

PhoenixTemp29451.tmp.txt
SEQUENCE LISTING

<110> Novo Nordisk A/S

<120> DI MERIC PROLACTIN RECEPTOR LIGANDS

<130> 7790.514-WO

<160> 3

<170> Patent In version 3.3

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<212> PRT

<213> Homo sapiens

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35 40 45

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

Thr Pro Glu Asp Lys Glu Gln Ala Gln Gln Met Asn Gln Lys Asp Phe
65 70 75 80

Leu Ser Leu Ile Val Ser Ile Leu Arg Ser Trp Asn Glu Pro Leu Tyr
85 90 95

His Leu Val Thr Glu Val Arg Gly Met Gln Glu Ala Pro Glu Ala Ile
100 105 110

Leu Ser Lys Ala Val Glu Ile Glu Glu Gln Thr Lys Arg Leu Leu Glu
115 120 125

Gly Met Glu Leu Ile Val Ser Gln Val His Pro Glu Thr Lys Glu Asn
130 135 140

Glu Ile Tyr Pro Val Trp Ser Gly Leu Pro Ser Leu Gln Met Ala Asp
145 150 155 160

Glu Glu Ser Arg Leu Ser Ala Tyr Tyr Asn Leu Leu His Cys Leu Arg
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Arg Asp Ser His Lys Ile Asp Asn Tyr Leu Lys Leu Leu Lys Cys Arg
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Ile Ile His Asn Asn Asn Cys

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Ala His Arg Leu His Gln Leu Ala Phe Asp Thr Tyr Gln Glu Phe Glu
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Glu Ala Tyr Ile Pro Lys Glu Gln Lys Tyr Ser Phe Leu Gln Asn Pro
 35 40 45

Gln Thr Ser Leu Cys Phe Ser Glu Ser Ile Pro Thr Pro Ser Asn Arg
 50 55 60

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

Leu Leu Ile Gln Ser Trp Leu Glu Pro Val Gln Phe Leu Arg Ser Val
 85 90 95

Phe Ala Asn Ser Leu Val Tyr Gly Ala Ser Asp Ser Asn Val Tyr Asp
 100 105 110

Leu Leu Lys Asp Leu Glu Glu Gly Ile Gln Thr Leu Met Gly Arg Leu
 115 120 125

Glu Asp Gly Ser Pro Arg Thr Gly Gln Ile Phe Lys Gln Thr Tyr Ser
 130 135 140

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