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<110> Imperial Innovations Limited

<120> Method of Diagnosis and Prognosis

<130> 15631WO

<150> GB 1107118.0

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<170> PatentIn version 3.5

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<220>

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<223> AMIDATION

<400> 69

Tyr Asp His Phe

1

<210> 70

<211> 4

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<223> AMIDATION

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Tyr Glu Arg Phe

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<210> 71

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<213> artificial sequence

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<400> 71

Tyr Asp His Phe
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Tyr Asp His Phe

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<400> 73

Tyr Glu Arg Phe Gly Tyr Asp Arg Phe
1 5

<210> 74
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<222> (9)..(9)

<223> AMIDATION

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Tyr Asp His Phe Gly Tyr Asp His Phe

1

5

<210> 75

<211> 7

<212> PRT

<213> artificial sequence

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<223> AMIDATION

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Tyr Arg Phe Gly Tyr Arg Phe

1 5

<210> 76

<211> 9

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<400> 76

Tyr Asp His Phe Gly Tyr Asp His Phe
1 5

<210> 77
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<223> benzyloxycarbonyl-Tyr

<220>

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<223> AMIDATION

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Tyr Glu Arg Phe Gly Tyr Glu Arg Phe

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<210> 78

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<223> AMIDATION

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Tyr Asp His Phe Gly Tyr Asp His Phe
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<210> 79
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<223> ACETYLATION

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<222> (4)..(4)
<223> AMIDATION

<400> 81

Tyr Asp His Gln
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<210> 82
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<222> (4)..(4)

<223> AMIDATION

<400> 82

Tyr Asp His Ala

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<210> 83

<211> 4

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<223> AMIDATION

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Tyr Glu Lys Trp

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Trp Asp Arg Phe

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Tyr Asp His Trp

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Trp Asp His Trp

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Trp Glu His Trp

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<210> 93

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Tyr Glu Lys Trp

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Ala Glu Arg Phe

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Tyr Ala Arg Phe

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Tyr Glu Ala Phe
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Tyr Glu Arg Ala

1

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Pro Glu Arg Phe

1

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<211> 4

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<400> 107

Tyr Glu Pro Phe

1

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Tyr Pro Arg Phe
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Tyr Glu Arg Pro

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<220>

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<400> 110

Tyr Asp His Gln
1

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<220>
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<223> 4-methyl ester aspartic acid (Asp(OMe))

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<222> (4)..(4)
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<400> 111

Tyr Asp His Gln

1

<210> 112

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<220>

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<222> (4)..(4)

<223> AMIDATION

<400> 112

Tyr Glu Arg Phe

1

<210> 113

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Tyr Asp His Phe

1

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<222> (4)..(4)

<223> AMIDATION

<400> 114

Tyr Asp His Gln

1

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<222> (4)..(4)

<223> (3-Methoxy,4-hydroxy-benzoyl)-Tyr

<400> 115

Tyr Asp His Gln

1

<210> 116

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<213> artificial sequence

<220>

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<220>

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<223> (3-Methoxy,4-hydroxy-benzoyl)-Tyr

<220>

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<223> AMIDATION

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Tyr Asp Phe Gln

1

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Tyr Tyr Arg Phe

1

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Tyr Tyr Glu Arg Phe

1 5

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Tyr Asn Arg Phe

1

<210> 120

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<223> benzyloxycarbonyl-Tyr

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Tyr Asn Arg Phe

1

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1

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Tyr Met Arg Phe
1

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<223> AMIDATION

<400> 123

Tyr Asn Arg Phe

1

<210> 124

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<223> AMIDATION

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Tyr Asn Arg Phe

1

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Tyr Leu Arg Phe
1

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Tyr Leu Arg Phe

1

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<223> AMIDATION

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Tyr Asp His Gln

1

<210> 128

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<212> PRT

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<221> MOD_RES

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<223> AMIDATION

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Tyr Glu Arg Phe

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<220>
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<223> AMIDATION

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<223> AMIDATION

<400> 156

Gln Ala Ala Ala

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<211> 4

<212> PRT

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<223> residues 2, 3 and 4 may be substituted by any amino acid

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<223> AMIDATION

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<223> AMIDATION

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Ala Ala Ala Ala

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Tyr Asp His Gln

1

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<223> AMIDATION

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<223> AMIDATION

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Tyr Asp His Gln Lys
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Trp Asn Lys Trp

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Asp Asp Asp Asp Tyr

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5

<210> 227

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His Pro Phe His Leu

1

5