

SEQUENCE LISTING

<110> Eli Lilly and Company

<120> IL-17 RECEPTOR MODULATORS

<130> X21903M

<160> 3

<170> PatentIn version 3.5

<210> 1

<211> 176

<212> PRT

<213> Artificial sequence

<220>

<223> Synthetic construct

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Met Thr Pro Gly Lys Thr Ser Leu Val Ser Leu Leu Leu Leu Leu Ser
1 5 10 15

Leu Glu Ala Ile Val Lys Ala Gly Ile Thr Ile Pro Arg Asn Pro Gly
20 25 30

Cys Pro Asn Ser Glu Asp Lys Asn Phe Pro Arg Thr Val Met Val Asn
35 40 45

Leu Asn Ile His Asn Arg Asn Thr Asn Thr Asn Pro Lys Arg Ser Ser
50 55 60

Asp Tyr Tyr Asn Arg Ser Thr Ser Pro Trp Asn Leu His Arg Asn Glu
65 70 75 80

Asp Pro Glu Arg Tyr Pro Ser Val Ile Trp Glu Ala Lys Cys Arg His
85 90 95

Leu Gly Cys Ile Asn Ala Asp Gly Asn Val Asp Tyr His Met Asn Ser
100 105 110

Val Pro Ile Gln Gln Glu Ile Leu Val Leu Arg Arg Glu Pro Pro His
 115 120 125

Cys Pro Asn Ser Phe Arg Leu Glu Lys Ile Leu Val Ser Val Gly Cys
 130 135 140

Thr Cys Val Thr Pro Ile Val His His Val Ala Gly Leu Asn Asp Ile
 145 150 155 160

Phe Glu Ala Gln Lys Ile Glu Trp His Glu His His His His His His
 165 170 175

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 <213> Artificial sequence

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<400> 2

Met Ala Leu Lys Asp Asn Thr Val Pro Leu Lys Leu Ile Ala Leu Leu
 1 5 10 15

Ala Asn Gly Glu Phe His Ser Gly Glu Gln Leu Gly Glu Thr Leu Gly
 20 25 30

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

Gly Val Asp Val Phe Thr Val Pro Gly Lys Gly Tyr Ser Leu Pro Glu
 50 55 60

Pro Ile Gln Leu Leu Asn Ala Lys Gln Ile Leu Gly Gln Leu Asp Gly
 65 70 75 80

Gly Ser Val Ala Val Leu Pro Val Ile Asp Ser Thr Asn Gln Tyr Leu
 85 90 95

Leu Asp Arg Ile Gly Glu Leu Lys Ser Gly Asp Ala Cys Ile Ala Glu
 100 105 110

Tyr Gln Gln Ala Gly Arg Gly Arg Arg Gly Arg Lys Trp Phe Ser Pro
 115 120 125

Phe Gly Ala Asn Leu Tyr Leu Ser Met Phe Trp Arg Leu Glu Gln Gly
 130 135 140

Pro Ala Ala Ala Ile Gly Leu Ser Leu Val Ile Gly Ile Val Met Ala
 145 150 155 160

Glu Val Leu Arg Lys Leu Gly Ala Asp Lys Val Arg Val Lys Trp Pro
 165 170 175

Asn Asp Leu Tyr Leu Gln Asp Arg Lys Leu Ala Gly Ile Leu Val Glu
 180 185 190

Leu Thr Gly Lys Thr Gly Asp Ala Ala Gln Ile Val Ile Gly Ala Gly
 195 200 205

Ile Asn Met Ala Met Arg Arg Val Glu Glu Ser Val Val Asn Gln Gly
 210 215 220

Trp Ile Thr Leu Gln Glu Ala Gly Ile Asn Leu Asp Arg Asn Thr Leu
 225 230 235 240

Ala Ala Met Leu Ile Arg Glu Leu Arg Ala Ala Leu Glu Leu Phe Glu
 245 250 255

Gln Glu Gly Leu Ala Pro Tyr Leu Ser Arg Trp Glu Lys Leu Asp Asn
 260 265 270

Phe Ile Asn Arg Pro Val Lys Leu Ile Ile Gly Asp Lys Glu Ile Phe
 275 280 285

Gly Ile Ser Arg Gly Ile Asp Lys Gln Gly Ala Leu Leu Leu Glu Gln
 290 295 300

Asp Gly Ile Ile Lys Pro Trp Met Gly Gly Glu Ile Ser Leu Arg Ser
 305 310 315 320

Ala Glu Lys

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<220>
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<400> 3

Met Gly Ala Ala Arg Ser Pro Pro Ser Ala Val Pro Gly Pro Leu Leu
 1 5 10 15

Gly Leu Leu Leu Leu Leu Gly Val Leu Ala Pro Gly Gly Ala Ser
 20 25 30

Leu Arg Leu Leu Asp His Arg Ala Leu Val Cys Ser Gln Pro Gly Leu
 35 40 45

Asn Cys Thr Val Lys Asn Ser Thr Cys Leu Asp Asp Ser Trp Ile His
 50 55 60

Pro Arg Asn Leu Thr Pro Ser Ser Pro Lys Asp Leu Gln Ile Gln Leu
 65 70 75 80

His Phe Ala His Thr Gln Gln Gly Asp Leu Phe Pro Val Ala His Ile
 85 90 95

Glu Trp Thr Leu Gln Thr Asp Ala Ser Ile Leu Tyr Leu Glu Gly Ala
 100 105 110

Glu Leu Ser Val Leu Gln Leu Asn Thr Asn Glu Arg Leu Cys Val Arg
 115 120 125

Phe Glu Phe Leu Ser Lys Leu Arg His His His Arg Arg Trp Arg Phe
 130 135 140

Thr Phe Ser His Phe Val Val Asp Pro Asp Gln Glu Tyr Glu Val Thr
 145 150 155 160

Val His His Leu Pro Lys Pro Ile Pro Asp Gly Asp Pro Asn His Gln
 165 170 175

Ser Lys Asn Phe Leu Val Pro Asp Cys Glu His Ala Arg Met Lys Val
 180 185 190

Thr Thr Pro Cys Met Ser Ser Gly Ser Leu Trp Asp Pro Asp Ile Thr
 195 200 205

Val Glu Thr Leu Glu Ala His Gln Leu Arg Val Ser Phe Thr Leu Trp
 210 215 220

Asn Glu Ser Thr His Tyr Gln Ile Leu Leu Thr Ser Phe Pro His Met
 225 230 235 240

Glu Asn His Ser Cys Phe Glu His Met His His Ile Pro Ala Pro Arg
 245 250 255

Pro Glu Glu Phe His Gln Arg Ser Asp Val Thr Leu Thr Leu Arg Asn
 260 265 270

Leu Lys Gly Cys Cys Arg His Gln Val Gln Ile Gln Pro Phe Phe Ser
 275 280 285

Ser Cys Leu Asn Asp Cys Leu Arg His Ser Ala Thr Val Ser Cys Pro
 290 295 300

Glu Met Pro Asp Thr Pro Glu Pro Ile Pro Asp Tyr Met Leu Val Pro
 305 310 315 320

Arg Gly Ser Asp Tyr Lys Asp Asp Asp Asp Lys
325 330