

PBD00056\_wo\_Seq.ST25.txt  
SEQUENCE LISTING

<110> Probiodrug AG

<120> Novel Inhibitors of GlutaminyI Cyclase

<130> PBD 00056/wo

<150> 60/912,525

<151> 2007-04-18

<160> 20

<170> PatentIn version 3.1

<210> 1

<211> 42

<212> PRT

<213> Homo sapiens

<400> 1

Asp Ala Glu Phe Arg His Asp Ser Gly Tyr Glu Val His His Gln Lys  
1 5 10 15

Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile  
20 25 30

Gly Leu Met Val Gly Gly Val Val Ile Ala  
35 40

<210> 2

<211> 40

<212> PRT

<213> Homo sapiens

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Asp Ala Glu Phe Arg His Asp Ser Gly Tyr Glu Val His His Gln Lys  
1 5 10 15

PBD00056\_WO\_Seq.ST25.txt

Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile  
 20 25 30

Gly Leu Met Val Gly Gly Val Val  
 35 40

<210> 3

<211> 40

<212> PRT

<213> Homo sapiens

<400> 3

Glu Phe Arg His Asp Ser Gly Tyr Glu Val His His Gln Lys Leu Val  
 1 5 10 15

Phe Phe Ala Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile Gly Leu  
 20 25 30

Met Val Gly Gly Val Val Ile Ala  
 35 40

<210> 4

<211> 38

<212> PRT

<213> Homo sapiens

<400> 4

Glu Phe Arg His Asp Ser Gly Tyr Glu Val His His Gln Lys Leu Val  
 1 5 10 15

Phe Phe Ala Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile Gly Leu  
 20 25 30

Met Val Gly Gly Val Val  
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<210> 5

<211> 17

<212> PRT

<213> Homo sapiens

<220>

<221> MOD\_RES

<222> (17)..(17)

<223> AMIDATION

<400> 5

Gln Gly Pro Trp Leu Glu Glu Glu Glu Glu Ala Tyr Gly Trp Met Asp  
1 5 10 15

Phe

<210> 6

<211> 13

<212> PRT

<213> Homo sapiens

<400> 6

Gln Leu Tyr Glu Asn Lys Pro Arg Arg Pro Tyr Ile Leu  
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<210> 7

<211> 10

<212> PRT

<213> Homo sapiens

<220>

<221> MOD\_RES

<222> (10)..(10)

<223> AMIDATION

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Gln His Trp Ser Tyr Gly Leu Arg Pro Gly  
1 5 10

<210> 8

<211> 97

<212> PRT

<213> Homo sapiens

PBD00056\_WO\_Seq.ST25.txt

<400> 8

Gln Pro Lys Val Pro Glu Trp Val Asn Thr Pro Ser Thr Cys Cys Leu  
1 5 10 15

Lys Tyr Tyr Glu Lys Val Leu Pro Arg Arg Leu Val Val Gly Tyr Arg  
20 25 30

Lys Ala Leu Asn Cys His Leu Pro Ala Ile Ile Phe Val Thr Lys Arg  
35 40 45

Asn Arg Glu Val Cys Thr Asn Pro Asn Asp Asp Trp Val Gln Glu Tyr  
50 55 60

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

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

Gln

<210> 9

<211> 76

<212> PRT

<213> Homo sapiens

<400> 9

Gln Pro Asp Ser Val Ser Ile Pro Ile Thr Cys Cys Phe Asn Val Ile  
1 5 10 15

Asn Arg Lys Ile Pro Ile Gln Arg Leu Glu Ser Tyr Thr Arg Ile Thr  
20 25 30

Asn Ile Gln Cys Pro Lys Glu Ala Val Ile Phe Lys Thr Lys Arg Gly  
35 40 45

Lys Glu Val Cys Ala Asp Pro Lys Glu Arg Trp Val Arg Asp Ser Met  
50 55 60

Lys His Leu Asp Gln Ile Phe Gln Asn Leu Lys Pro  
65 70 75

<210> 10

<211> 76

<212> PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 10

Gln Pro Asp Ala Ile Asn Ala Pro Val Thr Cys Cys Tyr Asn Phe Thr  
 1 5 10 15

Asn Arg Lys Ile Ser Val Gln Arg Leu Ala Ser Tyr Arg Arg Ile Thr  
 20 25 30

Ser Ser Lys Cys Pro Lys Glu Ala Val Ile Phe Lys Thr Ile Val Ala  
 35 40 45

Lys Glu Ile Cys Ala Asp Pro Lys Gln Lys Trp Val Gln Asp Ser Met  
 50 55 60

Asp His Leu Asp Lys Gln Thr Gln Thr Pro Lys Thr  
 65 70 75

&lt;210&gt; 11

&lt;211&gt; 68

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 11

Gln Val Gly Thr Asn Lys Glu Leu Cys Cys Leu Val Tyr Thr Ser Trp  
 1 5 10 15

Gln Ile Pro Gln Lys Phe Ile Val Asp Tyr Ser Glu Thr Ser Pro Gln  
 20 25 30

Cys Pro Lys Pro Gly Val Ile Leu Leu Thr Lys Arg Gly Arg Gln Ile  
 35 40 45

Cys Ala Asp Pro Asn Lys Lys Trp Val Gln Lys Tyr Ile Ser Asp Leu  
 50 55 60

Lys Leu Asn Ala  
 65

&lt;210&gt; 12

&lt;211&gt; 373

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 12

PBD00056\_WO\_Seq.ST25.txt

Gln His His Gly Val Thr Lys Cys Asn Ile Thr Cys Ser Lys Met Thr  
 1 5 10 15  
 Ser Lys Ile Pro Val Ala Leu Leu Ile His Tyr Gln Gln Asn Gln Ala  
 20 25 30  
 Ser Cys Gly Lys Arg Ala Ile Ile Leu Glu Thr Arg Gln His Arg Leu  
 35 40 45  
 Phe Cys Ala Asp Pro Lys Glu Gln Trp Val Lys Asp Ala Met Gln His  
 50 55 60  
 Leu Asp Arg Gln Ala Ala Ala Leu Thr Arg Asn Gly Gly Thr Phe Glu  
 65 70 75 80  
 Lys Gln Ile Gly Glu Val Lys Pro Arg Thr Thr Pro Ala Ala Gly Gly  
 85 90 95  
 Met Asp Glu Ser Val Val Leu Glu Pro Glu Ala Thr Gly Glu Ser Ser  
 100 105 110  
 Ser Leu Glu Pro Thr Pro Ser Ser Gln Glu Ala Gln Arg Ala Leu Gly  
 115 120 125  
 Thr Ser Pro Glu Leu Pro Thr Gly Val Thr Gly Ser Ser Gly Thr Arg  
 130 135 140  
 Leu Pro Pro Thr Pro Lys Ala Gln Asp Gly Gly Pro Val Gly Thr Glu  
 145 150 155 160  
 Leu Phe Arg Val Pro Pro Val Ser Thr Ala Ala Thr Trp Gln Ser Ser  
 165 170 175  
 Ala Pro His Gln Pro Gly Pro Ser Leu Trp Ala Glu Ala Lys Thr Ser  
 180 185 190  
 Glu Ala Pro Ser Thr Gln Asp Pro Ser Thr Gln Ala Ser Thr Ala Ser  
 195 200 205  
 Ser Pro Ala Pro Glu Glu Asn Ala Pro Ser Glu Gly Gln Arg Val Trp  
 210 215 220  
 Gly Gln Gly Gln Ser Pro Arg Pro Glu Asn Ser Leu Glu Arg Glu Glu  
 225 230 235 240  
 Met Gly Pro Val Pro Ala His Thr Asp Ala Phe Gln Asp Trp Gly Pro  
 245 250 255  
 Gly Ser Met Ala His Val Ser Val Val Pro Val Ser Ser Glu Gly Thr  
 260 265 270

PBD00056\_wo\_Seq.ST25.txt  
 Pro Ser Arg Glu Pro Val Ala Ser Gly Ser Trp Thr Pro Lys Ala Glu  
           275                                  280                                  285

Glu Pro Ile His Ala Thr Met Asp Pro Gln Arg Leu Gly Val Leu Ile  
           290                                  295                                  300

Thr Pro Val Pro Asp Ala Gln Ala Ala Thr Arg Arg Gln Ala Val Gly  
   305                                  310                                  315                                  320

Leu Leu Ala Phe Leu Gly Leu Leu Phe Cys Leu Gly Val Ala Met Phe  
                                   325                                  330                                  335

Thr Tyr Gln Ser Leu Gln Gly Cys Pro Arg Lys Met Ala Gly Glu Met  
                                   340                                  345                                  350

Ala Glu Gly Leu Arg Tyr Ile Pro Arg Ser Cys Gly Ser Asn Ser Tyr  
                                   355                                  360                                  365

Val Leu Val Pro Val  
       370

<210> 13

<211> 76

<212> PRT

<213> Homo sapiens

<400> 13

Gln Pro Val Gly Ile Asn Thr Ser Thr Thr Cys Cys Tyr Arg Phe Ile  
   1                                  5                                  10                                  15

Asn Lys Lys Ile Pro Lys Gln Arg Leu Glu Ser Tyr Arg Arg Thr Thr  
                                   20                                  25                                  30

Ser Ser His Cys Pro Arg Glu Ala Val Ile Phe Lys Thr Lys Leu Asp  
                                   35                                  40                                  45

Lys Glu Ile Cys Ala Asp Pro Thr Gln Lys Trp Val Gln Asp Phe Met  
   50                                  55                                  60

Lys His Leu Asp Lys Lys Thr Gln Thr Pro Lys Leu  
   65                                  70                                  75

<210> 14

<211> 33

<212> PRT

<213> Homo sapiens

PBD00056\_WO\_Seq.ST25.txt

<400> 14

Gln Pro Leu Pro Asp Cys Cys Arg Gln Lys Thr Cys Ser Cys Arg Leu  
1 5 10 15

Tyr Glu Leu Leu His Gly Ala Gly Asn His Ala Ala Gly Ile Leu Thr  
20 25 30

Leu

<210> 15

<211> 11

<212> PRT

<213> Homo sapiens

<400> 15

Arg Pro Lys Pro Gln Gln Phe Phe Gly Leu Met  
1 5 10

<210> 16

<211> 32

<212> PRT

<213> Artificial sequence

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<223> Synthetic peptide

<400> 16

Glu Val His His Gln Lys Leu Val Phe Phe Ala Glu Asp Val Gly Ser  
1 5 10 15

Asn Lys Gly Ala Ile Ile Gly Leu Met Val Gly Gly Val Val Ile Ala  
20 25 30

<210> 17

<211> 30

<212> PRT

<213> Artificial sequence

<220>

<223> Synthetic peptide



<400> 17

Glu Val His His Gln Lys Leu Val Phe Phe Ala Glu Asp Val Gly Ser  
1 5 10 15

Asn Lys Gly Ala Ile Ile Gly Leu Met Val Gly Gly Val Val  
20 25 30

<210> 18

<211> 34

<212> PRT

<213> Artificial sequence

<220>

<223> Synthetic peptide

<400> 18

Glu Ala Ser Asn Cys Phe Ala Ile Arg His Phe Glu Asn Lys Phe Ala  
1 5 10 15

Val Glu Thr Leu Ile Cys Ser Arg Thr Val Lys Lys Asn Ile Ile Glu  
20 25 30

Glu Asn

<210> 19

<211> 34

<212> PRT

<213> Artificial sequence

<220>

<223> Synthetic peptide

<400> 19

Glu Ala Ser Asn Cys Phe Ala Ile Arg His Phe Glu Asn Lys Phe Ala  
1 5 10 15

Val Glu Thr Leu Ile Cys Phe Asn Leu Phe Leu Asn Ser Gln Glu Lys  
20 25 30

His Tyr

<210> 20

<211> 5

PBD00056\_WO\_Seq.ST25.txt

<212> PRT

<213> Artificial sequence

<220>

<223> Synthetic peptide

<400> 20

Gln Tyr Asn Ala Asp  
1 5