SEQUENCE LISTING

<110> Jain, Arjun

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<141> 2016-01-22

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Gln Pro Thr Asn Leu Val Leu Pro Ser Asn Gly Ser Met His Asn Tyr
35 40 45

Cys Pro Gln Gln Thr Lys Ile Thr Ser Ala Phe Lys Lys Leu Leu Ala
50 55 60

Gly Arg Trp Pro Phe Asp His Asn Asp Phe Gly Val Phe Leu Cys Pro
65 70 75 80

Phe Glu Tyr Arg Gly Glu Gln His Lys Thr Cys Met Leu Asn Ala Thr
85 90 95

Ser Lys Phe Met Glu Phe Tyr Gln Asp Val Lys Asp Thr Cys Glu Met
100 105 110

Leu Asn Arg Arg Asn Gly Ser Leu Arg Ile Ala Leu Ser Glu His Leu
115 120 125

Lys Gln Arg Arg Glu Val Ala Lys Thr Val Phe Cys Val Tyr Asn Glu
130 135 140

Met Asp Lys Asn Arg Cys Glu Leu Leu Ser Phe Leu Ser Lys Lys Phe
145 150 155 160

Lys Asn Cys Phe Gln Ser Cys Leu Cys Cys Cys Cys Tyr Gln Ser Lys
165 170 175

Ser Leu Met Thr Ser Val Pro Met Asn Gly Thr Ser Ile Gln Trp Lys
180 185 190

Asn His Asp Gln Asn Asn His Asn Thr Asp Arg Ser Ser His Lys Asp

195 200 205

Ser Met Asn Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu
210 215 220

Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr
225 230 235 240

Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val
245 250 255

Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val
260 265 270

Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser
275 280 285

Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu
290 295 300

Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala
305 310 315 320

Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro
325 330 335

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Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln
            340                 345                 350

Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala
        355                 360                 365

Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr
    370                 375                 380

Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu
385                 390                 395                 400

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

Gln Pro Thr Asn Leu Val Leu Pro Ser Asn Gly Ser Met His Asn Tyr
35 40 45

Cys Pro Gln Gln Thr Lys Ile Thr Ser Ala Phe Lys Lys Leu Leu Ala
50 55 60

Gly Arg Trp Pro Phe Asp His Asn Asp Phe Gly Val Phe Leu Cys Pro
65 70 75 80

Phe Glu Tyr Arg Gly Glu Gln His Lys Thr Cys Met Leu Asn Ala Thr
85 90 95

Ser Lys Phe Met Glu Phe Tyr Gln Asp Val Lys Asp Ser Glu His Leu
100 105 110

Lys Gln Arg Arg Glu Val Ala Lys Thr Val Phe Cys Val Tyr Asn Glu
115 120 125

Met Asp Lys Asn Arg Cys Glu Leu Leu Ser Phe Leu Ser Lys Lys Phe
130 135 140

Lys Asn Cys Phe Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu
145 150 155 160

Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp
165 170 175

Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp
180 185 190

Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly
195 200 205

Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn
210 215 220

Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp
225 230 235 240

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

Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu
260 265 270

Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn
275 280 285

Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile
290 295 300

Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr
305 310 315 320

Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys
325 330 335

Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys
340 345 350

Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu
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Ser Leu Ser Pro Gly Lys
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Phe Thr Thr Phe Arg Gly Thr Glu Leu Ser Phe Leu Val Thr Thr His
20 25 30

Gln Pro Thr Asn Leu Val Leu Pro Ser Asn Gly Ser Met His Asn Tyr

35 40 45

Cys Pro Gln Gln Thr Lys Ile Thr Ser Ala Phe Lys Lys Leu Leu Ala
50 55 60

Gly Arg Trp Pro Phe Asp His Asn Asp Phe Gly Val Phe Leu Cys Pro
65 70 75 80

Phe Glu Tyr Arg Gly Glu Gln His Lys Thr Cys Met Leu Asn Ala Thr
85 90 95

Ser Lys Phe Met Glu Phe Tyr Gln Asp Val Lys Asp Thr Cys Glu Met
100 105 110

Leu Asn Arg Arg Asn Gly Ser Leu Arg Ile Ala Leu Ser Glu His Leu
115 120 125

Lys Gln Arg Arg Glu Val Ala Lys Thr Val Phe Cys Ser Lys Lys Phe
130 135 140

Lys Asn Cys Phe Gln Ser Cys Leu Cys Cys Cys Cys Tyr Gln Ser Lys
145 150 155 160

Ser Leu Met Thr Ser Val Pro Met Asn Gly Thr Ser Ile Gln Trp Lys
165 170 175

Asn His Asp Gln Asn Asn His Asn Thr Asp Arg Ser Ser His Lys Asp
180 185 190

Ser Met Asn Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu
195 200 205

Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr
210 215 220

Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val
225 230 235 240

Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val
245 250 255

Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser
260 265 270

Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu
275 280 285

Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala
290 295 300

Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro
305 310 315 320

Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln
325 330 335

Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala
340 345 350

Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr
355 360 365

Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu
370 375 380

Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser
385 390 395 400

Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser
405 410 415

Leu Ser Pro Gly Lys
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20 25 30

Gln Pro Thr Asn Leu Val Leu Pro Ser Asn Gly Ser Met His Asn Tyr
35 40 45

Cys Pro Gln Gln Thr Lys Ile Thr Ser Ala Phe Lys Lys Leu Leu Ala
50 55 60

Gly Arg Trp Pro Phe Asp His Asn Asp Phe Gly Val Phe Leu Cys Pro
65 70 75 80

Phe Glu Tyr Arg Gly Glu Gln His Lys Thr Cys Met Leu Asn Ala Thr
85 90 95

Ser Lys Phe Met Glu Phe Tyr Gln Asp Val Lys Asp Ser Glu His Leu
100 105 110

Lys Gln Arg Arg Glu Val Ala Lys Thr Val Phe Cys Ser Lys Lys Phe
115 120 125

Lys Asn Cys Phe Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu
130 135 140

Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp
145 150 155 160

Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp
165 170 175

Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly
180 185 190

Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn
195 200 205

Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp
210 215 220

Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro
225 230 235 240

Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu
245 250 255

Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn
260 265 270

Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile
275 280 285

Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr
290 295 300

Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys
305 310 315 320

Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys
325 330 335

Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu
340 345 350

Ser Leu Ser Pro Gly Lys
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Gln Pro Thr Asn Leu Val Leu Pro Ser Asn Gly Ser Met His Asn Tyr
35 40 45

Cys Pro Gln Gln Thr Lys Ile Thr Ser Ala Phe Lys Lys Leu Leu Ala
50 55 60

Gly Arg Trp Pro Phe Asp His Asn Asp Phe Gly Val Phe Leu Cys Thr

65 70 75 80

Cys Glu Met Leu Asn Arg Arg Asn Gly Ser Leu Arg Ile Ala Leu Ser
85 90 95

Glu His Leu Lys Gln Arg Arg Glu Val Ala Lys Thr Val Phe Cys Val
100 105 110

Tyr Asn Glu Met Asp Lys Asn Arg Cys Glu Leu Leu Ser Phe Leu Ser
115 120 125

Lys Lys Phe Lys Asn Cys Phe Gln Ser Cys Leu Cys Cys Cys Cys Tyr
130 135 140

Gln Ser Lys Ser Leu Met Thr Ser Val Pro Met Asn Gly Thr Ser Ile
145 150 155 160

Gln Trp Lys Asn His Asp Gln Asn Asn His Asn Thr Asp Arg Ser Ser
165 170 175

His Lys Asp Ser Met Asn Lys Thr His Thr Cys Pro Pro Cys Pro Ala
180 185 190

Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro
195 200 205

Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val
210 215 220

Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val
225 230 235 240

Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln
245 250 255

Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln
260 265 270

Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala
275 280 285

Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro
290 295 300

Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr
305 310 315 320

Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser
325 330 335

Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr
340 345 350

Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr
355 360 365

Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe
370 375 380

Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys
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Ser Leu Ser Leu Ser Pro Gly Lys
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Phe Thr Thr Phe Arg Gly Thr Glu Leu Ser Phe Leu Val Thr Thr His
            20                  25                  30

Gln Pro Thr Asn Leu Val Leu Pro Ser Asn Gly Ser Met His Asn Tyr
        35                  40                  45

Cys Pro Gln Gln Thr Lys Ile Thr Ser Ala Phe Lys Lys Leu Leu Ala
    50                  55                  60

Gly Arg Trp Pro Phe Asp His Asn Asp Phe Gly Val Phe Leu Cys Ser
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65 70 75 80

Glu His Leu Lys Gln Arg Arg Glu Val Ala Lys Thr Val Phe Cys Val
85 90 95

Tyr Asn Glu Met Asp Lys Asn Arg Cys Glu Leu Leu Ser Phe Leu Ser
100 105 110

Lys Lys Phe Lys Asn Cys Phe Lys Thr His Thr Cys Pro Pro Cys Pro
115 120 125

Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys
130 135 140

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

Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr
165 170 175

Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu
180 185 190

Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His
195 200 205

Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys
210 215 220

Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln
225 230 235 240

Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met
245 250 255

Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro
260 265 270

Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn
275 280 285

Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu
290 295 300

Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val
305 310 315 320

Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln
325 330 335

Lys Ser Leu Ser Leu Ser Pro Gly Lys
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Cys Leu Val Phe Val Leu Gly Ile Ile Gly Asn Ser Thr Leu Leu Arg
20 25 30

Ile Ile Tyr Lys Asn Lys Cys Ser Lys Arg Phe Lys Asn Cys Phe Lys
35 40 45

Ser Cys Leu Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu

50 55 60

Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr
65 70 75 80

Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val
85 90 95

Ser His Glu Asp Pro Glu Val Lys Phe Asp Trp Tyr Val Asp Gly Val
100 105 110

Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser
115 120 125

Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu
130 135 140

Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala
145 150 155 160

Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro
165 170 175

Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln
180 185 190

Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala
195 200 205

Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr
210 215 220

Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu
225 230 235 240

Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser
245 250 255

Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser
260 265 270

Leu Ser Pro Gly Lys
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Cys Leu Val Phe Val Leu Gly Ile Ile Gly Asn Ser Thr Leu Leu Arg
20 25 30

Ile Ile Tyr Lys Asn Lys Cys Ile Leu Ile Ala Ser Leu Ala Leu Gly
35 40 45

Asp Leu Leu His Ile Val Ile Asp Ile Pro Ile Asn Val Tyr Lys Leu
50 55 60

Leu Ala Glu Asp Trp Pro Phe Gly Ala Glu Met Cys Lys Leu Val Pro
65 70 75 80

Phe Ile Gln Lys Ala Ser Val Gly Ile Thr Val Leu Ser Leu Cys Ala
85 90 95

Leu Ser Ile Ser Lys Arg Phe Lys Asn Cys Phe Lys Ser Cys Leu Lys
100 105 110

Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro
115 120 125

Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser
130 135 140

Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu Asp
145 150 155 160

Pro Glu Val Lys Phe Asp Trp Tyr Val Asp Gly Val Glu Val His Asn
165 170 175

Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg Val
180 185 190

Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu
195 200 205

Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys
210 215 220

Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr
225 230 235 240

Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln Val Ser Leu Thr
245 250 255

Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu
260 265 270

Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu
275 280 285

Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys
290 295 300

Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu
305 310 315 320

Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly
325 330 335

Lys

<210> 9
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Cys Leu Val Phe Val Leu Gly Ile Ile Gly Asn Ser Thr Leu Leu Arg
20 25 30

Ile Ile Tyr Lys Asn Lys Cys Lys Thr His Thr Cys Pro Pro Cys Pro
35 40 45

Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys
50 55 60

Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val
65 70 75 80

Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asp Trp Tyr

85 90 95

Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu
100 105 110

Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His
115 120 125

Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys
130 135 140

Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln
145 150 155 160

Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met
165 170 175

Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro
180 185 190

Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn
195 200 205

Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu
210 215 220

Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val
225 230 235 240

Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln
245 250 255

Lys Ser Leu Ser Leu Ser Pro Gly Lys
260 265

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Lys Leu Leu Ala Gly
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Asp Asn Pro Glu Arg Tyr Ser Thr Asn Leu Ser Asn His Val Asp Asp
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Phe Thr Thr Phe Arg Gly Thr Glu Leu Ser Phe Leu Val Thr Thr His
20 25 30

Gln Pro Thr Asn Leu Val Leu Pro Ser Asn Gly Ser Met His Asn Tyr
35 40 45

Cys Pro Gln Gln Thr Lys Ile Thr Ser Ala Phe Lys
50 55 60

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<400> 12

Lys Leu Leu Ala Gly Arg Trp Pro Phe Asp His Asn Asp Phe Gly Val
1 5 10 15

Phe Leu Cys

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<400> 13

Pro Phe Glu Tyr Arg Gly Glu Gln His Lys Thr Cys Met Leu Asn Ala
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Thr Ser Lys Phe Met Glu Phe Tyr Gln Asp Val Lys Asp
20 25

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<400> 14

Val Tyr Asn Glu Met Asp Lys Asn Arg Cys Glu Leu Leu Ser Phe Leu
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Thr Cys Glu Met Leu Asn Arg Arg Asn Gly Ser Leu Arg Ile Ala Leu
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Ser Glu His Leu Lys Gln Arg Arg Glu Val Ala Lys Thr Val Phe Cys
20 25 30

<210> 16
<211> 55
<212> PRT
<213> Artificial Sequence

<220>
<223> ETA receptor ligand binding domain C terminus

<400> 16

Ser Lys Lys Phe Lys Asn Cys Phe Gln Ser Cys Leu Cys Cys Cys Cys
1 5 10 15

Tyr Gln Ser Lys Ser Leu Met Thr Ser Val Pro Met Asn Gly Thr Ser
20 25 30

Ile Gln Trp Lys Asn His Asp Gln Asn Asn His Asn Thr Asp Arg Ser
35 40 45

Ser His Lys Asp Ser Met Asn
50 55

<210> 17
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<400> 17

Pro Ile Glu Ile Lys Glu Thr Phe Lys Tyr Ile Asn Thr Val Val Ser
1 5 10 15

Cys Leu Val Phe Val Leu Gly Ile Ile Gly Asn Ser Thr Leu Leu Arg
20 25 30

Ile Ile Tyr Lys Asn Lys Cys
35

<210> 18
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<223> ETB receptor ligand binding domain C terminus

<400> 18

Ser Lys Arg Phe Lys Asn Cys Phe Lys Ser Cys Leu
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<210> 19

<211> 60

<212> PRT

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<223> ETB receptor ligand binding domain transmembrane domains 2 + 3

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Ile Leu Ile Ala Ser Leu Ala Leu Gly Asp Leu Leu His Ile Val Ile
1 5 10 15

Asp Ile Pro Ile Asn Val Tyr Lys Leu Leu Ala Glu Asp Trp Pro Phe
20 25 30

Gly Ala Glu Met Cys Lys Leu Val Pro Phe Ile Gln Lys Ala Ser Val
35 40 45

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