150 155 160
Ile Ile Thr Asn Leu Ser Lys Gln Ser Thr Leu Asn Phe Asp Ser Pro
165 170 175
Leu Tyr Val Gly Gly Met Pro Gly Lys Ser Asn Val Ala Ser Leu Arg
180 185 190
Gln Ala Pro Gly Gln Asn Gly Thr Ser Phe His Gly Cys Ile Arg Asn
195 200 205
Leu Tyr Ile Asn Ser Glu Leu Gln Asp Phe Gln Lys Val Pro Met Gln
210 215 220
Thr Gly Ile Leu Pro Gly Cys Glu Pro Cys His Lys Lys Val Cys Ala
225 230 235 240
His Gly Thr Cys Gln Pro Ser Ser Gln Ala Gly Phe Thr Cys Glu Cys
245 250 255
Gln Glu Gly Trp Met Gly Pro Leu Cys Asp Gln Arg Thr Asn Asp Pro
260 265 270
Cys Leu Gly Asn Lys Cys Val His Gly Thr Cys Leu Pro Ile Asn Ala
275 280 285
Phe Ser Tyr Ser Cys Lys Cys Leu Glu Gly His Gly Gly Val Leu Cys
290 295 300
Asp Glu Glu Glu Asp Leu Phe Asn Pro Cys Gln Ala Ile Lys Cys Lys
305 310 315 320

SP102075SGM SEQUENCE LISTING

His Gly Lys Cys Arg 325 Leu Ser Gly Leu Gly 330 Gln Pro Tyr Cys Glu 335 Cys
 Ser Ser Gly Tyr 340 Thr Gly Asp Ser Cys 345 Asp Arg Glu Ile Ser 350 Cys Arg
 Gly Glu Arg 355 Ile Arg Asp Tyr Tyr 360 Gln Lys Gln Gln Gly 365 Tyr Ala Ala
 Cys Gln 370 Thr Thr Lys Lys Val 375 Ser Arg Leu Glu Cys 380 Arg Gly Gly Cys
 Ala Gly Gly Gln Cys Cys 390 Gly Pro Leu Arg Ser 395 Lys Arg Arg Lys Tyr 400
 Ser Phe Glu Cys Thr 405 Asp Gly Ser Ser Phe 410 Val Asp Glu Val Glu 415 Lys
 Val Val Lys Cys 420 Gly Cys Thr Arg Cys 425 Val Ser Glu Gln Lys 430 Leu Ile
 Ser Glu Glu 435 Asp Leu Gly Gly Gly 440 Pro Arg Lys Pro Thr 445 Thr Thr Pro
 Ala Pro Arg Pro Pro Thr Pro 455 Ala Pro Thr Ile Ala 460 Ser Gln Pro Leu
 Ser Leu Arg Pro Glu Ala 470 Cys Arg Pro Ala Ala 475 Gly Gly Ala Val His 480
 Thr Arg Gly Leu Asp 485 Phe Ala Cys Asp Ile 490 Tyr Ile Trp Ala Pro 495 Leu
 Ala Gly Thr Cys 500 Gly Val Leu Leu Leu 505 Ser Leu Val Ile Thr 510 Lys Arg
 Gly Arg Lys 515 Lys Leu Leu Tyr Ile 520 Phe Lys Gln Pro Phe 525 Met Arg Pro
 Val Gln Thr Thr Gln Glu Glu 535 Asp Gly Cys Ser Cys 540 Arg Phe Pro Glu
 Glu Glu Glu Gly Gly Cys 550 Glu Leu Arg Val Lys 555 Phe Ser Arg Ser Ala 560
 Asp Ala Pro Ala Tyr 565 Gln Gln Gly Gln Asn 570 Gln Leu Tyr Asn Glu 575 Leu
 Asn Leu Gly Arg 580 Arg Glu Glu Tyr Asp 585 Val Leu Asp Lys Arg 590 Arg Gly
 Arg Asp Pro Glu Met Gly Gly Lys 600 Pro Gln Arg Arg Lys 605 Asn Pro Gln
 Glu Gly Leu Tyr Asn Glu Leu Gln Lys Asp Lys Met 620 Ala Glu Ala Tyr
 Ser Glu Ile Gly Met Lys 630 Gly Glu Arg Arg Arg 635 Gly Lys Gly His Asp 640
 Gly Leu Tyr Gln Gly 645 Leu Ser Thr Ala Thr 650 Lys Asp Thr Tyr Asp 655 Ala
 Leu His Met Gln 660 Ala Leu Pro Pro Arg 665

SP102075SGM SEQUENCE LISTING

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<212> PRT
<213> Unknown

<220>
<223> HUMAN

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Gln Cys Ile Val Arg Ile Asn Glu Pro Ile Cys Gln Cys Leu Pro Gly
35 40 45
Tyr Gln Gly Glu Lys Cys Glu Lys Leu Val Ser Val Asn Phe Ile Asn
50 55 60
Lys Glu Ser Tyr Leu Gln Ile Pro Ser Ala Lys Val Arg Pro Gln Thr
65 70 75 80
Asn Ile Thr Leu Gln Ile Ala Thr Asp Glu Asp Ser Gly Ile Leu Leu
85 90 95
Tyr Lys Gly Asp Lys Asp His Ile Ala Val Glu Leu Tyr Arg Gly Arg
100 105 110
Val Arg Ala Ser Tyr Asp Thr Gly Ser His Pro Ala Ser Ala Ile Tyr
115 120 125
Ser Val Glu Thr Ile Asn Asp Gly Asn Phe His Ile Val Glu Leu Leu
130 135 140
Ala Leu Asp Gln Ser Leu Ser Leu Ser Val Asp Gly Gly Asn Pro Lys
145 150 155 160
Ile Ile Thr Asn Leu Ser Lys Gln Ser Thr Leu Asn Phe Asp Ser Pro
165 170 175
Leu Tyr Val Gly Gly Met Pro Gly Lys Ser Asn Val Ala Ser Leu Arg
180 185 190
Gln Ala Pro Gly Gln Asn Gly Thr Ser Phe His Gly Cys Ile Arg Asn
195 200 205
Leu Tyr Ile Asn Ser Glu Leu Gln Asp Phe Gln Lys Val Pro Met Gln
210 215 220
Thr Gly Ile Leu Pro Gly Cys Glu Pro Cys His Lys Lys Val Cys Ala
225 230 235 240
His Gly Thr Cys Gln Pro Ser Ser Gln Ala Gly Phe Thr Cys Glu Cys
245 250 255
Gln Glu Gly Trp Met Gly Pro Leu Cys Asp Gln Arg Thr Asn Asp Pro
260 265 270
Cys Leu Gly Asn Lys Cys Val His Gly Thr Cys Leu Pro Ile Asn Ala
275 280 285
Phe Ser Tyr Ser Cys Lys Cys Leu Glu Gly His Gly Gly Val Leu Cys
290 295 300
Asp Glu Glu Glu Asp Leu Phe Asn Pro Cys Gln Ala Ile Lys Cys Lys
305 310 315 320

SP102075SGM SEQUENCE LISTING

His Gly Lys Cys Arg₃₂₅ Leu Ser Gly Leu Gly₃₃₀ Gln Pro Tyr Cys Glu₃₃₅ Cys
Ser Ser Gly Tyr₃₄₀ Thr Gly Asp Ser Cys₃₄₅ Asp Arg Glu Ile Ser₃₅₀ Cys Arg
Gly Glu Arg₃₅₅ Ile Arg Asp Tyr Tyr₃₆₀ Gln Lys Gln Gln Gly₃₆₅ Tyr Ala Ala
Cys Gln Thr Thr Lys Lys Val₃₇₅ Ser Arg Leu Glu Cys₃₈₀ Arg Gly Gly Cys
Ala₃₈₅ Gly Gly Gln Cys Cys₃₉₀ Gly Pro Leu Arg Ser₃₉₅ Lys Arg Arg Lys Tyr₄₀₀
Ser Phe Glu Cys Thr₄₀₅ Asp Gly Ser Ser Phe Val₄₁₀ Asp Glu Val Glu₄₁₅ Lys
Val Val Lys Cys₄₂₀ Gly Cys Thr Arg Cys₄₂₅ Val Ser Glu Gln Lys₄₃₀ Leu Ile
Ser Glu Glu₄₃₅ Asp Leu Gly Gly Gly₄₄₀ Pro Arg Lys Pro Thr₄₄₅ Thr Thr Pro
Ala Pro₄₅₀ Arg Pro Pro Thr Pro₄₅₅ Ala Pro Thr Ile Ala₄₆₀ Ser Gln Pro Leu
Ser₄₆₅ Leu Arg Pro Glu Ala₄₇₀ Cys Arg Pro Ala Ala₄₇₅ Gly Gly Ala Val His₄₈₀
Thr Arg Gly Leu Asp₄₈₅ Phe Ala Cys Asp Ile₄₉₀ Tyr Ile Trp Ala Pro₄₉₅ Leu
Ala Gly Thr Cys₅₀₀ Gly Val Leu Leu Leu₅₀₅ Ser Leu Val Ile Thr₅₁₀ Arg Ser
Lys Arg Ser₅₁₅ Arg Gly Gly His Ser₅₂₀ Asp Tyr Met Asn Met₅₂₅ Thr Pro Arg
Arg Pro₅₃₀ Gly Pro Thr Arg Lys₅₃₅ His Tyr Gln Pro Tyr₅₄₀ Ala Pro Pro Arg
Asp₅₄₅ Phe Ala Ala Tyr Arg₅₅₀ Ser Lys Arg Gly Arg₅₅₅ Lys Lys Leu Leu Tyr₅₆₀
Ile Phe Lys Gln Pro₅₆₅ Phe Met Arg Pro Val₅₇₀ Gln Thr Thr Gln Glu₅₇₅ Glu
Asp Gly Cys Ser₅₈₀ Cys Arg Phe Pro Glu₅₈₅ Glu Glu Glu Gly₅₉₀ Gly Cys Glu
Leu Arg Val₅₉₅ Lys Phe Ser Arg Ser₆₀₀ Ala Asp Ala Pro Ala₆₀₅ Tyr Gln Gln
Gly Gln Asn Gln Leu Tyr Asn₆₁₅ Glu Leu Asn Leu Gly₆₂₀ Arg Arg Glu Glu
Tyr₆₂₅ Asp Val Leu Asp Lys₆₃₀ Arg Arg Gly Arg Asp₆₃₅ Pro Glu Met Gly Gly₆₄₀
Lys Pro Gln Arg Arg₆₄₅ Lys Asn Pro Gln Glu₆₅₀ Gly Leu Tyr Asn Glu₆₅₅ Leu
Gln Lys Asp Lys₆₆₀ Met Ala Glu Ala Tyr₆₆₅ Ser Glu Ile Gly Met₆₇₀ Lys Gly
Glu Arg Arg Arg Gly Lys Gly His Asp Gly Leu Tyr Gln Gly Leu Ser

SP102075SGM SEQUENCE LISTING															
675					680					685					
Thr	Ala	Thr	Lys	Asp	Thr	Tyr	Asp	Ala	Leu	His	Met	Gln	Ala	Leu	Pro
690						695					700				
Pro	Arg														
705															