

(12) STANDARD PATENT
(19) AUSTRALIAN PATENT OFFICE

(11) Application No. **AU 2009305113 B2**

(54) Title
Dengue virus neutralizing antibodies and uses thereof

(51) International Patent Classification(s)
C07K 14/18 (2006.01) **C07K 16/10** (2006.01)
A61K 39/12 (2006.01) **A61P 31/14** (2006.01)

(21) Application No: **2009305113** (22) Date of Filing: **2009.10.13**

(87) WIPO No: **WO10/043977**

(30) Priority Data

(31) Number	(32) Date	(33) Country
61/104,911	2008.10.13	US

(43) Publication Date: **2010.04.22**

(44) Accepted Journal Date: **2015.07.16**

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(56) Related Art
CHAMBERS T.J. et al., Journal of Virology, 2003, Vol. 77, pages 3655-3668
GONCALVEZ A. P. et al., PNAS, 2007, Vol. 104, pages 9422-9427
PUTTIKHUNT C. et al., Journal of Medical Virology, 2008, Vol. 80, pages 125-133
WO 2004/067567 A2 (NOVARTIS AG) 12 August 2004
WO 2005/056600 A3 , 23 June 2005

(12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property Organization
International Bureau



(43) International Publication Date
22 April 2010 (22.04.2010)

(10) International Publication Number
WO 2010/043977 A3

(51) International Patent Classification:
C07K 16/10 (2006.01) A61P 31/14 (2006.01)

(21) International Application Number:
PCT/IB2009/007372

(22) International Filing Date:
13 October 2009 (13.10.2009)

(25) Filing Language: English

(26) Publication Language: English

(30) Priority Data:
61/104,911 13 October 2008 (13.10.2008) US

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(81) Designated States (unless otherwise indicated, for every kind of national protection available): AE, AG, AL, AM, AO, AT, AU, AZ, BA, BB, BG, BH, BR, BW, BY, BZ, CA, CH, CL, CN, CO, CR, CU, CZ, DE, DK, DM, DO,

DZ, EC, EE, EG, ES, FI, GB, GD, GE, GH, GM, GT, HN, HR, HU, ID, IL, IN, IS, JP, KE, KG, KM, KN, KP, KR, KZ, LA, LC, LK, LR, LS, LT, LU, LY, MA, MD, ME, MG, MK, MN, MW, MX, MY, MZ, NA, NG, NI, NO, NZ, OM, PE, PG, PH, PL, PT, RO, RS, RU, SC, SD, SE, SG, SK, SL, SM, ST, SV, SY, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VC, VN, ZA, ZM, ZW.

(84) Designated States (unless otherwise indicated, for every kind of regional protection available): ARIPO (BW, GH, GM, KE, LS, MW, MZ, NA, SD, SL, SZ, TZ, UG, ZM, ZW), Eurasian (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European (AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HR, HU, IE, IS, IT, LT, LU, LV, MC, MK, MT, NL, NO, PL, PT, RO, SE, SI, SK, SM, TR), OAPI (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG).

Published:

- with international search report (Art. 21(3))
- before the expiration of the time limit for amending the claims and to be republished in the event of receipt of amendments (Rule 48.2(h))
- with sequence listing part of description (Rule 5.2(a))

(88) Date of publication of the international search report:
30 September 2010

(54) Title: DENGUE VIRUS NEUTRALIZING ANTIBODIES AND USES THEREOF

(57) Abstract: The invention relates to antibodies and antigen binding fragments thereof and to cocktails of antibodies and antigen binding fragments that neutralize dengue virus infection without contributing to antibody-dependent enhancement of dengue virus infection. The invention also relates to immortalized B cells that produce, and to epitopes that bind to, such antibodies and antigen binding fragments. In addition, the invention relates to the use of the antibodies, antigen binding fragments, and epitopes in screening methods as well as in the diagnosis and therapy of dengue virus infection.



WO 2010/043977 A3

DENGUE VIRUS NEUTRALIZING ANTIBODIES AND USES THEREOF

This application claims priority to U.S. Provisional Application Serial No. 61/104,911, entitled "Dengue Virus Neutralizing Antibodies and Use Thereof," filed October 13, 2008, which is incorporated herein by reference in its entirety.

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BACKGROUND

Dengue viruses (DENV) are human pathogens with a significant threat to world health. These viruses are estimated to cause several hundred thousand cases of dengue fever, dengue hemorrhagic fever and dengue shock syndrome annually. There are four closely related serotypes of dengue viruses, DENV-1, DENV-2, DENV-3 and DENV-4, of the genus *Flavivirus*.
10 The four viruses are spread from human to human through the bite of *Aedes aegypti*, a highly urbanized mosquito species that has successfully resisted all attempts at eradication and control. Vaccination is considered to be the only efficient method of control of dengue. To this end, several tetravalent dengue candidate vaccines are in late stages of development.

A first infection with one Dengue virus serotype induces a life-long protective immunity
15 to the homologous serotype. However, there is no cross-protection against infection by a different serotype. Indeed, pre-existing immunity against one serotype is associated with increased risk for dengue infection and dengue hemorrhagic fever caused by a different serotype due to antibody-dependent enhancement (ADE) of infection. In ADE, antibodies raised by prior dengue infection or passively transferred from mother form infectious immune complexes that
20 attach to Fc-receptor-bearing cells in the mononuclear phagocyte lineage resulting in efficient infection.

Accordingly, there is a need for materials and methods for preventing dengue virus infection without increasing the risk of antibody-dependent enhancement of infection.

SUMMARY

25 The invention is based, in part, on the discovery of antibodies and cocktails of antibodies that neutralize dengue virus infection without contributing to antibody-dependent enhancement of dengue virus infection. Accordingly, in one aspect of the invention, the invention comprises a human antibody, an antibody variant, or an antigen binding fragment thereof, that neutralize a dengue virus, wherein the antibody, antibody variant, or antigen binding fragment does not
30 contribute to antibody-dependent enhancement of dengue virus infection. In one embodiment,

the invention comprises a human antibody, an antibody variant, or an antigen binding fragment thereof, that neutralize a dengue virus, wherein the antibody, antibody variant, or antigen binding fragment comprises a mutation in the Fc region, and wherein the mutation reduces binding of the antibody to an Fc receptor.

5 In another embodiment of the invention, the invention comprises a pharmaceutical composition comprising two or more human antibodies, or antigen binding fragments thereof. The antibodies or antigen binding fragments neutralize dengue virus serotypes DENV-1, DENV-2, DENV-3, and DENV-4 by binding at least two distinct epitopes on each dengue virus serotype. The antibodies of the pharmaceutical composition do not contribute to antibody-
10 dependent enhancement of dengue virus infection.

 In yet another embodiment, the invention comprises an antibody, or an antigen binding fragment thereof, comprising at least one complementarity determining region (CDR) sequence having the sequence of any one of SEQ ID NOs: 1-6, 17-22, 33-38, 49-54, 67-72, 83-88, 99, 100, 105-110, 121-123, 124, 125, 135-139, 149, 153-158, 169-174, 185-188, or 189, wherein the
15 antibody neutralizes dengue virus infection.

 In yet another embodiment, the invention comprises an antibody, or an antigen binding fragment thereof, wherein the antibody comprises a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 13 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 14; or a heavy chain variable region comprising the amino acid
20 sequence of SEQ ID NO: 29 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 30; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 45 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 46; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 61 and a light chain variable region comprising the amino acid
25 sequence of SEQ ID NO: 62; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 65 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 62; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 79 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 80; or a heavy chain variable region comprising the amino acid
30 sequence of SEQ ID NO: 95 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 96; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 95 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 103; or a heavy chain variable region comprising the amino acid

sequence of SEQ ID NO: 117 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 118; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 131 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 132; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 145 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 146; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 151 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 146; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 165 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 166; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 181 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 182; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 195 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 196, wherein the antibody neutralizes dengue virus infection.

15 In a further embodiment, the invention comprises a recombinant antibody, antibody variant, or antigen binding fragment thereof, that can neutralize a dengue virus. The recombinant antibody, antibody variant, or antigen binding fragment does not contribute to antibody-dependent enhancement of dengue virus infection.

20 In another aspect, the invention comprises a nucleic acid molecule comprising a polynucleotide encoding an antibody or antibody fragment of the invention that neutralizes dengue virus infection. In yet another aspect, the invention comprises a cell expressing an antibody of the invention. In still another aspect, the invention comprises an isolated or purified immunogenic polypeptide comprising an epitope that binds to an antibody of the invention.

25 The invention also comprises a pharmaceutical composition comprising an antibody, an antibody variant or an antigen binding fragment of the invention, a nucleic acid of the invention, or an immunogenic polypeptide of the invention and a pharmaceutically acceptable diluent or carrier and, optionally, an agent useful for extending the half life of the antibody or antigen binding fragment thereof.

30 In another aspect of the invention, the invention provides a method of inhibiting or preventing dengue virus infection or a dengue virus-related disease or a method of treating dengue virus infection or a dengue virus-related disease. The method comprises administering to a subject in need thereof, a therapeutically effective amount of at least one antibody, antibody variant, antigen binding fragment, or a pharmaceutical composition of the invention.

In yet another aspect of the invention, the invention comprises a method of screening for polypeptides that can induce or reveal an immune response against dengue virus, comprising screening polypeptide libraries using an antibody, an antibody fragment or variant of the invention.

5 In yet another aspect of the invention, the invention comprises a method of monitoring the quality of anti-dengue virus vaccines. The method comprises using an antibody, an antibody variant, or an antigen binding fragment thereof of the invention to check that the antigen of the vaccine contains the specific epitope in the correct conformation.

10 In a further aspect of the invention, the invention comprises a vaccine comprising an epitope which specifically binds to an antibody, an antibody fragment or variant of the invention.

Use of an antibody of the invention, or an antigen binding fragment thereof, a nucleic acid of the invention, an immunogenic polypeptide of the invention, or a pharmaceutical composition of the invention (i) in the manufacture of a medicament for the treatment of dengue virus infection, (ii) in a vaccine, or (iii) in diagnosis of dengue virus infection is also
15 contemplated to be within the scope of the invention. Further, use of an antibody of the invention, or an antigen binding fragment thereof, for monitoring the quality of anti-DENV vaccines by checking that the antigen of said vaccine contains the specific epitope in the correct conformation is also contemplated to be within the scope of the invention.

In a further aspect, the invention comprises an epitope which specifically binds to an
20 antibody of any one of the invention, or an antigen binding fragment thereof, for use (i) in therapy, (ii) in the manufacture of a medicament for treating dengue virus infection, (iii) as a vaccine, or (iv) in screening for ligands able to neutralise dengue virus infection.

BRIEF DESCRIPTION OF FIGURES

Figure 1. VERO cells and K562 cells were used in virus neutralization and enhancement
25 assays with each serotype of dengue virus using wild-type anti-dengue virus antibodies. Antibodies to dengue virus inhibit infection of the target virus on VERO cells in a dose-dependent manner. On K562 cells, the antibodies lead to a dose dependent antibody-dependent enhancement (ADE) of infection.

Figure 2. Anti-dengue virus antibodies that have a CH2 L4A and L5A substitution
30 (LALA variants) in the heavy chain neutralize target virus infection on VERO cells as did the unmodified antibodies. However, the LALA variants completely abolished the antibody-dependent enhancement of infection by the target virus on K562 cells.

DETAILED DESCRIPTION OF THE INVENTION

The invention is based on the discovery of antibodies and cocktails of antibodies that neutralize dengue virus (DENV) infection without contributing to antibody-dependent enhancement (ADE) of dengue virus infection. In one aspect of the invention, the invention
5 comprises a human antibody, a variant antibody, or an antigen binding fragment thereof, that neutralizes a dengue virus without contributing to antibody-dependent enhancement of dengue virus infection. The antibodies or antibody fragments can neutralize more than one dengue virus serotype, for example, 2, 3 or all 4 dengue virus serotypes DENV-1, DENV-2, DENV-3, and DENV-4.

10 The invention also comprises a pharmaceutical composition comprising, for example, an antibody cocktail that comprises two or more human antibodies, antibody variants or antigen binding fragments thereof. The pharmaceutical compositions of the invention comprising a cocktail of human antibodies, antibody fragments or variants neutralize all four dengue virus serotypes, *i.e.*, DENV-1, DENV-2, DENV-3, and DENV-4. In one embodiment, the cocktail of
15 antibodies, antibody fragments or variants neutralize dengue virus by binding at least two distinct epitopes on each dengue virus serotype. It is noted that the antibodies, variants and fragments of the pharmaceutical composition do not contribute to antibody-dependent enhancement of dengue virus infection. In one embodiment, the cocktail comprises two antibodies, fragments or variants thereof. In another embodiment, the cocktail comprises three antibodies, fragments or variants
20 thereof. In yet another embodiment, the cocktail comprises more than 3 antibodies, *e.g.*, 4, 5, 6, 7 or 8 antibodies.

As used herein, the terms “fragment,” “antibody fragment,” and “antigen binding fragment” are used interchangeably to refer to any fragment of an antibody of the invention that retains the antigen-binding activity of the antibody. Exemplary antibody fragments include, but
25 are not limited to, Fab, Fab', F(ab')₂, Fv, and scFv fragments.

The terms “mutation,” and “substitution” are used interchangeably to refer to a change in one or more nucleic acid or amino acid residues.

As used herein, the terms “variant,” and “antibody variant” are used interchangeably to refer to any variant of an antibody of the invention that retains the antigen-binding activity of the
30 antibodies. The term variant includes antibodies that comprise mutations and/or substitutions. Exemplary antibody variants include, but are not limited to, those that have an L to A substitution at position CH2 4, 5, or both.

Antibodies of the invention

The invention provides antibodies that neutralize dengue virus, but do not contribute to ADE of dengue virus infection. A “neutralizing antibody” is one that can neutralize the ability of a pathogen to initiate and/or perpetuate an infection in a host. The antibodies of the invention
5 are able to neutralize one or more dengue virus serotypes DENV-1, DENV-2, DENV-3, and DENV-4. In one embodiment, the antibody of the invention neutralizes more than one, *e.g.*, 2, 3, or all 4 dengue virus serotypes. In another embodiment, a pharmaceutical composition comprising two or more antibodies, antibody fragments or variants can neutralize all 4 dengue virus serotypes. In yet another embodiment, the pharmaceutical composition comprising two or
10 more antibodies, antibody fragments or variants neutralizes dengue virus infection by targeting two distinct epitopes on each dengue virus serotype. These antibodies, antigen binding fragment and variants can be used as prophylactic or therapeutic agents upon appropriate formulation, or as a diagnostic tool, as described herein.

The antibodies of the invention may be monoclonal, for example, human monoclonal
15 antibodies, or recombinant antibodies. The invention also provides fragments of the antibodies of the invention, particularly fragments that retain the antigen-binding activity of the antibodies. Although the specification, including the claims, may, in some places, refer explicitly to antibody fragment(s), variant(s) and/or derivative(s) of antibodies, it is understood that the term “antibody” or “antibody of the invention” includes all categories of antibodies, namely, antibody
20 fragment(s), variant(s) and derivative(s) of antibodies.

Without being bound to any theory, it is believed that antibody-dependent enhancement of dengue virus infection is brought about by the binding of the Fc region of the antibody, in particular, the Fc region of the heavy chain of an IgG molecule, to an Fc receptor, *e.g.*, an Fc γ receptor on a host cell. The invention, on the other hand, provides antibodies, including IgG
25 molecules, that have reduced binding to the Fc receptors (FcR). In one embodiment, the antibody of the invention comprises one or more mutations in the Fc region. The mutation(s) may be any mutation that reduces binding of the antibody to an Fc receptor. In one embodiment, the Fc region of an antibody of the invention comprises a substitution at positions CH2 4, 5, or both. In general, the amino acid at positions 4 and 5 of CH2 of the wild-type IgG1 and IgG3 is a
30 leucine (“L”). In one embodiment, the antibodies of the invention comprise an amino acid at position CH2 4, 5, or both, that is not an L. In another embodiment, the antibodies of the invention comprise an alanine (“A”) at position CH2 4, or 5, or both. An antibody comprising a CH2 L4A and an L5A substitution is referred to herein as a “LALA” variant.

Alternatively, the invention provides antibody fragments that do not comprise an Fc region and thus do not bind to an FcR. Exemplary antibody fragments include, but are not limited to, Fab, Fab', F(ab')₂, Fv and scFv.

The sequences of the heavy chains and light chains of several exemplary antibodies of the invention, each comprising three CDRs on the heavy chain and three CDRs on the light chain have been determined. The position of the CDR amino acids are defined according to the IMGT numbering system [1, 2, 3]. The sequences of the CDRs, heavy chains, light chains as well as the sequences of the nucleic acid molecules encoding the CDRs, heavy chains, light chains of many exemplary antibodies of the invention are disclosed in the sequence listing. Table 1 provides the SEQ ID NOs for the amino acid sequences of the six CDRs, the variable region of the heavy and light chains, respectively, of exemplary antibodies of the invention. Table 2 provides the SEQ ID NOs for the sequences of the nucleic acid molecules encoding the CDRs, heavy chains and light chains of exemplary antibodies of the invention.

Table 1. Amino Acid SEQ IDs for Antibody CDRs, Heavy and Light Chains

Antibody	CDRs	Heavy Chain Variable Region	Light Chain Variable Region
HMB-DV-1	1-6	13	14
HMB-DV-2	17-22	29	30
HMB-DV-3	33-38	45	46
HMB-DV-4	49-54	61, 65	62
HMB-DV-5	67-72	79	80
HMB-DV-6	83-88	95	96
HMB-DV-7	83-85, 99, 53, 100	95	103
HMB-DV-8	105-110	117	118
HMB-DV-9	121-123, 70 124, 125	131	132
HMB-DV-10	135-139, 109	145	146
HMB-DV-11	149, 136-139, 109	151	146
HMB-DV-12	153-158	165	166
HMB-DV-13	169-174	181	182
HMB-DV-14	185-188, 37, 189	195	196

Table 2. Nucleic Acid SEQ IDs for Antibody CDRs, Heavy and Light Chains

Antibody	CDRs	Heavy Chain Variable Region	Light Chain Variable Region
HMB-DV-1	7-12	15	16
HMB-DV-2	23-28	31	32
HMB-DV-3	39-44	47	48
HMB-DV-4	55-60	63, 66	64
HMB-DV-5	73-78	81	82
HMB-DV-6	89-94	97	98
HMB-DV-7	89-91, 101, 59, 102	97	104
HMB-DV-8	111-116	119	120

HMB-DV-9	126-128, 76, 129, 130	133	134
HMB-DV-10	140-143, 115, 144	147	148
HMB-DV-11	150, 141-143, 115, 144	152	148
HMB-DV-12	159-164	167	168
HMB-DV-13	175-180	183	184
HMB-DV-14	190-193, 43, 194	197	198

In one embodiment, the antibodies or antigen-binding fragments of the invention comprise one or more heavy or light chain CDRs of the exemplary antibodies of the invention. In an exemplary embodiment, the antibodies or antigen-binding fragments of the invention neutralize dengue virus infection and comprise at least one CDR sequence having the sequence of any one of SEQ ID NOs: 1-6, 17-22, 33-38, 49-54, 67-72, 83-88, 99, 100, 105-110, 121-123, 124, 125, 135-139, 149, 153-158, 169-174, 185-188, or 189.

In another embodiment, the antibodies, antibody variants or antigen binding fragments of the invention comprise a heavy chain comprising an amino acid sequence of one or more of SEQ ID NOs: 1-3, 17-19, 33-35, 49-51, 67-69, 83-85, 105-107, 121-123, 135-137, 149, 153-155, 169-171, or 185-187. In yet another embodiment, the antibodies, antibody variants or antigen binding fragments of the invention comprise a heavy chain CDR1 selected from the group consisting of SEQ ID NOs: 1, 17, 33, 49, 67, 83, 105, 121, 135, 149, 153, 169, and 185; a heavy chain CDR2 selected from the group consisting of SEQ ID NOs: 2, 18, 34, 50, 68, 84, 106, 122, 136, 154, 170, and 186; and a heavy chain CDR3 selected from the group consisting of SEQ ID NOs: 3, 19, 35, 51, 69, 85, 107, 123, 137, 155, 171, and 187.

For example, the antibodies of the invention comprise a heavy chain comprising SEQ ID NO: 1 for CDRH1, SEQ ID NO: 2 for CDRH2, SEQ ID NO: 3 for CDRH3; SEQ ID NO: 17 for CDRH1, SEQ ID NO: 18 for CDRH2 and SEQ ID NO: 19 for CDRH3; SEQ ID NO: 33 for CDRH1, SEQ ID NO: 34 for CDRH2 and SEQ ID NO: 35 for CDRH3; SEQ ID NO: 49 for CDRH1, SEQ ID NO: 50 for CDRH2 and SEQ ID NO: 51 for CDRH3; SEQ ID NO: 67 for CDRH1, SEQ ID NO: 68 for CDRH2 and SEQ ID NO: 69 for CDRH3; SEQ ID NO: 83 for CDRH1, SEQ ID NO: 84 for CDRH2 and SEQ ID NO: 85 for CDRH3; SEQ ID NO: 105 for CDRH1, SEQ ID NO: 106 for CDRH2 and SEQ ID NO: 107 for CDRH3; SEQ ID NO: 121 for CDRH1, SEQ ID NO: 122 for CDRH2 and SEQ ID NO: 123 for CDRH3; SEQ ID NO: 135 for CDRH1, SEQ ID NO: 136 for CDRH2 and SEQ ID NO: 137 for CDRH3; SEQ ID NO: 149 for CDRH1, SEQ ID NO: 136 for CDRH2 and SEQ ID NO: 137 for CDRH3; SEQ ID NO: 153 for CDRH1, SEQ ID NO: 154 for CDRH2 and SEQ ID NO: 155 for CDRH3; SEQ ID NO: 169 for CDRH1, SEQ ID NO: 170 for CDRH2 and SEQ ID NO: 171 for CDRH3; and SEQ ID NO: 185 for CDRH1, SEQ ID NO: 186 for CDRH2 and SEQ ID NO: 187 for CDRH3.

In yet another embodiment, the antibodies, antibody variants or antibody fragments of the invention comprise a light chain comprising an amino acid sequence of one or more of SEQ ID NOs: 4-6, 20-22, 36-38, 52-54, 70-72, 86-88, 99, 100, 108-110, 124, 125, 138, 139, 156-158, 172-174, 188, or 189. In a further embodiment, the antibodies, antibody variants or antibody fragments of the invention comprise a light chain CDR1 selected from the group consisting of SEQ ID NOs: 4, 20, 36, 52, 70, 86, 99, 108, 138, 156, 172, and 188; a light chain CDR2 selected from the group consisting of SEQ ID NOs: 5, 21, 37, 53, 71, 87, 109, 124, 157, and 173; and a light chain CDR3 selected from the group consisting of SEQ ID NOs: 6, 22, 38, 54, 72, 88, 100, 110, 125, 139, 158, 174, and 189.

For example, the antibodies of the invention comprise a light chain comprising SEQ ID NO: 4 for CDRL1, SEQ ID NO: 5 for CDRL2; SEQ ID NO: 6 for CDRL3; SEQ ID NO: 20 for CDRL1, SEQ ID NO: 21 for CDRL2; SEQ ID NO: 22 for CDRL3; SEQ ID NO: 36 for CDRL1, SEQ ID NO: 37 for CDRL2; SEQ ID NO: 38 for CDRL3; SEQ ID NO: 52 for CDRL1, SEQ ID NO: 53 for CDRL2; SEQ ID NO: 54 for CDRL3; SEQ ID NO: 70 for CDRL1, SEQ ID NO: 71 for CDRL2; SEQ ID NO: 72 for CDRL3; SEQ ID NO: 86 for CDRL1, SEQ ID NO: 87 for CDRL2; SEQ ID NO: 88 for CDRL3; SEQ ID NO: 99 for CDRL1, SEQ ID NO: 53 for CDRL2; SEQ ID NO: 100 for CDRL3; SEQ ID NO: 108 for CDRL1, SEQ ID NO: 109 for CDRL2; SEQ ID NO: 110 for CDRL3; SEQ ID NO: 70 for CDRL1, SEQ ID NO: 124 for CDRL2; SEQ ID NO: 125 for CDRL3; SEQ ID NO: 138 for CDRL1, SEQ ID NO: 109 for CDRL2; SEQ ID NO: 139 for CDRL3; SEQ ID NO: 156 for CDRL1, SEQ ID NO: 157 for CDRL2; SEQ ID NO: 158 for CDRL3; SEQ ID NO: 172 for CDRL1, SEQ ID NO: 173 for CDRL2; SEQ ID NO: 174 for CDRL3; and SEQ ID NO: 188 for CDRL1, SEQ ID NO: 37 for CDRL2; SEQ ID NO: 189 for CDRL3.

In one embodiment, an antibody of the invention, or antigen binding fragment thereof, comprises all of the CDRs of antibody HMB-DV-1 as listed in Table 1, and neutralizes dengue virus infection in a human host. In another embodiment, an antibody of the invention, or antigen binding fragment thereof, comprises all of the CDRs of antibody HMB-DV-2 as listed in Table 1, and neutralizes dengue virus infection in a human host. In another embodiment, an antibody of the invention, or antigen binding fragment thereof, comprises all of the CDRs of antibody HMB-DV-3 as listed in Table 1, and neutralizes dengue virus infection in a human host. In yet another embodiment, an antibody of the invention, or antigen binding fragment thereof, comprises all of the CDRs of antibody HMB-DV-4 as listed in Table 1, and neutralizes dengue virus infection in a human host. In yet another embodiment, an antibody of the invention, or antigen binding fragment thereof, comprises all of the CDRs of antibody HMB-DV-5 as listed in Table 1, and

neutralizes dengue virus infection in a human host. In yet another embodiment, an antibody of the invention, or antigen binding fragment thereof, comprises all of the CDRs of antibody HMB-DV-6 as listed in Table 1, and neutralizes dengue virus infection in a human host. In yet another embodiment, an antibody of the invention, or antigen binding fragment thereof, comprises all of the CDRs of antibody HMB-DV-7 as listed in Table 1, and neutralizes dengue virus infection in a human host.

In a further embodiment, an antibody of the invention, or antigen binding fragment thereof, comprises all of the CDRs of antibody HMB-DV-8 as listed in Table 1, and neutralizes dengue virus infection in a human host. In another embodiment, an antibody of the invention, or antigen binding fragment thereof, comprises all of the CDRs of antibody HMB-DV-9 as listed in Table 1, and neutralizes dengue virus infection in a human host. In still another embodiment, an antibody of the invention, or antigen binding fragment thereof, comprises all of the CDRs of antibody HMB-DV-10 as listed in Table 1, and neutralizes dengue virus infection in a human host. In yet another embodiment, an antibody of the invention, or antigen binding fragment thereof, comprises all of the CDRs of antibody HMB-DV-11 as listed in Table 1, and neutralizes dengue virus infection in a human host. In yet another embodiment, an antibody of the invention, or antigen binding fragment thereof, comprises all of the CDRs of antibody HMB-DV-12 as listed in Table 1, and neutralizes dengue virus infection in a human host. In yet another embodiment, an antibody of the invention, or antigen binding fragment thereof, comprises all of the CDRs of antibody HMB-DV-13 as listed in Table 1, and neutralizes dengue virus infection in a human host. In yet another embodiment, an antibody of the invention, or antigen binding fragment thereof, comprises all of the CDRs of antibody HMB-DV-14 as listed in Table 1, and neutralizes dengue virus infection in a human host.

In still another embodiment, the antibodies of the invention comprise a heavy chain with an amino acid sequence that is at least 70%, at least 75%, at least 80%, at least 85%, at least 90%, at least 95%, at least 98%, or at least 99% identical to those of SEQ ID NOs: 13, 29, 45, 61, 65, 79, 95, 117, 131, 145, 151, 165, 181, or 195. In yet another embodiment, the antibodies of the invention comprise a light chain with an amino acid sequence that is at least 70%, at least 75%, at least 80%, at least 85%, at least 90%, at least 95%, at least 98%, or at least 99% identical to those of SEQ ID NOs: 14, 30, 46, 62, 80, 96, 103, 118, 132, 146, 166, 182, or 196.

In a further embodiment, the antibodies, antibody variants or antibody fragments of the invention comprise a heavy chain variable region comprising the amino acid sequence of any one of SEQ ID NOs: 13, 29, 45, 61, 65, 79, 95, 117, 131, 145, 151, 165, 181, or 195, and a light chain

variable region comprising the amino acid sequence of any one of SEQ ID NOs: 14, 30, 46, 62, 80, 96, 103, 118, 132, 146, 166, 182, or 196.

In yet another embodiment, the antibodies, antibody variants or antibody fragments of the invention neutralize dengue virus infection and comprise a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 13 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 14; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 29 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 30; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 45 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 46; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 61 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 62; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 65 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 62; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 79 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 80; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 95 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 96; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 95 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 103; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 117 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 118; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 131 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 132; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 145 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 146; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 151 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 146; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 165 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 166; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 181 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 182; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 195 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 196.

Methods for chain replacement and for CDR grafting are well known in the art. Originally these methods were developed to humanize non-human antibodies (generally mouse antibodies) or to select human antibody counterparts having equivalent bioactivity to the non-

human antibodies. These methods include replacement techniques where only one of the CDRs, for example, the CDR3s, of the non-human antibody are retained and the remainder of the V-region, including the framework and the other two CDRs, for example, the CDRs 1 and 2, are individually replaced in steps performed sequentially (e.g. U.S. Patent Application No.

5 20030166871; Rader, et al., *Proc Natl Acad Sci USA* 95:8910-15, 1998; Steinberg, et al., *J Biol Chem* 275:36073-78, 2000; Rader, et al., *J Biol Chem* 275:13668-76, 2000).

In addition, methods of creating antibodies with the binding specificities of a reference antibody for a target antigen are described in Patent Application No. WO05/069,970. The methods include transferring, from the reference antibody to a recipient antibody or antibody
10 fragment, the minimal essential binding specificity of the reference antibody. Examples of regions that can be transferred include, but are not limited to, the transfer of a single CDR segment, for example a CDR3 segment, from the heavy and/or from the light chain, or a D segment, or a CDR3-FR4 segment, or any CDR3-FR4 segment that comprises the minimal essential binding specificity determinant. Antibodies created using these methods retain the
15 binding specificity, and often affinity, of the reference antibody.

The antibodies, antibody variants or antibody fragments of the invention include antibodies that comprise, *inter alia*, one or more CDRs, a heavy chain or a light chain of an exemplary antibody of the invention and retain their specificity and ability to neutralize dengue virus infection.

20 Exemplary antibodies of the invention include, but are not limited to, HMB-DV1, HMB-DV2, HMB-DV3, HMB-DV4, HMB-DV5, HMB-DV6, HMB-DV7, HMB-DV8, HMB-DV9, HMB-DV10, HMB-DV11, HMB-DV12, HMB-DV13, and HMB-DV14.

Variants of HMB-DV4 consist of a heavy chain variants having amino acid sequence recited in SEQ ID NO: 61 and SEQ ID NO: 65, and a light chain having the amino acid sequence
25 recited in SEQ ID NO: 62. The nucleic acid sequences encoding the heavy chain variants are recited in SEQ ID NO: 63 and SEQ ID NO: 66. The nucleic acid encoding the light chain is recited in SEQ ID NO: 64. Thus, antibodies comprising the HMB-DV4 variant heavy chains (SEQ ID NOs: 61, 65) and light chain (SEQ ID NO: 62) are included within the scope of the invention.

30 As used herein, the term "HMB-DV4" is used to refer to any and/or all variants of HMB-DV4, for example, those with heavy chains corresponding to SEQ ID NOs: 61 and 65 and light chain corresponding to SEQ ID NO: 62.

In one embodiment, an antibody cocktail of the invention comprises two or more antibodies selected from the group consisting of HMB-DV1, HMB-DV2, HMB-DV3, HMB-DV4, HMB-DV5, HMB-DV6, HMB-DV7, HMB-DV8, HMB-DV9, HMB-DV10, HMB-DV11, HMB-DV12, HMB-DV13, and HMB-DV14. In another embodiment, a cocktail of the invention
5 comprises three antibodies selected from the group consisting of HMB-DV1, HMB-DV2, HMB-DV3, HMB-DV4, HMB-DV5, HMB-DV6, HMB-DV7, HMB-DV8, HMB-DV9, HMB-DV10, HMB-DV11, HMB-DV12, HMB-DV13, and HMB-DV14. In yet another embodiment, an antibody cocktail of the invention comprises more than three, for example, 4, 5, 6, 7, or 8 antibodies selected from the group consisting of HMB-DV1, HMB-DV2, HMB-DV3, HMB-
10 DV4, HMB-DV5, HMB-DV6, HMB-DV7, HMB-DV8, HMB-DV9, HMB-DV10, HMB-DV11, HMB-DV12, HMB-DV13, and HMB-DV14. In an exemplary embodiment, a cocktail of the invention comprises HMB-DV5, HMB-DV6, and HMB-DV8.

The invention further comprises an antibody, or fragment thereof, that binds to an epitope capable of binding to an antibody of the invention. The invention also comprises an antibody or
15 an antibody fragment that competes with an antibody of the invention.

In another aspect, the invention also includes nucleic acid sequences encoding part or all of the light and heavy chains and CDRs of the antibodies of the present invention. In one embodiment, nucleic acid sequences according to the invention include nucleic acid sequences having at least 70%, at least 75%, at least 80%, at least 85%, at least 90%, at least 95%, at least
20 98%, or at least 99% identity to the nucleic acid encoding a heavy or light chain of an antibody of the invention. In another embodiment, a nucleic acid sequence of the invention has the sequence of a nucleic acid encoding a heavy or light chain CDR of an antibody of the invention. For example, a nucleic acid sequence according to the invention comprises a sequence that is at least 70%, at least 75%, at least 80%, at least 85%, at least 90%, at least 95%, at least 98%, or at
25 least 99% identical to the nucleic acid sequences of SEQ ID NOs: 7-12, 23-28, 39-44, 55-60, 73-78, 89-94, 101, 102, 111-116, 126-128, 129, 130, 140-144, 150, 159-164, 175-180, and 190-194.

Due to the redundancy of the genetic code, variants of these sequences will exist that encode the same amino acid sequences. These variants are included within the scope of the invention.

30 Variant antibodies are also included within the scope of the invention. Thus, variants of the sequences recited in the application are also included within the scope of the invention. Such variants include natural variants generated by somatic mutation *in vivo* during the immune response or *in vitro* upon culture of immortalized B cell clones. Alternatively, variants may arise

due to the degeneracy of the genetic code, as mentioned above or may be produced due to errors in transcription or translation. Variants may also be introduced to modify the antibody effector function, for instance in the Fc region to reduce the binding of the antibody to an Fc receptor.

Further variants of the antibody sequences having improved affinity and/or potency may be obtained using methods known in the art and are included within the scope of the invention. For example, amino acid substitutions may be used to obtain antibodies with further improved affinity. Alternatively, codon optimisation of the nucleotide sequence may be used to improve the efficiency of translation in expression systems for the production of the antibody. Further, polynucleotides comprising a sequence optimized for antibody specificity or neutralizing activity by the application of a directed evolution method to any of the nucleic acid sequences of the invention are also within the scope of the invention.

In one embodiment variant antibody sequences may share 70% or more (*i.e.* 75%, 80%, 85%, 90%, 95%, 97%, 98%, 99% or more) amino acid sequence identity with the sequences recited in the application. In some embodiments such sequence identity is calculated with regard to the full length of the reference sequence (*i.e.* the sequence recited in the application). In some further embodiments, percentage identity, as referred to herein, is as determined using BLAST version 2.1.3 using the default parameters specified by the NCBI (the National Center for Biotechnology Information; <http://www.ncbi.nlm.nih.gov/>) [Blosum 62 matrix; gap open penalty=11 and gap extension penalty=1].

Further included within the scope of the invention are vectors, for example expression vectors, comprising a nucleic acid sequence according to the invention. Cells transformed with such vectors are also included within the scope of the invention. Examples of such cells include but are not limited to, eukaryotic cells, *e.g.* yeast cells, animal cells or plant cells. In one embodiment the cells are mammalian, *e.g.* human, CHO, HEK293T, PER.C6, NS0, myeloma or hybridoma cells.

The invention also relates to monoclonal antibodies that bind to an epitope capable of binding an antibody of the invention. In one embodiment, the invention includes a monoclonal antibody that binds to an epitope capable of binding a monoclonal antibody selected from the group consisting of HMB-DV1, HMB-DV2, HMB-DV3, HMB-DV4, HMB-DV5, HMB-DV6, HMB-DV7, HMB-DV8, HMB-DV9, HMB-DV10, HMB-DV11, HMB-DV12, HMB-DV13, and HMB-DV14.

Monoclonal and recombinant antibodies are particularly useful in identification and purification of the individual polypeptides or other antigens against which they are directed. The

antibodies of the invention have additional utility in that they may be employed as reagents in immunoassays, radioimmunoassays (RIA) or enzyme-linked immunosorbent assays (ELISA). In these applications, the antibodies can be labelled with an analytically-detectable reagent such as a radioisotope, a fluorescent molecule or an enzyme. The antibodies may also be used for the
5 molecular identification and characterisation (epitope mapping) of antigens.

Antibodies of the invention will typically be glycosylated. N-linked glycans attached to the C_H2 domain of a heavy chain, for instance, can influence C1q and FcR binding, with aglycosylated antibodies having lower affinity for these receptors. The glycan structure can also affect activity *e.g.* differences in complement-mediated cell death may be seen depending on the
10 number of galactose sugars (0, 1 or 2) at the terminus of a glycan's biantennary chain. An antibody's glycans preferably do not lead to a human immunogenic response after administration.

Antibodies of the invention can be coupled to a drug for delivery to a treatment site or coupled to a detectable label to facilitate imaging of a site comprising cells of interest, such as
15 cells infected with dengue virus. Methods for coupling antibodies to drugs and detectable labels are well known in the art, as are methods for imaging using detectable labels. Labelled antibodies may be employed in a wide variety of assays, employing a wide variety of labels. Detection of the formation of an antibody-antigen complex between an antibody of the invention and an epitope of interest (a DENV epitope) can be facilitated by attaching a detectable
20 substance to the antibody. Suitable detection means include the use of labels such as radionuclides, enzymes, coenzymes, fluorescers, chemiluminescers, chromogens, enzyme substrates or co-factors, enzyme inhibitors, prosthetic group complexes, free radicals, particles, dyes, and the like. Examples of suitable enzymes include horseradish peroxidase, alkaline phosphatase, β -galactosidase, or acetylcholinesterase; examples of suitable prosthetic group
25 complexes include streptavidin/biotin and avidin/biotin; examples of suitable fluorescent materials include umbelliferone, fluorescein, fluorescein isothiocyanate, rhodamine, dichlorotriazinylamine fluorescein, dansyl chloride or phycoerythrin; an example of a luminescent material is luminol; examples of bioluminescent materials include luciferase, luciferin, and aequorin; and examples of suitable radioactive material include ¹²⁵I, ¹³¹I, ³⁵S, or ³H.
30 Such labeled reagents may be used in a variety of well-known assays, such as radioimmunoassays, enzyme immunoassays, *e.g.*, ELISA, fluorescent immunoassays, and the like.

An antibody according to the invention may be conjugated to a therapeutic moiety such as a cytotoxin, a therapeutic agent, or a radioactive metal ion or radioisotope. Examples of radioisotopes include, but are not limited to, I-131, I-123, I-125, Y-90, Re-188, Re-186, At-211, Cu-67, Bi-212, Bi-213, Pd-109, Tc-99, In-111, and the like. Such antibody conjugates can be used for modifying a given biological response; the drug moiety is not to be construed as limited to classical chemical therapeutic agents. For example, the drug moiety may be a protein or polypeptide possessing a desired biological activity. Such proteins may include, for example, a toxin such as abrin, ricin A, pseudomonas exotoxin, calicheamicin bacterial toxin, or diphtheria toxin.

Techniques for conjugating such therapeutic moiety to antibodies are well known. See, for example, Arnon *et al.* (1985) "Monoclonal Antibodies for Immunotargeting of Drugs in Cancer Therapy," in *Monoclonal Antibodies and Cancer Therapy*, ed. Reisfeld *et al.* (Alan R. Liss, Inc.), pp. 243-256; ed. Hellstrom *et al.* (1987) "Antibodies for Drug Delivery," in *Controlled Drug Delivery*, ed. Robinson *et al.* (2d ed; Marcel Dekker, Inc.), pp. 623-653; Thorpe (1985) "Antibody Carriers of Cytotoxic Agents in Cancer Therapy: A Review," in *Monoclonal Antibodies '84: Biological and Clinical Applications*, ed. Pinchera *et al.* pp. 475-506 (Editrice Kurtis, Milano, Italy, 1985); "Analysis, Results, and Future Prospective of the Therapeutic Use of Radiolabeled Antibody in Cancer Therapy," in *Monoclonal Antibodies for Cancer Detection and Therapy*, ed. Baldwin *et al.* (Academic Press, New York, 1985), pp. 303-316; and Thorpe *et al.* (1982) *Immunol. Rev.* 62:119-158.

Alternatively, an antibody can be conjugated to a second antibody to form an antibody heteroconjugate as described in reference 4. In addition, linkers may be used between the labels and the antibodies of the invention [5]. Antibodies or, antigen-binding fragments thereof may be directly labelled with radioactive iodine, indium, yttrium, or other radioactive particle known in the art [6]. Treatment may consist of a combination of treatment with conjugated and non-conjugated antibodies administered simultaneously or subsequently [7, 8].

Antibodies of the invention may also be attached to a solid support.

Additionally, antibodies of the invention, or functional antibody fragments thereof, can be chemically modified by covalent conjugation to a polymer to, for example, increase their circulating half-life, for example. Examples of polymers, and methods to attach them to peptides, are shown in references 9-12. In some embodiments the polymers may be selected from polyoxyethylated polyols and polyethylene glycol (PEG). PEG is soluble in water at room temperature and has the general formula: $R(O-CH_2-CH_2)_nO-R$ where R can be hydrogen, or a

protective group such as an alkyl or alkanol group. In one embodiment the protective group may have between 1 and 8 carbons. In a further embodiment the protective group is methyl. The symbol n is a positive integer. In one embodiment n is between 1 and 1,000. In another embodiment n is between 2 and 500. In one embodiment the PEG has an average molecular weight between 1,000 and 40,000. In a further embodiment the PEG has a molecular weight between 2,000 and 20,000. In yet a further embodiment the PEG has a molecular weight of between 3,000 and 12,000. In one embodiment PEG has at least one hydroxy group. In another embodiment the PEG has a terminal hydroxy group. In yet another embodiment it is the terminal hydroxy group which is activated to react with a free amino group on the inhibitor. However, it will be understood that the type and amount of the reactive groups may be varied to achieve a covalently conjugated PEG/antibody of the present invention.

Antibodies of the invention can be modified by introducing random amino acid mutations into particular region of the CH2 or CH3 domain of the heavy chain in order to alter their binding affinity for FcRn and/or their serum half-life in comparison to the unmodified antibodies. Examples of such modifications include, but are not limited to, substitutions of at least one amino acid from the heavy chain constant region selected from the group consisting of amino acid residues 250, 314, and 428.

Water-soluble polyoxyethylated polyols are also useful in the present invention. They include polyoxyethylated sorbitol, polyoxyethylated glucose, polyoxyethylated glycerol (POG), and the like. In one embodiment, POG is used. Without being bound by any theory, because the glycerol backbone of polyoxyethylated glycerol is the same backbone occurring naturally in, for example, animals and humans in mono-, di-, triglycerides, this branching would not necessarily be seen as a foreign agent in the body. In some embodiments POG has a molecular weight in the same range as PEG. The structure for POG is shown in reference 13, and a discussion of POG/IL-2 conjugates is found in reference 9.

Another drug delivery system that can be used for increasing circulatory half-life is the liposome. Methods of preparing liposome delivery systems are discussed in references 14, 15 and 16. Other drug delivery systems are known in the art and are described in, for example, references 17 and 18.

Antibodies of the invention may be provided in purified form. Typically, the antibody will be present in a composition that is substantially free of other polypeptides *e.g.* where less than 90% (by weight), usually less than 60% and more usually less than 50% of the composition is made up of other polypeptides.

Antibodies of the invention may be immunogenic in non-human (or heterologous) hosts *e.g.* in mice. In particular, the antibodies may have an idiotope that is immunogenic in non-human hosts, but not in a human host. Antibodies of the invention for human use include those that cannot be easily isolated from hosts such as mice, goats, rabbits, rats, non-primate mammals, *etc.* and cannot generally be obtained by humanisation or from xeno-mice.

Antibodies of the invention can be of any isotype (*e.g.* IgA, IgG, IgM *i.e.* an α , γ or μ heavy chain), but will generally be IgG. Within the IgG isotype, antibodies may be IgG1, IgG2, IgG3 or IgG4 subclass. In one embodiment, the antibody is IgG1. Antibodies of the invention may have a κ or a λ light chain.

Included within the scope of the invention are DENV-neutralizing recombinant or engineered bispecific antibody molecules or antigen binding fragments thereof. Such antibodies and fragments may comprise a first binding site for an epitope on a first Dengue virus serotype and a second binding site for a second epitope on the same dengue virus serotype or on a different, for example, a second, third or fourth, dengue virus serotype. The variable domains of the respective binding sites can be formed as immunoglobulin isotypes of the invention or as heterodimeric Fab, Fab', F(ab')₂, ScFv or diabodies that can be linked together via one or more peptide linkers.

Production of antibodies

Monoclonal antibodies according to the invention can be made by any method known in the art. The general methodology for making monoclonal antibodies using hybridoma technology is well known [19, 20]. Preferably, the alternative EBV immortalisation method described in reference 21 is used.

Using the method described in reference 21, B cells producing the antibody of the invention can be transformed with EBV in the presence of a polyclonal B cell activator. Transformation with EBV is a standard technique and can easily be adapted to include polyclonal B cell activators.

Additional stimulants of cellular growth and differentiation may optionally be added during the transformation step to further enhance the efficiency. These stimulants may be cytokines such as IL-2 and IL-15. In one aspect, IL-2 is added during the immortalisation step to further improve the efficiency of immortalisation, but its use is not essential.

The immortalised B cells produced using these methods can then be cultured using methods known in the art and antibodies isolated therefrom.

The antibodies of the invention can also be made by culturing single plasma cells in microwell culture plates using the method described in UK Patent Application 0819376.5. Further, from single plasma cell cultures, RNA can be extracted and single cell PCR can be performed using methods known in the art. The VH and VL regions of the antibodies can be amplified by RT-PCR, sequenced and cloned into an expression vector that is then transfected into HEK293T cells or other host cells. The cloning of nucleic acid in expression vectors, the transfection of host cells, the culture of the transfected host cells and the isolation of the produced antibody can be done using any methods known to one of skill in the art.

Monoclonal antibodies may be further purified, if desired, using filtration, centrifugation and various chromatographic methods such as HPLC or affinity chromatography. Techniques for purification of monoclonal antibodies, including techniques for producing pharmaceutical-grade antibodies, are well known in the art.

Fragments of the monoclonal antibodies of the invention can be obtained from the monoclonal antibodies by methods that include digestion with enzymes, such as pepsin or papain, and/or by cleavage of disulfide bonds by chemical reduction. Alternatively, fragments of the monoclonal antibodies can be obtained by cloning and expression of part of the sequences of the heavy or light chains. Antibody "fragments" may include Fab, Fab', F(ab')₂ and Fv fragments. The invention also encompasses single-chain Fv fragments (scFv) derived from the heavy and light chains of a monoclonal antibody of the invention *e.g.* the invention includes a scFv comprising the CDRs from an antibody of the invention. Also included are heavy or light chain monomers and dimers as well as single chain antibodies, *e.g.* single chain Fv in which the heavy and light chain variable domains are joined by a peptide linker.

Standard techniques of molecular biology may be used to prepare DNA sequences coding for the antibodies or fragments or variants of the antibodies of the present invention. Desired DNA sequences may be synthesised completely or in part using oligonucleotide synthesis techniques. Site-directed mutagenesis and polymerase chain reaction (PCR) techniques may be used as appropriate.

Any suitable host cell/vector system may be used for expression of the DNA sequences encoding the antibody molecules of the present invention or fragments thereof. Bacterial, for example *E. coli*, and other microbial systems may be used, in part, for expression of antibody fragments such as Fab and F(ab')₂ fragments, and especially Fv fragments and single chain antibody fragments, for example, single chain Fvs. Eukaryotic, *e.g.* mammalian, host cell expression systems may be used for production of larger antibody molecules, including complete

antibody molecules. Suitable mammalian host cells include CHO, HEK293T, PER.C6, NS0, myeloma or hybridoma cells.

The present invention also provides a process for the production of an antibody of the invention comprising culturing a host cell comprising a vector of the present invention under
5 conditions suitable for leading to expression of protein from DNA encoding the antibody of the present invention, and isolating the antibody molecule.

The antibody molecule may comprise only a heavy or light chain polypeptide, in which case only a heavy chain or light chain polypeptide coding sequence needs to be used to transfect the host cells. For production of products comprising both heavy and light chains, the cell line
10 may be transfected with two vectors, a first vector encoding a light chain polypeptide and a second vector encoding a heavy chain polypeptide. Alternatively, a single vector may be used, the vector including sequences encoding light chain and heavy chain polypeptides.

Alternatively, antibodies according to the invention may be produced by i) expressing a nucleic acid sequence according to the invention in a cell, and ii) isolating the expressed
15 antibody product. Additionally, the method may include iii) purifying the antibody.

Screening and isolation of B cells

Transformed B cells may be screened for those producing antibodies of the desired antigen specificity, and individual B cell clones may then be produced from the positive cells.

The screening step may be carried out by ELISA, by staining of tissues or cells (including
20 infected or transfected cells), a neutralisation assay or one of a number of other methods known in the art for identifying desired antigen specificity. The assay may select on the basis of simple antigen recognition, or may select on the additional basis of a desired function *e.g.* to select neutralizing antibodies rather than just antigen-binding antibodies, to select antibodies that can change characteristics of targeted cells, such as their signalling cascades, their shape, their
25 growth rate, their capability of influencing other cells, their response to the influence by other cells or by other reagents or by a change in conditions, their differentiation status, *etc.*

The cloning step for separating individual clones from the mixture of positive cells may be carried out using limiting dilution, micromanipulation, single cell deposition by cell sorting or another method known in the art.

30 The immortalised B cell clones of the invention can be used in various ways *e.g.* as a source of monoclonal antibodies, as a source of nucleic acid (DNA or mRNA) encoding a monoclonal antibody of interest, for research, *etc.*

The invention provides a composition comprising immortalised B memory cells, wherein the cells produce antibodies that neutralize one or more dengue virus serotypes, and wherein the antibodies are produced at ≥ 5 pg per cell per day. The invention also provides a composition comprising clones of an immortalised B memory cell, wherein the clones produce a monoclonal antibody that neutralizes one or more dengue virus serotypes, and wherein the antibody is produced at ≥ 5 pg per cell per day.

Exemplary immortalised B cell clone according to the invention include, but are not limited to, HMB-DV1, HMB-DV2, HMB-DV3, HMB-DV4, HMB-DV5, HMB-DV6, HMB-DV7, HMB-DV8, HMB-DV9, HMB-DV10, HMB-DV11, HMB-DV12, HMB-DV13, and HMB-DV14.

Epitopes

As mentioned above, the antibodies of the invention can be used to map the epitopes to which they bind. The epitopes recognised by the antibodies of the present invention may have a number of uses. The epitope and mimotopes thereof in purified or synthetic form can be used to raise immune responses (*i.e.* as a vaccine, or for the production of antibodies for other uses) or for screening patient serum for antibodies that immunoreact with the epitope or mimotopes thereof. In one embodiment such an epitope or mimotope, or antigen comprising such an epitope or mimotope may be used as a vaccine for raising an immune response. The antibodies and antigen binding fragments of the invention can also be used in a method of monitoring the quality of vaccines. In particular the antibodies can be used to check that the antigen in a vaccine contains the specific epitope in the correct conformation.

The epitope may also be useful in screening for ligands that bind to said epitope. Such ligands, include but are not limited to antibodies, including those from camels, sharks and other species, fragments of antibodies, peptides, phage display technology products, aptamers, adnectins, synthetic compounds, or fragments of other viral or cellular proteins, that may block the epitope and so prevent infection. Such ligands are encompassed within the scope of the invention.

Recombinant expression

The immortalised B memory cells of the invention may also be used as a source of nucleic acid for the cloning of antibody genes for subsequent recombinant expression. Expression from recombinant sources is more common for pharmaceutical purposes than

expression from B cells or hybridomas *e.g.* for reasons of stability, reproducibility, culture ease, *etc.*

Thus the invention provides a method for preparing a recombinant cell, comprising the steps of: (i) obtaining one or more nucleic acids (*e.g.* heavy and/or light chain genes) from the B
5 cell clone that encodes the antibody of interest; and (ii) inserting the nucleic acid into an expression host in order to permit expression of the antibody of interest in that host.

Similarly, the invention provides a method for preparing a recombinant cell, comprising the steps of: (i) sequencing nucleic acid(s) from the B cell clone that encodes the antibody of interest; and (ii) using the sequence information from step (i) to prepare nucleic acid(s) for
10 insertion into an expression host in order to permit expression of the antibody of interest in that host. The nucleic acid may, but need not, be manipulated between steps (i) and (ii) to introduce restriction sites, to change codon usage, to optimise transcription and/or translation regulatory sequences, and/or to modify effector function.

The invention also provides a method of preparing a recombinant cell, comprising the
15 step of transforming a host cell with one or more nucleic acids that encode a monoclonal antibody of interest, wherein the nucleic acids are nucleic acids that were derived from an immortalised B cell clone of the invention. Thus the procedures for first preparing the nucleic acid(s) and then using it to transform a host cell can be performed at different times by different people in different places (*e.g.*, in different countries).

20 These recombinant cells of the invention can then be used for expression and culture purposes. They are particularly useful for expression of antibodies for large-scale pharmaceutical production. They can also be used as the active ingredient of a pharmaceutical composition. Any suitable culture techniques can be used, including but not limited to static culture, roller bottle culture, ascites fluid, hollow-fiber type bioreactor cartridge, modular
25 minifermenter, stirred tank, microcarrier culture, ceramic core perfusion, *etc.*

Methods for obtaining and sequencing immunoglobulin genes from B cells are well known in the art (*e.g.*, see reference 22).

The expression host is preferably a eukaryotic cell, including yeast and animal cells, particularly mammalian cells (*e.g.* CHO cells, NS0 cells, human cells such as PER.C6 [Crucell;
30 reference 23] or HKB-11 [Bayer; references 24 & 25] cells, myeloma cells [26 & 27], *etc.*), as well as plant cells. Preferred expression hosts can glycosylate the antibody of the invention, particularly with carbohydrate structures that are not themselves immunogenic in humans. In one embodiment the expression host may be able to grow in serum-free media. In a further

embodiment the expression host may be able to grow in culture without the presence of animal-derived products.

The expression host may be cultured to give a cell line.

The invention provides a method for preparing one or more nucleic acid molecules (*e.g.* heavy and light chain genes) that encode an antibody of interest, comprising the steps of:

5 (i) preparing an immortalised B cell clone according to the invention; (ii) obtaining from the B cell clone nucleic acid that encodes the antibody of interest. The invention also provides a method for obtaining a nucleic acid sequence that encodes an antibody of interest, comprising the steps of: (i) preparing an immortalised B cell clone according to the invention; (ii)

10 sequencing nucleic acid from the B cell clone that encodes the antibody of interest.

The invention also provides a method of preparing nucleic acid molecule(s) that encodes an antibody of interest, comprising the step of obtaining the nucleic acid from a B cell clone that was obtained from a transformed B cell of the invention. Thus the procedures for first obtaining the B cell clone and then preparing nucleic acid(s) from it can be performed at very different

15 times by different people in different places (*e.g.* in different countries).

The invention provides a method for preparing an antibody (*e.g.* for pharmaceutical use), comprising the steps of: (i) obtaining and/or sequencing one or more nucleic acids (*e.g.* heavy and light chain genes); (ii) using the sequence information from step (i) to prepare nucleic acid(s) for insertion into an expression host in order to permit expression of the antibody of interest in

20 that host; (iii) culturing or sub-culturing the expression host under conditions where the antibody of interest is expressed; and, optionally, (iv) purifying the antibody of the interest. The nucleic acid can, but need not be, obtained and/or sequenced from a B cell clone expressing the antibody of interest. In one embodiment, the nucleic acid from step (i) may, optionally be modified so as to introduce desired substitutions in the amino acid sequence of the antibody.

25 The invention also provides a method of preparing an antibody comprising the steps of: culturing or sub-culturing an expression host cell population under conditions where the antibody of interest is expressed and, optionally, purifying the antibody of the interest, wherein said expression host cell population has been prepared by (i) providing nucleic acid(s) encoding an antibody of interest; (ii) inserting the nucleic acid(s) into an expression host that can express the

30 antibody of interest, and (iii) culturing or sub-culturing expression hosts comprising said inserted nucleic acids to produce said expression host cell population.

Pharmaceutical compositions

The invention provides a pharmaceutical composition containing the antibodies and/or antibody fragments of the invention and/or nucleic acid encoding such antibodies and/or immortalised B cells that express such antibodies and/or the epitopes recognised by the antibodies of the invention. A pharmaceutical composition may also contain a pharmaceutically acceptable carrier to allow administration. The carrier should not itself induce the production of antibodies harmful to the individual receiving the composition and should not be toxic. Suitable carriers may be large, slowly metabolised macromolecules such as proteins, polypeptides, liposomes, polysaccharides, polylactic acids, polyglycolic acids, polymeric amino acids, amino acid copolymers and inactive virus particles.

Pharmaceutically acceptable salts can be used, for example mineral acid salts, such as hydrochlorides, hydrobromides, phosphates and sulphates, or salts of organic acids, such as acetates, propionates, malonates and benzoates.

Pharmaceutically acceptable carriers in therapeutic compositions may additionally contain liquids such as water, saline, glycerol and ethanol. Additionally, auxiliary substances, such as wetting or emulsifying agents or pH buffering substances, may be present in such compositions. Such carriers enable the pharmaceutical compositions to be formulated as tablets, pills, dragees, capsules, liquids, gels, syrups, slurries and suspensions, for ingestion by the patient.

Within the scope of the invention, forms of administration may include those forms suitable for parenteral administration, *e.g.* by injection or infusion, for example by bolus injection or continuous infusion. Where the product is for injection or infusion, it may take the form of a suspension, solution or emulsion in an oily or aqueous vehicle and it may contain formulatory agents, such as suspending, preservative, stabilising and/or dispersing agents. Alternatively, the antibody molecule may be in dry form, for reconstitution before use with an appropriate sterile liquid.

Once formulated, the compositions of the invention can be administered directly to the subject. In one embodiment the compositions are adapted for administration to human subjects.

The pharmaceutical compositions of this invention may be administered by any number of routes including, but not limited to, oral, intravenous, intramuscular, intra-arterial, intramedullary, intraperitoneal, intrathecal, intraventricular, transdermal, transcutaneous, topical, subcutaneous, intranasal, enteral, sublingual, intravaginal or rectal routes. Hyposprays may also be used to administer the pharmaceutical compositions of the invention. Typically, the

therapeutic compositions may be prepared as injectables, either as liquid solutions or suspensions. Solid forms suitable for solution in, or suspension in, liquid vehicles prior to injection may also be prepared.

Direct delivery of the compositions will generally be accomplished by injection, subcutaneously, intraperitoneally, intravenously or intramuscularly, or delivered to the interstitial space of a tissue. Dosage treatment may be a single dose schedule or a multiple dose schedule. Known antibody-based pharmaceuticals provide guidance relating to frequency of administration *e.g.* whether a pharmaceutical should be delivered daily, weekly, monthly, *etc.* Frequency and dosage may also depend on the severity of symptoms.

Compositions of the invention may be prepared in various forms. For example, the compositions may be prepared as injectables, either as liquid solutions or suspensions. Solid forms suitable for solution, or suspension, in liquid vehicles prior to injection can also be prepared (*e.g.* a lyophilised composition, like Synagis™ and Herceptin™, for reconstitution with sterile water containing a preservative). The composition may be prepared for topical administration *e.g.* as an ointment, cream or powder. The composition may be prepared for oral administration *e.g.* as a tablet or capsule, as a spray, or as a syrup (optionally flavoured). The composition may be prepared for pulmonary administration *e.g.* as an inhaler, using a fine powder or a spray. The composition may be prepared as a suppository or pessary. The composition may be prepared for nasal, aural or ocular administration *e.g.* as drops. The composition may be in kit form, designed such that a combined composition is reconstituted just prior to administration to a patient. For example, a lyophilised antibody can be provided in kit form with sterile water or a sterile buffer.

It will be appreciated that the active ingredient in the composition will be an antibody molecule, an antibody fragment or variants and derivatives thereof. As such, it will be susceptible to degradation in the gastrointestinal tract. Thus, if the composition is to be administered by a route using the gastrointestinal tract, the composition will need to contain agents which protect the antibody from degradation but which release the antibody once it has been absorbed from the gastrointestinal tract.

A thorough discussion of pharmaceutically acceptable carriers is available in Gennaro (2000) *Remington: The Science and Practice of Pharmacy*, 20th edition, ISBN: 0683306472.

Pharmaceutical compositions of the invention generally have a pH between 5.5 and 8.5, in some embodiments this may be between 6 and 8, and in further embodiments about 7. The pH may be maintained by the use of a buffer. The composition may be sterile and/or pyrogen free.

The composition may be isotonic with respect to humans. In one embodiment pharmaceutical compositions of the invention are supplied in hermetically-sealed containers.

Pharmaceutical compositions will include a therapeutically effective amount of one or more antibodies of the invention and/or a polypeptide comprising an epitope that binds an antibody of the invention *i.e.* an amount that is sufficient to treat, ameliorate, or prevent a desired disease or condition, or to exhibit a detectable therapeutic effect. Therapeutic effects also include reduction in physical symptoms. The precise effective amount for any particular subject will depend upon their size and health, the nature and extent of the condition, and the therapeutics or combination of therapeutics selected for administration. The effective amount for a given situation is determined by routine experimentation and is within the judgment of a clinician. For purposes of the present invention, an effective dose will generally be from about 0.01mg/kg to about 50mg/kg, or about 0.05 mg/kg to about 10 mg/kg of the compositions of the present invention in the individual to which it is administered. Known antibody-based pharmaceuticals provide guidance in this respect *e.g.*, Herceptin™ is administered by intravenous infusion of a 21 mg/ml solution, with an initial loading dose of 4mg/kg body weight and a weekly maintenance dose of 2mg/kg body weight; Rituxan™ is administered weekly at 375mg/m²; *etc.*

In one embodiment pharmaceutical compositions can include more than one (*e.g.* 2, 3, 4, 5, 6, 7, 8, *etc.*) antibody of the invention. In another embodiment the composition comprises two or more (*e.g.* 2, 3, 4, 5, *etc.*) antibodies, wherein the first antibody is specific for a first DENV epitope, and the second antibody is specific for a second DENV epitope. In yet another embodiment, the pharmaceutical composition comprises three antibodies of the invention. In another embodiment, the composition comprises two or more (*e.g.* 2, 3, 4, 5, *etc.*) antibodies, that together neutralise more than one dengue virus serotype. In yet another embodiment, the two or more antibodies of the invention together neutralise all four dengue virus serotypes, DENV-1, DENV-2, DENV-3 and DENV-4. In a further embodiment two or more antibodies of the invention together neutralise all four dengue virus serotypes by binding at least two distinct epitopes on each dengue virus serotype.

Exemplary antibodies of the invention for use in a pharmaceutical composition that neutralize a dengue virus without contributing to antibody-dependent enhancement of dengue virus infection include, but are not limited to, HMB-DV1, HMB-DV2, HMB-DV3, HMB-DV4, HMB-DV5, HMB-DV6, HMB-DV7, HMB-DV8, HMB-DV9, HMB-DV10, HMB-DV11, HMB-DV12, HMB-DV13, and HMB-DV14.

In one embodiment, a pharmaceutical composition includes two exemplary antibodies of the invention, for example, HMB-DV3 and HMB-DV7; HMB-DV3 and HMB-DV9; HMB-DV3 and HMB-DV12; HMB-DV3 and HMB-DV14; HMB-DV6 and HMB-DV7; HMB-DV6 and HMB-DV8. In another embodiment, a pharmaceutical composition includes three exemplary
5 antibodies of the invention, for example, HMB-DV2, HMB-DV3 and HMB-DV6; HMB-DV2, HMB-DV6 and HMB-DV8; HMB-DV2, HMB-DV8 and HMB-DV9; HMB-DV2, HMB-DV8 and HMB-DV12; HMB-DV2, HMB-DV8 and HMB-DV14; HMB-DV5, HMB-DV6 and HMB-DV8. Based on the teachings herein, one of skill in the art can determine other combinations of antibodies for use in a pharmaceutical composition.

10 In one embodiment, the invention provides a pharmaceutical composition comprising the antibody HMB-DV1 or an antigen binding fragment thereof, and a pharmaceutically acceptable diluent or carrier. In another embodiment, the invention provides a pharmaceutical composition comprising the antibody HMB-DV2 or an antigen binding fragment thereof, and a pharmaceutically acceptable diluent or carrier. In another embodiment, the invention provides a
15 pharmaceutical composition comprising the antibody HMB-DV3 or an antigen binding fragment thereof, and a pharmaceutically acceptable diluent or carrier. In another embodiment, the invention provides a pharmaceutical composition comprising the antibody HMB-DV4 or an antigen binding fragment thereof, and a pharmaceutically acceptable diluent or carrier. In yet another embodiment, the invention provides a pharmaceutical composition comprising the
20 antibody HMB-DV5 or an antigen binding fragment thereof, and a pharmaceutically acceptable diluent or carrier. In another embodiment, the invention provides a pharmaceutical composition comprising the antibody HMB-DV6 or an antigen binding fragment thereof, and a pharmaceutically acceptable diluent or carrier. In another embodiment, the invention provides a pharmaceutical composition comprising the antibody HMB-DV7 or an antigen binding fragment
25 thereof, and a pharmaceutically acceptable diluent or carrier.

In yet another embodiment, the invention provides a pharmaceutical composition comprising the antibody HMB-DV8 or an antigen binding fragment thereof, and a pharmaceutically acceptable diluent or carrier. In another embodiment, the invention provides a pharmaceutical composition comprising the antibody HMB-DV9 or an antigen binding fragment
30 thereof, and a pharmaceutically acceptable diluent or carrier. In another embodiment, the invention provides a pharmaceutical composition comprising the antibody HMB-DV10 or an antigen binding fragment thereof, and a pharmaceutically acceptable diluent or carrier. In yet another embodiment, the invention provides a pharmaceutical composition comprising the antibody HMB-DV11 or an antigen binding fragment thereof, and a pharmaceutically acceptable

diluent or carrier. In another embodiment, the invention provides a pharmaceutical composition comprising the antibody HMB-DV12 or an antigen binding fragment thereof, and a pharmaceutically acceptable diluent or carrier. In another embodiment, the invention provides a pharmaceutical composition comprising the antibody HMB-DV13 or an antigen binding
5 fragment thereof, and a pharmaceutically acceptable diluent or carrier. In yet another embodiment, the invention provides a pharmaceutical composition comprising the antibody HMB-DV14 or an antigen binding fragment thereof, and a pharmaceutically acceptable diluent or carrier.

Antibodies of the invention may be administered (either combined or separately) with
10 other therapeutics *e.g.* with chemotherapeutic compounds, with radiotherapy, *etc.* Preferred therapeutic compounds include anti-viral compounds. Such combination therapy provides an additive or synergistic improvement in therapeutic efficacy relative to the individual therapeutic agents when administered alone. The term “synergy” is used to describe a combined effect of two or more active agents that is greater than the sum of the individual effects of each respective
15 active agent. Thus, where the combined effect of two or more agents results in “synergistic inhibition” of an activity or process, it is intended that the inhibition of the activity or process is greater than the sum of the inhibitory effects of each respective active agent. The term “synergistic therapeutic effect” refers to a therapeutic effect observed with a combination of two or more therapies wherein the therapeutic effect (as measured by any of a number of parameters)
20 is greater than the sum of the individual therapeutic effects observed with the respective individual therapies.

In compositions of the invention that include antibodies of the invention, the antibodies may make up at least 50% by weight (*e.g.* 60%, 70%, 75%, 80%, 85%, 90%, 95%, 97%, 98%, 99% or more) of the total protein in the composition. The antibodies are thus in purified form.

25 The invention provides a method of preparing a pharmaceutical, comprising the steps of: (i) preparing an antibody of the invention; and (ii) admixing the purified antibody with one or more pharmaceutically-acceptable carriers.

The invention also provides a method of preparing a pharmaceutical, comprising the step of admixing an antibody with one or more pharmaceutically-acceptable carriers, wherein the
30 antibody is a monoclonal antibody that was obtained from a transformed B cell of the invention. Thus the procedures for first obtaining the monoclonal antibody and then preparing the pharmaceutical can be performed at very different times by different people in different places (*e.g.* in different countries).

As an alternative to delivering antibodies for therapeutic purposes, it is possible to deliver nucleic acid (typically DNA) that encodes the monoclonal antibody (or active fragment thereof) of interest to a subject, such that the nucleic acid can be expressed in the subject *in situ* to provide a desired therapeutic effect. Suitable gene therapy and nucleic acid delivery vectors are known in the art.

Compositions of the invention may be immunogenic compositions, and in some embodiments may be vaccine compositions comprising an antigen comprising a DENV epitope. Vaccines according to the invention may either be prophylactic (*i.e.* to prevent infection) or therapeutic (*i.e.* to treat infection).

Compositions may include an antimicrobial, particularly if packaged in a multiple dose format. Compositions may comprise detergent *e.g.* a Tween (polysorbate), such as Tween 80. Detergents are generally present at low levels *e.g.* <0.01%. Compositions may include sodium salts (*e.g.* sodium chloride) to give tonicity. A concentration of 10 ± 2 mg/ml NaCl is typical.

Compositions may comprise a sugar alcohol (*e.g.* mannitol) or a disaccharide (*e.g.* sucrose or trehalose) *e.g.* at around 15-30 mg/ml (*e.g.* 25 mg/ml), particularly if they are to be lyophilised or if they include material which has been reconstituted from lyophilised material. The pH of a composition for lyophilisation may be adjusted to around 6.1 prior to lyophilisation.

The compositions of the invention may also comprise one or more immunoregulatory agents. In one embodiment, one or more of the immunoregulatory agents include(s) an adjuvant.

Medical treatments and uses

The antibodies, antigen binding fragments, derivatives and variants thereof, or the cocktails and pharmaceutical compositions of the invention can be used for the treatment of DENV infection, for the prevention of DENV infection or for the diagnosis of DENV infection.

Methods of diagnosis may include contacting an antibody or an antibody fragment with a sample. Such samples may be tissue samples taken from, for example, salivary glands, lung, liver, pancreas, kidney, ear, eye, placenta, alimentary tract, heart, ovaries, pituitary, adrenals, thyroid, brain or skin. The methods of diagnosis may also include the detection of an antigen/antibody complex.

The invention therefore provides (i) an antibody, an antibody fragment, or variants and derivatives thereof according to the invention, (ii) an immortalised B cell clone according to the invention, (iii) an epitope capable of binding an antibody of the invention or (iv) a ligand,

preferably an antibody, capable of binding an epitope that binds an antibody of the invention for use in therapy.

Also provided is a method of treating a subject comprising administering to that subject (i) an antibody, an antibody fragment, variants and derivatives thereof, or a pharmaceutical composition according to the invention, or, a ligand, preferably an antibody, capable of binding an epitope that binds an antibody of the invention.

The invention also provides the use of (i) an antibody, an antibody fragment, or variants and derivatives thereof according to the invention, (ii) an immortalised B cell clone according to the invention, (iii) an epitope capable of binding an antibody of the invention, or (iv) a ligand, preferably an antibody, that binds to an epitope capable of binding an antibody of the invention, in the manufacture of a medicament for the prevention or treatment of DENV infection.

The invention provides a pharmaceutical composition for use as a medicament for the prevention or treatment of DENV infection. It also provides the use of an antibody of the invention and/or a protein comprising an epitope to which such an antibody binds in the manufacture of a medicament for treatment of a patient and/or diagnosis in a patient. It also provides a method for treating a subject, *e.g.*, a human subject. The method comprises the step of administering to the subject a therapeutically effective dose of a composition of the invention. One way of checking efficacy of therapeutic treatment involves monitoring disease symptoms after administration of the composition of the invention. Treatment can be a single dose schedule or a multiple dose schedule.

In one embodiment, an antibody, antibody fragment, antibody variant, epitope or pharmaceutical composition according to the invention is administered to a subject in need of such treatment. Such a subject includes, but is not limited to, one who is particularly at risk of or susceptible to DENV infection.

Antibodies of the invention can be used in passive immunisation. Antibodies and fragments or variants thereof, or a nucleic acid encoding an antibody or an antibody fragment or variant as described in the present invention may also be used in a kit for the diagnosis of dengue virus infection.

Epitopes capable of binding an antibody of the invention, *e.g.*, the monoclonal antibodies HMB-DV1, HMB-DV2, HMB-DV3, HMB-DV4, HMB-DV5, HMB-DV6, HMB-DV7, HMB-DV8, HMB-DV9, HMB-DV10, HMB-DV11, HMB-DV12, HMB-DV13, and HMB-DV14, may be used in a kit for monitoring the efficacy of vaccination procedures by detecting the presence of protective anti-DENV antibodies.

Antibodies, antibody fragments, or variants and derivatives thereof, as described in the present invention may also be used in a kit for monitoring vaccine manufacture with the desired immunogenicity.

The invention also provides a method of preparing a pharmaceutical composition,
5 comprising the step of admixing a monoclonal antibody with one or more pharmaceutically-acceptable carriers, wherein the monoclonal antibody is a monoclonal antibody that was obtained from an expression host of the invention. Thus the procedures for first obtaining the monoclonal antibody (*e.g.* expressing it and/or purifying it) and then admixing it with the pharmaceutical carrier(s) can be performed at very different times by different people in different places (*e.g.* in
10 different countries).

Starting with a transformed B cell of the invention, various steps of culturing, sub-culturing, cloning, sub-cloning, sequencing, nucleic acid preparation *etc.* can be performed in order to perpetuate the antibody expressed by the transformed B cell, with optional optimisation at each step. In a preferred embodiment, the above methods further comprise
15 techniques of optimisation (*e.g.* affinity maturation or optimisation) applied to the nucleic acids encoding the antibody. The invention encompasses all cells, nucleic acids, vectors, sequences, antibodies *etc.* used and prepared during such steps.

In all these methods, the nucleic acid used in the expression host may be manipulated to insert, delete or amend certain nucleic acid sequences. Changes from such manipulation include,
20 but are not limited to, changes to introduce restriction sites, to amend codon usage, to add or optimise transcription and/or translation regulatory sequences, *etc.* It is also possible to change the nucleic acid to alter the encoded amino acids. For example, it may be useful to introduce one or more (*e.g.* 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, *etc.*) amino acid substitutions, deletions and/or insertions into the antibody's amino acid sequence. Such point mutations can modify effector functions,
25 antigen-binding affinity, post-translational modifications, immunogenicity, *etc.*, can introduce amino acids for the attachment of covalent groups (*e.g.* labels) or can introduce tags (*e.g.* for purification purposes). Mutations can be introduced in specific sites or can be introduced at random, followed by selection (*e.g.* molecular evolution). For instance, one or more nucleic acids encoding any of the CDR regions, heavy chain variable regions or light chain variable
30 regions of antibodies of the invention can be randomly or directionally mutated to introduce different properties in the encoded amino acids. Such changes can be the result of an iterative process wherein initial changes are retained and new changes at other nucleotide positions are introduced. Moreover, changes achieved in independent steps may be combined. Different

properties introduced into the encoded amino acids may include, but are not limited to, enhanced affinity.

General

The term “comprising” encompasses “including” as well as “consisting” *e.g.* a composition “comprising” X may consist exclusively of X or may include something additional *e.g.* X + Y.

The word “substantially” does not exclude “completely” *e.g.* a composition which is “substantially free” from Y may be completely free from Y. Where necessary, the word “substantially” may be omitted from the definition of the invention.

The term “about” in relation to a numerical value x means, for example, $x \pm 10\%$.

The term “disease” as used herein is intended to be generally synonymous, and is used interchangeably with, the terms “disorder” and “condition” (as in medical condition), in that all reflect an abnormal condition of the human or animal body or of one of its parts that impairs normal functioning, is typically manifested by distinguishing signs and symptoms, and causes the human or animal to have a reduced duration or quality of life.

As used herein, reference to “treatment” of a patient is intended to include prevention and prophylaxis as well as therapy. The term “patient” means all mammals including humans. Generally, the patient is a human.

EXAMPLES

Exemplary embodiments of the present invention are provided in the following examples. The following examples are presented only by way of illustration and to assist one of ordinary skill in using the invention. The examples are not intended in any way to limit the scope of the invention.

Example 1. Cloning of B cells and screening for identification of Dengue virus specific Abs.

Memory B cells were isolated from the blood of DENV immune donors and immortalized using EBV and CpG as described in reference. Briefly, IgG⁺ memory B cells were isolated using CD22 beads, followed by removal of IgM⁺, IgD⁺ IgA⁺ B cells using specific antibodies and cell sorting. The sorted cells (IgG⁺) were immortalized with EBV in the presence of CpG 2006 and irradiated allogeneic mononuclear cells. Replicate cultures each containing 30-50 memory B cells were set up in several 96 well U-bottom plates. After two weeks the culture supernatants were collected and tested for their capacity to stain C6/36 cells infected with DENV

of serotypes 1, 2, 3 or 4 by immunofluorescence analysis and/or to bind to recombinant DENV1-4 E2 proteins by ELISA. Supernatants were tested for their capacity to neutralize DENV infection of either VERO cells or DC-SIGN-transfected Raji cells and to enhance infection of K562 cells. B cell clones were isolated from positive polyclonal cultures as described previously [28]. IgG concentrations in the supernatants of selected clones were determined using an IgG-specific ELISA.

Example 2. Human mAbs from immortalized B cells recognize Dengue virus proteins and neutralize infection.

For the viral neutralization and viral enhancement assay, titrated amounts of attenuated DENV of serotypes 1, 2, 3 or 4 were mixed with an equal volume of culture supernatants. Viruses and multiplicity of infection (MOI) used were: rDEN1Δ30 (03JB186-1A+V2) MOI 0.04; rDEN2/4Δ30 (04JBV351-1A-V2) MOI 0.04; rDEN3/4Δ30 (DEN3#107C) MOI 0.02; rDEN4Δ30 (06JBV591-V3+1A1+v2) MOI 0.04. After 1-hour incubation at room temperature the mixture was added to target cells (e.g. VERO cells, DC-SIGN-Raji cells or K562 cells) in 96 well flat bottom plates and incubated at 37°C for 72-96 hours. The cells were then stained with a mouse monoclonal antibody to Dengue virus 1-4 E proteins (clone 4G2), followed by a fluorescein-labeled goat anti mouse Ig and analyzed by FACS. The neutralizing titer is indicated as the concentration of antibody (μg/ml) that gives a 50% reduction of DENV infection.

For identification of the target antigens recognized by the monoclonal antibodies yeasts displaying Dengue virus E protein domains III or domain I-II were stained with the monoclonal antibodies followed by Cy5-labeled goat anti human IgG antibodies and analyzed by FACS. Western blotting experiments were performed using lysates of DENV-infected cells.

Table 3 shows that three different types of antibodies have been identified. They include those that are specific for domain III (DIII) of E protein, those that are specific for domains I-II (DI-II) of E protein and those specific for prM. The antibodies show different degrees of cross-reactivity with the 4 different DENV serotypes and neutralize those serotypes to which they bind.

Table 3. Target Antigen Specificity of Neutralizing anti-Dengue Virus Antibodies

Antibody	Target Antigen	Dengue Virus Serotypes Neutralized
HMB-DV-1	E, DIII	1,2,3
HMB-DV-2	E, DIII	1,3
HMB-DV-3	prM	1,2,3,4
HMB-DV-4	E, DI-II	1,2,3,4
HMB-DV-5	E, DI-II	1,2,3,4
HMB-DV-6	E, DIII	1,2,3
HMB-DV-7	E, DIII	1,2,3
HMB-DV-8	E	4
HMB-DV-9	E, DIII	2
HMB-DV-10	E, DIII	1,2,3,4
HMB-DV-11	E, DIII	1,2,3,4
HMB-DV-12	E, DI-DII	2
HMB-DV-13	E, DIII	1,2,3,4
HMB-DV-14	E, DIII	2

Table 4 shows the results of virus neutralization assays on VERO cells and DC-SIGN-transfected Raji cells.

Table 4. Neutralization of Dengue Virus (serotypes DENV1-DENV4) by Antibodies

Antibody	Cell type	Neutralization			
		EC ₅₀ values (µg/ml)			
		DENV1	DENV2	DENV3	DENV4
HMB-DV-1	VERO	0.013	0.577	0.014	> 20
	DC-SIGN-Raji	0.032	5.340	0.055	> 20
HMB-DV-2	VERO	0.006	> 20	0.006	> 20
	DC-SIGN-Raji	0.014	> 20	0.013	> 20
HMB-DV-3	VERO	0.912	1.615	0.120	0.070
	DC-SIGN-Raji	ND	ND	ND	ND
HMB-DV-4	VERO	0.591	0.251	0.809	0.367
	DC-SIGN-Raji	2.250	1.370	0.613	> 20
HMB-DV-5	VERO	0.066	0.034	0.118	0.200
	DC-SIGN-Raji	2.390	0.504	0.348	> 20
HMB-DV-6	VERO	0.008	0.002	0.011	> 20
	DC-SIGN-Raji	0.027	0.440	0.332	> 20
HMB-DV-7	VERO	0.016	0.004	0.020	> 20
	DC-SIGN-Raji	ND	ND	ND	ND
HMB-DV-8	VERO	> 20	> 20	> 20	0.006
	DC-SIGN-Raji	ND	ND	ND	ND

HMB-DV-9	VERO	> 20	0.002	> 20	> 20
	DC-SIGN-Raji	ND	ND	ND	ND
HMB-DV-10	VERO	> 20	0.084	> 20	0.466
	DC-SIGN-Raji	ND	ND	ND	ND
HMB-DV-11	VERO	> 20	0.048	> 20	0.520
	DC-SIGN-Raji	ND	ND	ND	ND
HMB-DV-12	VERO	ND	0.003	ND	ND
	DC-SIGN-Raji	ND	ND	ND	ND
HMB-DV-13	VERO	0.993	3.326	1.513	> 20
	DC-SIGN-Raji	ND	ND	ND	ND
HMB-DV-14	VERO	> 20	0.002	> 20	> 20
	DC-SIGN-Raji	ND	ND	ND	ND

ND: not determined

Example 3. Neutralizing recombinant anti-Dengue virus antibodies with mutations in the Fc region do not cause enhancement of virus infection on K562 cells.

Antibody-dependent enhancement (ADE) of dengue virus infection has been described in the literature. This property could limit the therapeutic effectiveness of anti-dengue virus antibodies for use in clinical situations. Therefore, mRNAs from the immortalized B cell lines expressing antibodies HMB-DV-5, HMB-DV-6 and HMB-DV-8 were isolated, cDNA was synthesized using oligo-dT specific primers, variable regions of heavy and light chain were sequenced and cloned into an expression vector using specific primers. Vectors were transfected into host cells for recombinant expression. In addition to recombinant production of the wild-type IgG1 antibodies, each of the heavy chains was mutated at amino acids 4 and 5 of CH2 domain by substituting an alanine in place of the natural leucine using site-directed mutagenesis thereby creating the LALA variant of each antibody. Both recombinant wild type and mutated antibodies were harvested from the expression cell lines and purified. Both wild-type IgG1 anti-dengue virus antibody and the LALA variant bound to the target protein in comparable manner (data not shown).

Virus neutralization and enhancement was determined as above on VERO cells and K562 cells. Each of the three antibodies has a defined molecular target as well as serotype target (see Table 3). Figure 1 shows that the unmodified recombinant antibodies neutralize target virus infection of VERO cells in a dose-dependent manner (DOTTED LINES). On K562 cells, a cell line that is not efficiently infected by Dengue viruses, the unmodified antibodies show an enhancement of viral infection at concentrations that are generally higher than those required for neutralization (SOLID LINES). The experiment was repeated using the LALA variants of each antibody. Figure 2 shows that each of the LALA variants of the recombinant anti-dengue virus antibodies also neutralized the target virus on VERO cells (DOTTED LINES) in a dose-dependent manner. However, each of the LALA antibodies did not show evidence of antibody-dependent enhancement of infection on K562

cells (SOLID LINES). Note, the dose-response is flat on the K562 cells at the concentrations of antibodies used in this experiment and the line appears very close to the X-axis.

All patents and publications referred to herein are expressly incorporated by reference in their entirety. A reference herein to a patent document or other matter which is given as prior art is not taken as an admission that that document or prior art was part of common general knowledge at the priority date of any of the claims.

With reference to the use of the word(s) "comprise" or "comprises" or "comprising" in the foregoing description and/or in the following claims, unless the context requires otherwise, those words are used on the basis and clear understanding that they are to be interpreted inclusively, rather than exclusively, and that each of those words is to be so interpreted in construing the foregoing description and/or the following claims.

It should be noted that there are alternative ways of implementing the present invention and that various modifications can be made without departing from the scope and spirit of the invention. Accordingly, the present embodiments are to be considered as illustrative and not restrictive, and the invention is not to be limited to the details given herein, but may be modified within the scope and equivalents of the appended claims.

REFERENCES (the contents of which are hereby incorporated by reference)

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CLAIMS

1. A pharmaceutical composition comprising one, two or three human antibodies, or antigen-binding fragments thereof, wherein said antibody, antibodies, or antigen-binding fragments thereof, bind to, and neutralize dengue virus, and wherein said antibody, antibodies or antigen-binding fragments thereof, comprise the heavy chain CDR1, CDR2 and CDR3 and the light chain CDR1, CDR2 and CDR3 sequences as set forth in: (i) SEQ ID NOs: 1-6, respectively; (ii) SEQ ID NOs: 17-22, respectively; (iii) SEQ ID NOs: 33-38, respectively; (iv) SEQ ID NOs: 49-54, respectively; (v) SEQ ID NOs: 67-72, respectively; (vi) SEQ ID NOs: 83-88, respectively; (vii) SEQ ID NOs: 83-85, 99, 53 and 100, respectively; (viii) SEQ ID NOs: 105-110, respectively; (ix) SEQ ID NOs: 121-123, 70, 124 and 125, respectively; (x) SEQ ID NOs: 135-137, 138, 109 and 139, respectively; (xi) SEQ ID NOs: 149, 136-138, 109 and 139, respectively; (xii) SEQ ID NOs: 153-158, respectively; (xiii) SEQ ID NOs: 169-174, respectively; or (xiv) SEQ ID NOs: 185-188, 37 and 189, respectively.
2. The pharmaceutical composition of claim 1, wherein the composition neutralizes dengue virus serotypes DENV-1, DENV-2, DENV-3, and DENV-4.
3. The pharmaceutical composition of claim 2, wherein the composition neutralizes dengue virus serotypes DENV-1, DENV-2, DENV-3, and DENV-4 by binding at least two distinct epitopes on each dengue virus serotype.
4. The pharmaceutical composition of any one of claims 1 to 3, wherein the Fc region of said antibodies, or antigen-binding fragments thereof, comprises a CH2 L4A mutation, a CH2 L5A mutation, or both.
5. The pharmaceutical composition of any one of claims 1 to 4 wherein said antibodies or said antigen-binding fragments thereof, neutralizes more than one dengue virus of two, three or four different dengue virus serotypes.

6. The pharmaceutical composition of any one of the previous claims, wherein at least one of said antibodies, or antigen-binding fragments thereof, comprise: (i) a heavy chain variable region having at least 70% sequence identity to any one of SEQ ID NOs: 13, 29, 45, 61, 65, 79, 95, 117, 131, 145, 151, 165, 181, or 195; or (ii) a light chain variable region having at least 70% sequence identity to any one of SEQ ID NOs: 14, 30, 46, 62, 80, 96, 103, 118, 132, 146, 166, 182, or 196.
7. The pharmaceutical composition of any one of the previous claims, wherein at least one of said antibodies, or antigen-binding fragments thereof, comprise: (i) a heavy chain variable region comprising the amino acid sequence of any one of SEQ ID NOs: 13, 29, 45, 61, 65, 79, 95, 117, 131, 145, 151, 165, 181, or 195; or (ii) a light chain variable region comprising the amino acid sequence of any one of SEQ ID NOs: 14, 30, 46, 62, 80, 96, 103, 118, 132, 146, 166, 182, or 196.
8. The pharmaceutical composition of any one of the preceding claims, wherein at least one of said antibodies or antigen-binding fragments thereof, comprise: a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 13 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 14; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 29 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 30; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 45 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 46; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 61 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 62; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 65 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 62; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 79 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 80; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 95 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 96; or a heavy chain variable region comprising the amino acid

sequence of SEQ ID NO: 95 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 103; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 117 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 118; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 131 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 132; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 145 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 146; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 151 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 146; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 165 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 166; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 181 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 182; or a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 195 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 196.

9. The pharmaceutical composition of any one of claims 1-8, comprising an antibody, antibodies or antigen-binding fragments thereof, comprising the heavy chain CDR1, CDR2 and CDR3 and the light chain CDR1, CDR2 and CDR3 sequences as set forth in: (v) SEQ ID NOs: 67-72, respectively; (vi) SEQ ID NOs: 83-88, respectively; or (viii) SEQ ID NOs: 105-110, respectively, or a combination thereof.
10. The pharmaceutical composition of any one of claims 1-9, wherein the pharmaceutical composition comprises a pharmaceutically acceptable diluent or carrier and, optionally, an agent useful for extending the half life of the antibody or antigen binding fragment thereof.
11. A pharmaceutical composition comprising one or more nucleic acid molecules comprising a polynucleotide encoding an antibody, antibodies, or antigen-binding

fragments thereof, binding to, and neutralizing dengue virus, wherein said antibody, antibodies or antigen-binding fragments thereof, comprise the heavy chain CDR1, CDR2 and CDR3 and the light chain CDR1, CDR2 and CDR3 sequences that is at least 95%, preferably at least 98%, more preferably at least 99% and most preferably 100% identical to the nucleic acid sequences of: (i) SEQ ID NOs: 7-12, respectively; (ii) SEQ ID NOs: 23-28, respectively; (iii) SEQ ID NOs: 39-44, respectively; (iv) SEQ ID NOs: 55-60, respectively; (v) SEQ ID NOs: 73-78, respectively; (vi) SEQ ID NOs: 89-94, respectively; (vii) SEQ ID NOs: 89-91, 101, 59, 102, respectively; (viii) SEQ ID NOs: 111-116, respectively; (ix) SEQ ID NOs: 126-128, 76, 129, 130, respectively; (x) SEQ ID NOs: 140-143, 115, 144, respectively; (xi) SEQ ID NOs: 150, 141-143, 115, 144, respectively; (xii) SEQ ID NOs: 159-164, respectively; (xiii) SEQ ID NOs: 175-180, respectively; or (xiv) SEQ ID NOs: 190-193, 43, 194, respectively.

12. The pharmaceutical composition of claim 11, wherein the polynucleotide encoding the heavy chain CDR1, CDR2 and CDR3 and the light chain CDR1, CDR2 and CDR3 has the nucleic acid sequence of: (i) SEQ ID NOs: 7-12, respectively; (ii) SEQ ID NOs: 23-28, respectively; (iii) SEQ ID NOs: 39-44, respectively; (iv) SEQ ID NOs: 55-60, respectively; (v) SEQ ID NOs: 73-78, respectively; (vi) SEQ ID NOs: 89-94, respectively; (vii) SEQ ID NOs: 89-91, 101, 59, 102, respectively; (viii) SEQ ID NOs: 111-116, respectively; (ix) SEQ ID NOs: 126-128, 76, 129, 130, respectively; (x) SEQ ID NOs: 140-143, 115, 144, respectively; (xi) SEQ ID NOs: 150, 141-143, 115, 144, respectively; (xii) SEQ ID NOs: 159-164, respectively; (xiii) SEQ ID NOs: 175-180, respectively; or (xiv) SEQ ID NOs: 190-193, 43, 194, respectively.
13. The pharmaceutical composition of claim 11 or 12, wherein the pharmaceutical composition comprises a pharmaceutically acceptable diluent or carrier.
14. A cell expressing the antibody, antibodies, or an antigen-binding fragment thereof, of the pharmaceutical composition of any one of claims 1-9, or a cell expressing a

- vector comprising one or more nucleic acid molecules of the pharmaceutical composition of claim 11 or 12.
15. A method of inhibiting or preventing, or a method of treating, a dengue virus infection or a dengue virus-related disease comprising the steps of: administering to a subject in need thereof, a therapeutically or prophylactically effective amount of the pharmaceutical composition of any one of claims 1-13.
 16. The method of claim 15, additionally comprising the administration of a second therapeutic agent.
 17. The method of claim 16, wherein said second therapeutic agent is an anti-viral agent.
 18. The pharmaceutical composition of any one of claims 1-13, for use in a method of inhibiting or preventing, or a method of treating, a dengue virus infection or a dengue virus-related disease comprising the steps of: administering to a subject in need thereof, a therapeutically or prophylactically effective amount of the pharmaceutical composition of any one of claims 1-13.
 19. Use of the pharmaceutical composition of any one of claims 1-13 (i) in the manufacture of a medicament for the treatment of dengue virus infection, (ii) in a vaccine, or (iii) in diagnosis of dengue virus infection.
 20. Use of the pharmaceutical composition of any one of claims 1-13, for monitoring the quality of anti-dengue virus vaccines by checking that the antigen of said vaccine contains the specific epitope in the correct conformation.

Figure 1. Neutralization and antibody-dependent enhancement of wild-type mAbs

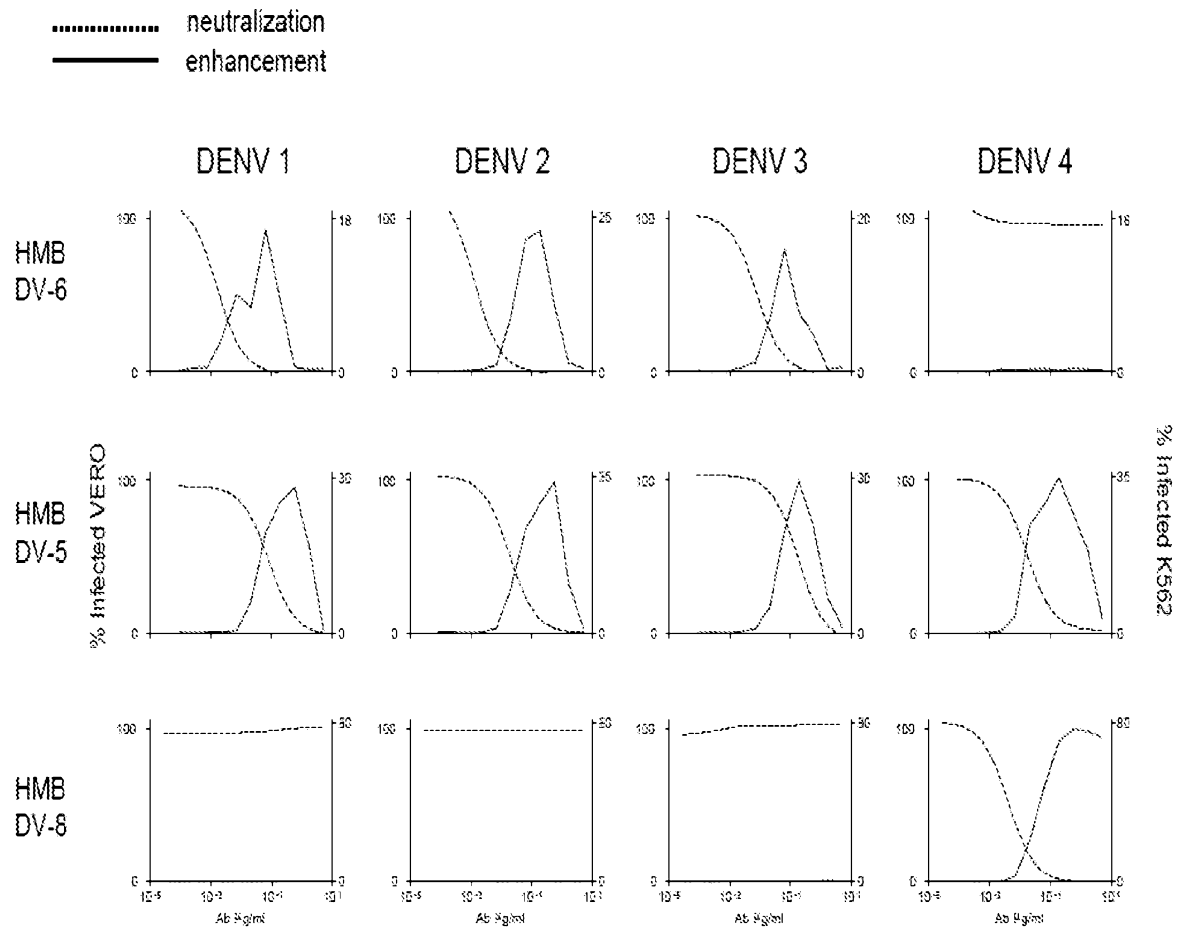
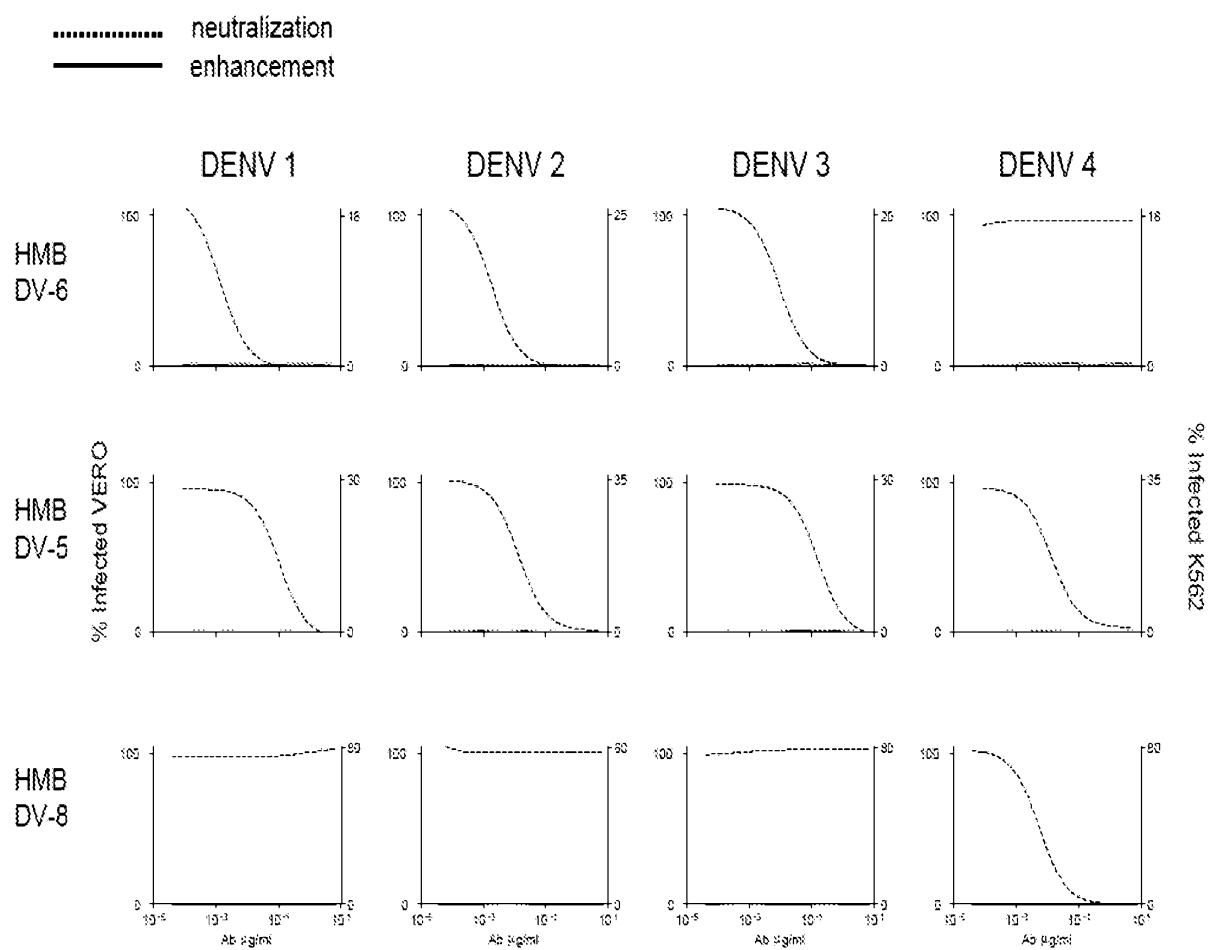


Figure 2. Neutralization and antibody-dependent enhancement of LALA-variant mAbs



2009274832 06 Jun 2011

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2009274832 06 Jun 2011

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2009274832 06 Jun 2011

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2009274832 06 Jun 2011

2009274832 06 Jun 2011

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2009274832 06 Jun 2011

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2009274832 06 Jun 2011

2009274832 06 Jun 2011

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2009274832 06 Jun 2011

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Tyr Phe Ile Pro Gly Gly Asn Arg Ala Phe Thr Pro Gly Arg Ile Asn
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Tyr Tyr Phe Lys Phe Ser Gly Pro Ser Val Ser Val Asp Thr Ala Cys
 35 40 45
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Ser Ser Ser Leu Ala Ala Ile His Val Ala Cys Asn Ser Leu Trp Arg
 50 55 60
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Pro Asp Asn

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2009274832 06 Jun 2011

2009274832 06 Jun 2011

14

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5 Ala Asn Gly Tyr Ala Arg Gly Glu Ala Val Gly Cys Leu Ile Leu Lys
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10 Pro Leu Ala Lys Ala Leu His Asp Gln Asn Lys Ile Arg Ala Val Ile
35 40 45

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30 Leu Ile Gln Ser Pro Glu Gln Gln Met Ile Ala Val Lys Ala Gly Ile
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35 Leu Ser Pro Asp Ser Met Cys His Thr Phe Asp Glu Ser Ala Asn
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55 Pro Gln Gln Arg Val Ser Leu Glu Val Ala Ser Glu Ala Leu Glu Asp
35 40 45

2009274832 06 Jun 2011

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Ala Gly Ile Pro Ala Lys Ser
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15 Lys Phe Ser Gly Pro Ser Phe Ser Ile Asp Thr Ala Cys Ser Ser Ser
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20 Leu Ala Thr Ile Gln Val
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35 Gly Met

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Thr Ala Gln Glu Ile Ser Thr Tyr Phe Ile Pro Gly
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55 <400> 36

Pro Glu Tyr Ser Gln Pro Leu Cys Thr Ala Ile Gln Ile Ala Leu Val

1 5 10 15

5 Glu Leu Leu Glu Ser Phe Gly Val Val Pro Lys Ala Val Val Gly His
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Ser Ser Gly Glu Ile Ala Ala
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15 <213> Penicillium coprobium PF1169

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20 Ile Ser Gln Pro Ala Cys Thr Ala Leu Gln Ile Ala Leu Val Asp Leu
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Leu Ala Glu Trp Ser Ile Thr Pro Ser Val Val Val Gly His Ser Ser
20 25 30

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Gly Glu Ile Ala
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1 5 10 15

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Glu Leu Leu Glu Ser Phe Gly Val Val Pro Lys Ala Val Val Gly His
20 25 30

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Ser Ser Gly Glu Ile Ala Ala
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Glu Glu Phe Trp Asp Leu Cys Ser Arg Gly Arg Gly Ala Trp Ser Pro

2009274832 06 Jun 2011

17

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5 Val Pro Lys Asp Arg Phe Asn Ala Gly Ser Phe Tyr His Pro Asn Ala
20 25 30

10 Asp Arg Pro Gly Ser Phe Asn Ala Ala Gly Ala His Phe Leu Thr Glu
35 40 45

15 Asp Ile Gly Leu Phe Asp Ala Pro Phe Phe Asn Ile Thr Leu Gln Glu
50 55 60

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20 25 30

35 Leu Phe Pro Asp Gly Ser Asn Ile Glu Thr Lys Leu Phe Val Gly Ser
35 40 45

40 Ile Lys Thr Val Ile Gly His Thr Glu Gly Ser Ala Gly Leu Ala Ser
50 55 60

45 Leu Ile Gly Ser Ser Leu Ala Met Lys His Gly Val Ile
65 70 75

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55 Lys Leu Ala Phe Val Phe Thr Gly Gln Gly Gly Gln Trp Ala Gly Met
1 5 10 15

2009274832 06 Jun 2011

18

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 20 25 30

5 Ser Gln Glu Ile Leu Ala Ser Leu Gly Cys Pro
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10 <210> 42
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15 <400> 42

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 1 5 10 15

20 Ile Phe Pro Trp Lys Ala Ala Val Val Gly Asp Thr Met Glu Gly Leu
 20 25 30

25 Ala Ala Ser Leu Ala Gln Asn Ile Lys Pro Arg Ser Val Leu Arg Met
 35 40 45

Pro Thr Leu Gly Phe Val Phe Thr Gly Gln Gly Ala Gln Trp Pro Gly
 50 55 60

30 Met Gly Lys Glu Leu Leu Gln
 65 70

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40 <400> 43

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45 Glu Glu Asp Ile Asp Ala Val Lys Ala Gln Ala Asp Gln Asp Gly Leu
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50 Phe Ala Gln Lys Leu Lys Thr Gly Val Ala Tyr His Ser Thr Ala Met
 35 40 45

55 Ser Ala Ile Ala Asn Asp Tyr
 50 55

2009274832 06 Jun 2011

19

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 10 Met Leu Ala Val Gly Ala Ser Ala Ser Asp Ile Gln Gln Ile Leu Asp
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 20 25 30
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 Ser Val Thr Leu Ser Gly Asp Leu Asp Val Ile Ala Asn Leu Gln Thr
 35 40 45
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 Ala Leu Asp Lys Glu Gly Ile Phe Thr Arg Lys Leu Lys Val Asp Val
 50 55 60

 25 Ala Tyr His Ser
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 1 5 10 15

 40 Leu Lys Gly Pro Ser Leu Ser Leu Asp Thr Ala Cys Ser Ser Ser Leu
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 Val Ala Leu His Leu Ala
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 55 Gly Pro Ser Met Thr Ile Asp Thr Ala Cys Ser Ser Ser Leu Ile Ala
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 20 25 30
 5 Val Ala Ala Gly Thr Asn Leu Leu Leu Gly Pro Glu Gln Tyr Ile Ala
 35 40 45
 10 Glu Ser Lys Leu Lys Met Leu Ser Pro Asn Gly Arg Ser Arg Met Trp
 50 55 60
 15 Asp Lys Asp Ala Asp Gly Tyr Ala Arg Gly Asp Gly Ile
 65 70 75
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 30 Val Met Gly Phe Ile Lys Ala Ile Leu Ser Ile Gln Lys Gly Val Leu
 20 25 30
 35 Ala Pro Gln Ala Asn Leu Thr Lys Leu Asn Ser Arg Ile Asp Trp Lys
 35 40 45
 Thr Ala Gly Val Lys Val Val Gln Glu Ala Thr Pro Trp
 50 55 60
 40 <210> 48
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 45 <400> 48
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 1 5 10 15
 50 Thr Met Asp Pro Gln Gln Arg Ile Phe Leu Glu Cys Val Tyr Glu Ala
 20 25 30
 55 Leu Glu Asn Gly Gly
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2009274832 06 Jun 2011

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

 20 Pro Leu Ala Lys Ala Leu His Asp Gln Asn Lys Ile Arg Ala Val Ile
 35 40 45

 25 Arg Gly Thr Gly Ser Asn Gln Asp Gly Arg Thr Ala Gly Ile Thr Val
 50 55 60

 30 Pro Asn Gly Ala Ala Gln
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 40 Thr Val Val Val Lys Pro Leu Ser Thr Ala Ile Arg Asp Gly Asp Thr
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 45 Ile Arg Ala Val Ile
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 50 <210> 51
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2009274832 06 Jun 2011

22

5 Ser Ala Ala Gly Gly Asn Thr Thr Val Ala Leu Glu Asp Ala Pro Ile
 20 25 30

10 Arg Thr Arg Ser Gly Ser Asp Pro Arg Ser Leu His Pro Ile Ala Ile
 35 40 45

15 Ser Ala Lys Ser Lys Val Ser Leu Arg Gly Asn Leu Glu Asn Leu Leu
 50 55 60

20 Ala Tyr Leu Asp Thr His Pro Asp Val Ser Leu Ser Asp Leu Ser Tyr
 65 70 75 80

Thr Thr Thr

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30 Val Tyr Ser Gly Ser Met Thr Asn Asp Tyr Glu Leu Leu Ser Thr Arg
 1 5 10 15

35 Asp Ile Tyr Asp Met Pro His Asn Ser Ala Thr Gly Asn Gly Arg Thr
 20 25 30

40 Met Leu Ala Asn Arg Leu Ser Trp Phe Phe Asp Leu Gln Gly Pro Ser
 35 40 45

45 Ile Met Met Asp Thr Ala Cys Ser Ser Ser Leu
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55 Leu Ser Pro Gln Asn Asn Pro Glu Asp Arg Cys Gln Tyr Phe Glu Ala
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Met Leu Ala Val Gly Ala Ser Ala Ser Asp Ile Gln Gln Ile Leu Asp
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2009274832 06 Jun 2011

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Ala Met Arg Gly Asn Lys Ala Val Ile Ala Cys Val Asn Ser Glu Ser
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5 Ser Val Thr Leu Ser Gly
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Ser Gly Cys Tyr Arg Glu Leu Ala Asp Cys Pro Gly Gln Arg Gly Ile
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20 Phe Thr Arg Lys Leu Lys Val Asp Val Ala Tyr His Ser
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30 <400> 57
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35 Leu Lys Gly Pro Ser Leu Ser Leu Asp Thr Ala Cys Ser Ser Ser Leu
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40 Val Ala Leu His Leu Ala
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45 <210> 58
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50 Ile Ser Glu Cys Val Thr Val Tyr Trp Lys Ala Ile Lys Ser Ala Gln
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55 Pro Asp Gly Pro Tyr Ala Leu Ala Gly Tyr Ser Tyr Gly Ser Met Leu
20 25 30

2009274832 06 Jun 2011

25

Ala Phe Glu Val Ala Lys Leu Leu Ile Lys Asn Gly Asp Lys Val Asp
 35 40 45

5 Phe Leu Gly Cys Phe Asn Leu Pro Pro His Ile
 50 55

10 <210> 59
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Gly Ala Ala Val Gln Leu Val Ile Glu Gly Gly Asn Gln Pro Lys Gly
 1 5 10 15

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

25 Asp Ala Met Lys Asp Lys His Ala Val Val Ala Cys Ile Asn Ser Asp
 35 40 45

Ser Ser Ile Thr Val Ser Gly Asp Glu Thr Ala Ile Glu Asp Leu Glu
 50 55 60

30 Ser Val Leu Lys Arg Gln Asp Ile
 65 70

35 <210> 60
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45 Thr Cys Val Gln Met Ala Leu Thr Lys Tyr Trp Thr Ser Leu Gly Val
 20 25 30

50 Thr Pro Ser Phe Val Met Gly His Ser Leu Gly Glu Phe Ala Ala Leu
 35 40 45

55 Asn Ala Ala Gly Val Leu Thr Ile Ser Asp Thr Ile Tyr Leu Ala Gly
 50 55 60

2009274832 06 Jun 2011

26

Arg Arg Ala Gln Leu Leu Thr Glu Gln Ile Lys Val Gly Thr His
 65 70 75

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

20
 Asp Arg Ile Met Glu Ser Trp Pro Thr Glu Gln Leu Trp Ala Tyr Val
 35 40 45

25
 Pro Gln Ile Pro Val Leu Ser Thr Gln Thr Gly Lys Ala Phe Gln Ala
 50 55 60

30
 Asp Ser Leu
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 Gly Pro Ser Met Thr Ile Asp Thr Ala Cys Ser Ser Ser Leu Ile Ala
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 Leu His Gln Ala Val Gln Ser Leu Arg Ser Gly Glu Thr Asp Val Ala
 20 25 30

50
 Val Ala Ala Gly Thr Asn Leu Leu Leu Gly Pro Glu Gln Tyr Ile Ala
 35 40 45

55
 Glu Ser Lys Leu Lys Met Leu Ser Pro Asn Gly Arg Ser Arg Met Trp
 50 55 60

Asp Lys Asp Ala Asp Gly Tyr Ala Arg Gly Asp Gly
 65 70 75

2009274832 06 Jun 2011

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10 Leu Phe Leu Phe Pro Asp Gly Ser Gly Ser Ala Thr Ser Tyr Ala Thr
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15 Ile Pro Gly Ile Ser Pro Asp Val Cys Val Tyr Gly Leu Asn Cys
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25 Ala Lys His Pro Pro Ala Thr Ser Ile Leu Leu Gln Gly Asn Pro Lys
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30 Thr Ala Thr Gln Ser Phe Ile Phe Val Pro
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40 Tyr Gln Ala Thr Gly Cys Ala Ala Ser Leu Gln Ser Asn Arg Ile Ser
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45 Tyr Phe Phe Asp Leu Arg Gly Pro Ser Ile Thr Ile Asp Thr Ala Cys
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50 Ser Ser Ser Leu Val Ala Leu His Tyr Ala Val Gln Ser Leu
 35 40 45

55 <210> 66
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2009274832 06 Jun 2011

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Tyr Ser Ala Thr Gly Ser Gly Leu Thr Val Leu Ala Asn Arg Ile Thr
 1 5 10 15
 5 His Cys Phe Asp Leu Arg Gly Pro Ser His Val Val Asp Thr Ala Cys
 20 25 30
 10 Ser Ser Ser Leu Tyr Ala Leu His Ser Ala Cys Leu Ala Leu Asp Ser
 35 40 45
 15 Arg Asp Cys Asp Gly Ala Val Val Ala Ala Ala Asn Leu Ile Gln Ser
 50 55 60
 Pro Glu
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 Thr Cys Val Gln Met Ala Leu Thr Lys Tyr Trp Thr Ser Leu Gly Val
 20 25 30
 35 Thr Pro Ser Phe Val Met Gly His Ser Leu Gly Glu Phe Ala Ala Leu
 35 40 45
 40 Asn Ala Ala Gly Val Leu Thr Ile Ser Asp Thr Ile Tyr Leu Ala Gly
 50 55 60
 45 Arg Arg Ala Gln Leu Leu Thr Glu Gln Ile Lys Val
 65 70 75
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 50 <211> 71
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1 5 10 15

Leu Leu Arg Asp Pro Glu Cys Val Pro Met Tyr Gln Cys Thr Asn Ala
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Leu Lys Gly Pro Ser Val Thr Val Asp Thr Ala Cys Ser Gly Ser Leu
50 55 60

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Tyr Ser Ala Thr Gly Ser Gly Leu Thr Val Leu Ala Asn Arg Ile Thr
1 5 10 15

30 His Cys Phe Asp Leu Arg Gly Pro Ser His Val Val Asp Thr Ala Cys
20 25 30

35 Ser Ser Ser Leu Tyr Ala Leu His Ser Ala Cys Phe Gly Pro Leu Asn
35 40 45

40 Ser Arg Asp Cys Asp Gly Ala Val Val Ala Ala Ala Asn Leu Ile Gln
50 55 60

45 Ser Pro Glu
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55 Ser Val Pro Ile Glu Glu His Ser Pro Val Val Thr Gln Leu Gly Thr
1 5 10 15

2009274832 06 Jun 2011

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Thr Cys Val Gln Met Ala Leu Thr Lys Tyr Trp Thr Ser Leu Gly Val
20 25 30

5 Thr Pro Ser Phe Val Met Gly His Ser Leu Gly Glu Phe Ala Ala Leu
35 40 45

10 Asn Ala Ala Gly Val Leu Thr Ile Ser Asp Thr Ile Tyr Leu Ala Gly
50 55 60

15 Arg Arg Ala Gln Leu Leu Thr Glu Gln Ile Glu Gly Gly Thr His
65 70 75

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25 Glu Ala Asn Leu His Val Pro Leu Glu Pro Thr Pro Trp Pro Ala Gly
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30 Arg Pro Glu Arg Ile Ser Val Asn Ser Phe Gly Ile Gly Gly Ser Asn
20 25 30

Ala His Ala Ile Leu Glu Ser Ala
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45 <223> Xaa can be any naturally occurring amino acid

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1 5 10 15

Ser Leu Ala Met Lys His Gly Val Ile Pro Pro Asn Leu His Phe Gly
20 25 30

55 Gln Leu Ser Glu Lys Val Ala Pro Phe Tyr Thr His Leu Asn Ile Pro

2009274832 06 Jun 2011

32

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5 Thr Glu Pro Val Pro Trp Pro Asn Ser Thr Ser Ser Gln Val Lys Arg
50 55 60

10 Ala Ser Ile Asn Ser Phe
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25 Thr Val Ala Ser Phe Arg Arg Gln Glu Asp Thr Trp Lys Val Leu Ser
20 25 30

30 Asn Ala Thr Ser Thr Leu Tyr Leu Ala Gly Ile Glu Ile
35 40 45

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<400> 76

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Ala Ala His

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55 <213> Penicillium coprobium PF1169

<400> 77

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 Asp Ile Tyr Asp Met Pro His Asn Ser Ala Thr Gly Asn Gly Arg Thr
 20 25 30
 10 Met Leu Ala Asn Arg Leu Ser Trp Phe Phe Asp Leu Gln Gly Pro Ser
 35 40 45
 15 Ile Met Met Asp Thr Ala Cys Ser Ser Ser Leu Thr Ala Val His Leu
 50 55 60
 Ala Ala Gln Ser Leu
 65
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 25 <213> Penicillium coprobium PF1169
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 30 Asp Ala Gln Phe Phe Gly Thr Lys Pro Val Glu Ala Asn Ser Ile Asp
 1 5 10 15
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 20 25 30
 35 Ser Gly Ile Pro Met Glu Arg Leu Gln Gly Ser Asn Thr Ala Val Tyr
 35 40 45
 40 Val Gly Leu Met Thr Asn Asp Tyr Ala Asp Met Leu Gly Arg Asp Met
 50 55 60
 45 Gln Asn Phe Pro Thr Tyr Phe Ala Ser Gly Thr Ala Arg Ser Ile Leu
 65 70 75 80
 50 Ser Asn Arg Val Ser
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2009274832 06 Jun 2011

2009274832 06 Jun 2011

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<400> 79

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

25

Ala Arg Lys Leu Arg Ile Asp Thr Ala Tyr His Ser Pro His Met Glu
 35 40 45

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Glu Leu Val
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Leu Lys Ser Ile Ser Pro Val Val Thr Gln Leu Gly Thr Thr Cys Val
 1 5 10 15

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Gln Met Ala Leu Thr Lys Tyr Trp
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<210> 82

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<400> 82

Gly Cys Phe Tyr Gly Met Thr Ser Asp Asp Tyr Arg Glu Val Asn Ser
 1 5 10 15

2009274832

2009274832

2009274832 06 Jun 2011

36

5 Pro Asn Arg Leu Trp Asp Met Val Ser Asn Gly Arg Ser Ala Leu Thr
 20 25 30

10 Glu Val Pro Lys Asp Arg Phe Asn Ile Asp Ala Phe Tyr His Pro His
 35 40 45

15 Ala Glu Arg Gln Gly Thr Met Asn Val Arg Arg Gly
 50 55 60

15 <210> 86
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20 <400> 86

25 Ser Val Pro Ile Glu Glu His Ser Pro Val Val Thr Gln Leu Gly Thr
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30 Thr Cys Val Gln Met Ala Leu Thr Lys Tyr Trp Thr Ser Leu Gly Val
 20 25 30

35 Thr Pro Ser Phe Val Met Gly His Ser Leu Gly Glu Phe Ala Ala Leu
 35 40 45

35 Asn Ala Ala Gly Val
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40 <210> 87
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45 <400> 87

50 Ser Val Pro Ile Glu Glu His Ser Pro Val Val Thr Gln Leu Gly Thr
 1 5 10 15

55 Thr Cys

55 <210> 88
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Phe Leu Asp Asp Leu Ala Phe Thr Val Asn Glu Arg Arg Ser Ile Phe
 1 5 10 15

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Pro Trp Lys Ala Ala Val Val Gly Asp Thr Met Glu Gly Leu Ala Ala
 20 25 30

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Ser Leu Ala Gln Asn Ile Lys Pro Arg Ser Val Leu Arg Met Pro Thr
 35 40 45

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Leu Gly Phe Val Phe Thr Gly Gln Gly Ala Gln Trp Pro Gly
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<400> 89

Ser Ser Phe Leu Thr Ser Thr Val Gln Gln Ile Val Glu Glu Thr Ile
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Gln Gly Gly Thr Gly Gln Val Val Met Glu Ser Asp Leu Met Gln Thr
 20 25 30

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Glu Phe Leu Glu Ala Ala Asn Gly His Arg Met Asn Asp Cys Gly Val
 35 40 45

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Val Thr Ser
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<400> 90

Glu Cys Gly Phe Val Glu Met His Gly Thr Gly Thr Lys Ala Gly Asp
 1 5 10 15

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

Leu Arg Asn Pro Leu Tyr Ile Gly Ser Val Lys Ser Asn Ile Gly His

2009274832 06 Jun 2011

2009274832 06 Jun 2011

38

35

40

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5 Leu Glu Gly Ala Ser Gly Ile Val Ala Val Ile Lys Ala Ala Met Met
50 55 60

10 Leu Asp Arg Asp Leu Met Leu Pro Asn Ala Glu Phe Lys
65 70 75

<210> 91
<211> 78
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15 <213> Penicillium coprobium PF1169

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<221> misc_feature
20 <222> (4)..(4)
<223> Xaa can be any naturally occurring amino acid

<400> 91
25 Phe Phe Lys Xaa Ser Gly Pro Ser Phe Ser Ile Asp Thr Ala Cys Ser
1 5 10 15

30 Ser Ser Leu Ala Thr Ile Gln Val Cys Thr His Leu Phe His Val His
20 25 30

35 Leu Asn Arg Gln Leu Thr Ile Ala Ala Cys Thr Ser Leu Trp Asn Gly
35 40 45

40 Glu Thr Asp Thr Val Val Ala Gly Gly Met Asn Ile Leu Thr Asn Ser
50 55 60

Asp Ala Phe Ala Gly Leu Ser His Gly His Phe Leu Thr Lys
65 70 75

45 <210> 92
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<213> Penicillium coprobium PF1169

50 <400> 92

Ser Val Pro Ile Glu Glu His Ser Pro Val Val Thr Gln Leu Gly Thr
1 5 10 15

55 Thr Cys Val Gln Met Ala Leu Thr Lys Tyr Trp Thr Ser Leu Gly Val
20 25 30

39

Thr Pro Ser Phe Val Met Gly His Ser Leu Gly Glu Phe Ala Ala Leu
 35 40 45

5

Asn Ala Ala Gly Val Leu Thr Ile Ser Asp Thr Ile Tyr Leu Ala Gly
 50 55 60

10

Arg Arg Ala Gln Leu Leu Thr Glu Gln Ile Glu Gly Gly Thr His
 65 70 75

15

<210> 93
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<400> 93

Leu Ser Ser Asp Gly Arg Cys His Thr Phe Asp Glu Lys Ala Asn Gly
 1 5 10 15

25

Tyr Ala Arg Gly Glu Ala Val Gly Cys Leu Ile Leu Lys Pro Leu Ala
 20 25 30

30

Lys Ala Leu His Asp Gln Asn Lys Ile Arg Ala Val Ile Arg Gly Thr
 35 40 45

35

Gly Ser Asn Gln Asp Gly Arg Thr Ala Gly Ile Thr Val Pro Asn Gly
 50 55 60

40

Ala Ala Gln Glu
 65

45

<210> 94
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 <213> Penicillium coprobium PF1169

<400> 94

50

Ser Pro Leu Phe Gly Leu Ala Arg Ile Ile Ala Ser Glu His Pro Asp
 1 5 10 15

55

Leu Gly Ser Leu Ile Asp Ile Glu Glu Pro Ile Ile Pro Leu Ser Thr
 20 25 30

Met Arg Tyr Ile Gln Gly Ala Asp Ile Val Arg Ile Ser Asp Gly Ile

2009274832 06 Jun 2011

2009274832 06 Jun 2011

40

35 40 45

5 Ala Arg Thr Ser Arg Phe Arg Ser Leu Pro Arg Thr Lys Leu Arg Pro
50 55 60

10 Val Ser Asp Gly Pro Arg Leu Leu Pro Arg Pro Glu Gly Thr Tyr Leu
65 70 75 80

<210> 95
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15 <213> Penicillium coprobium PF1169

<400> 95

20 Asn Arg Ile Ser Tyr Tyr Phe Asp Trp Gln Gly Pro Ser Met Ala Val
1 5 10 15

25 Asp Thr Gly Cys Ser Ser Ser Leu Leu Ala Val His Leu Gly Val Glu
20 25 30

30 Ala Leu Gln Asn Asp Asp Cys Ser Met Ala Val Ala Val Gly Ser Asn
35 40 45

35 Leu Ile Leu Ser Pro Asn Ala Tyr Ile Ala Asp Ser Lys Thr Arg Met
50 55 60

40 <210> 96
<211> 81
<212> PRT
<213> Penicillium coprobium PF1169

<400> 96

45 Val Asp Val Asn Pro Ala Val Leu Lys Asp Ala Pro Leu Pro Trp Asp
1 5 10 15

50 Pro Ser Ser Trp Ala Pro Ile Leu Asp Ala Ala Thr Ser Val Gly Ser
20 25 30

55 Thr Ile Phe Gln Thr Ala Ala Leu Arg Met Pro Ala Gln Ile Glu Arg
35 40 45

2009274832 06 Jun 2011

41

Val Glu Ile Phe Thr Ser Glu Asn Pro Pro Lys Thr Ser Trp Leu Tyr
50 55 60

5 Val Gln Glu Ala Ser Asp Ala Val Pro Thr Ser His Val Ser Val Val
65 70 75 80

10 Ser

15 <210> 97
<211> 37
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<213> Penicillium coprobium PF1169

<400> 97

20 Pro Leu Phe Gly Leu Ala Arg Ile Ile Ala Ser Glu His Pro Asp Leu
1 5 10 15

25 Gly Ser Leu Ile Asp Ile Glu Glu Pro Ile Ile Pro Leu Ser Thr Met
20 25 30

30 Arg Tyr Ile Arg Gly
35

35 <210> 98
<211> 84
<212> PRT
<213> Penicillium coprobium PF1169

<400> 98

40 Ala Val Ile Arg Gly Thr Gly Ser Asn Gln Asp Gly Arg Thr Ala Gly
1 5 10 15

45 Ile Thr Val Pro Asn Gly Ala Ala Gln Glu Ser Leu Ile Arg Ser Val
20 25 30

Tyr Ala Gln Ala Asp Leu Asp Pro Ser Glu Thr Asp Phe Val Glu Ala
35 40 45

50 His Gly Thr Gly Thr Leu Ala Gly Asp Pro Val Glu Thr Gly Ala Ile
50 55 60

55 Ala Arg Val Phe Gly Thr Asp Arg Pro Pro Gly Asp Pro Val Arg Ile
65 70 75 80

42

Gly Ser Ile Lys

5

<210> 99

<211> 69

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<213> Penicillium coprobium PF1169

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<400> 99

15

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

20

Arg Gly Lys Glu Ile Gly Cys Phe Val Gly Val Phe Gly Glu Asp Trp
 20 25 30

25

Leu Glu Met Ser His Lys Asp Pro Gln His Leu Asn Gln Met Phe Pro
 35 40 45

30

Asp Leu Thr Gly Pro
 65

35

<210> 100

<211> 79

<212> PRT

<213> Penicillium coprobium PF1169

<400> 100

40

Gly Gly Ala Thr Asp Thr Glu Lys Phe Trp Asp Leu Leu Ala Ser Gly
 1 5 10 15

45

Val Asp Val His Arg Lys Ile Pro Ala Asp Arg Phe Asp Val Glu Thr
 20 25 30

50

His Tyr Asp Pro Asn Gly Lys Arg Met Asn Ala Ser His Thr Pro Tyr
 35 40 45

55

Gly Cys Phe Ile Asp Glu Pro Gly Leu Phe Asp Ala Ala Phe Phe Asn
 50 55 60

Met Ser Pro Arg Glu Ala Gln Gln Thr Asp Pro Met Gln Arg Leu
 65 70 75

2009274832 06 Jun 2011

2009274832 06 Jun 2011

43

5 <210> 101
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 <213> Penicillium coprobium PF1169

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 10 Glu Leu Arg His Gly Lys Asn Ile Asp Lys Pro Glu Tyr Ser Gln Pro
 1 5 10 15

 15 Leu Cys Thr Ala Ile Gln Ile Ala Leu Val Glu Leu Leu Glu Ser Phe
 20 25 30

 Gly Val Val Pro Lys Ala Val Val Gly His Ser Ser Gly Glu Ile Ala
 35 40 45
 20
 Ala Ala Tyr Val
 50
 25
 <210> 102
 <211> 34
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 <213> Penicillium coprobium PF1169
 30
 <400> 102

 Val Gly Phe Val Phe Thr Gly Gln Gly Ala Gln Trp His Gly Met Gly
 1 5 10 15
 35
 Lys Glu Leu Leu Ser Thr Tyr Pro Ile Phe Arg Gln Thr Met Gln Asp
 20 25 30
 40
 Val Asp

 45 <210> 103
 <211> 63
 <212> PRT
 <213> Penicillium coprobium PF1169
 50
 <400> 103

 Phe Asp Ala Ala Phe Phe Asn Met Ser Pro Arg Glu Ala Gln Gln Thr
 1 5 10 15
 55
 Asp Pro Met Gln Arg Leu Ala Ile Val Thr Ala Tyr Glu Ala Leu Glu
 20 25 30

44

Arg Ala Gly Tyr Val Ala Asn Arg Thr Ala Ala Thr Asn Leu His Arg
 35 40 45

5

Ile Gly Thr Phe Tyr Gly Gln Ala Ser Asp Asp Tyr Arg Glu Val
 50 55 60

10

<210> 104
 <211> 43
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 <213> Penicillium coprobium PF1169

15

<400> 104

Ala Val Val Ser Gly Val Ser Ile Leu Glu Asn Pro Val Glu Thr Ile
 1 5 10 15

20

Gly Met Ser His His Gly Leu Leu Gly Pro Gln Gly Arg Ser Phe Ser
 20 25 30

25

Phe Asp Ser Arg Ala Glu Gly Tyr Ala Arg Gly
 35 40

30

<210> 105
 <211> 71
 <212> PRT
 <213> Penicillium coprobium PF1169

35

<400> 105

Lys Ala Ser Leu Ser Leu Gln His Gly Met Ile Ala Pro Asn Leu Leu
 1 5 10 15

40

Met Gln His Leu Asn Pro Lys Ile Lys Pro Phe Ala Ala Lys Leu Ser
 20 25 30

45

Val Pro Thr Glu Cys Ile Pro Trp Pro Ala Val Pro Asp Gly Cys Pro
 35 40 45

50

Arg Arg Ala Ser Val Asn Ser Phe Gly Phe Gly Gly Ala Asn Val His
 50 55 60

55

Val Val Leu Glu Ser Tyr Thr
 65 70

<210> 106

2009274832 06 Jun 2011

2009274832 06 Jun 2011

45

<211> 28
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 <213> Penicillium coprobium PF1169

5 <400> 106

Pro Trp Pro Thr Thr Gly Leu Arg Arg Ala Ser Val Asn Ser Phe Gly
 1 5 10 15

10

Tyr Gly Gly Thr Asn Ala His Cys Val Leu Asp Asp
 20 25

15

<210> 107
 <211> 71
 <212> PRT
 <213> Penicillium coprobium PF1169

20 <400> 107

Lys Ala Ser Leu Ser Leu Gln His Gly Met Ile Ala Pro Asn Leu Leu
 1 5 10 15

25

Met Gln His Leu Asn Pro Lys Ile Lys Pro Phe Ala Ala Lys Leu Ser
 20 25 30

30

Val Pro Thr Glu Cys Ile Pro Trp Pro Ala Val Pro Asp Gly Cys Pro
 35 40 45

35

Arg Arg Ala Ser Val Asn Ser Phe Gly Phe Gly Gly Ala Asn Val His
 50 55 60

40

Val Val Leu Glu Ser Tyr Thr
 65 70

<210> 108

<211> 50

<212> PRT

45 <213> Penicillium coprobium PF1169

<400> 108

Asp Arg Leu Phe Leu Gln Met Ser His Glu Glu Trp Glu Ala Ala Leu
 1 5 10 15

Ala Pro Lys Val Thr Gly Thr Trp Asn Leu His His Ala Thr Ala Gln
 20 25 30

55

His Ser Leu Asp Phe Phe Val Val Phe Gly Ser Ile Ala Gly Val Cys

2009274832 06 Jun 2011

46

35 40 45
 5 Gly Asn
 50
 10 <210> 109
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 <213> Penicillium coprobium PF1169
 <400> 109
 15 Thr Phe Leu Lys Gly Thr Gly Gly Gln Met Leu Gln Asn Val Val Leu
 1 5 10 15
 20 Arg Val Pro Val Ala Ile Asn Ala Pro Arg Ser Val Gln Val Val Val
 20 25 30
 25 Gln Gln Asp Gln Val Lys Val Val Ser Arg Leu Ile Pro Ser Glu Ala
 35 40 45
 30 Ser Val Leu Asp Asp Asp Ala Ser Trp Val Thr His Thr Thr Ala Tyr
 50 55 60
 35 Trp Asp Arg Arg Val Leu Gly Ser Glu Asp Arg Ile Asp Leu Ala Ala
 65 70 75 80
 40 Val Lys
 45 <210> 110
 <211> 30
 <212> PRT
 <213> Penicillium coprobium PF1169
 <400> 110
 50 Asp Thr Ala Cys Ser Ser Ser Leu Val Ala Leu His Tyr Ala Val Gln
 1 5 10 15
 55 Ser Leu Arg Asn Gly Glu Ser Thr Glu Ala Leu Ile Ala Gly
 20 25 30
 60 <210> 111
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 <212> PRT
 <213> Penicillium coprobium PF1169

47

<400> 111

5 Gly Thr Gly Asn Gly Ser Ala Met Ile Ser Asn Arg Ile Ser Trp Phe
1 5 10 15

10 Phe Asp Leu Lys Gly Pro Ser Leu Ser Leu Asp Thr Ala Cys Ser Ser
20 25 30

15 Ser Leu Val Ala Leu His
35

<210> 112

<211> 72

<212> PRT

20 <213> Penicillium coprobium PF1169

<400> 112

25 Thr Ser Thr Gln Leu Asn Asp Leu Asn Glu Thr Asn Ala Ile Lys Lys
1 5 10 15

30 Val Phe Gly Lys Gln Ala Tyr Asn Ile Pro Ile Ser Ser Thr Lys Ser
20 25 30

35 Tyr Thr Gly His Leu Ile Gly Ala Ala Gly Thr Met Glu Thr Ile Phe
35 40 45

35 Cys Ile Lys Thr Met Gln Glu Lys Ile Ala Pro Ala Thr Thr Asn Leu
50 55 60

40 Lys Glu Arg Asp Ser Asn Cys Asp
65 70

<210> 113

<211> 50

45 <212> PRT

<213> Penicillium coprobium PF1169

<400> 113

50 Val Ile Val Gly Ser Ala Ala Asn Gln Asn Leu Asn Leu Ser His Ile
1 5 10 15

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

2009274832 06 Jun 2011

48

Ser Gln Ala Gly Val His Pro His Ser Val Thr Tyr Val Glu Ala His
 35 40 45

5 Gly Thr
 50

10 <210> 114
 <211> 48
 <212> PRT
 <213> Penicillium coprobium PF1169

15 <400> 114

Leu Pro Thr Ala Ile Gln Pro Leu Phe Arg Ala Asn Val Ser Tyr Leu
 1 5 10 15

20 Leu Val Gly Gly Leu Gly Gly Ile Gly Lys Glu Val Ala Leu Trp Met
 20 25 30

25 Val Gln Asn Gly Ala Lys Ser Leu Ile Phe Val Asn Arg Ser Gly Leu
 35 40 45

30 <210> 115
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 <213> Penicillium coprobium PF1169

<400> 115

35 Val Ala Ile Val Gly Gly Val Asn Ala Leu Cys Gly Pro Gly Leu Thr
 1 5 10 15

40 Arg Val Leu Asp Lys Ala Gly Ala Ile Ser Ser Asp Gly Ser Cys Lys
 20 25 30

45 Ser Phe Asp Asp Asp Ala His Gly Tyr Ala Arg Gly Glu Gly Ala Gly
 35 40 45

Ala Leu Val Leu Lys
 50

50 <210> 116
 <211> 28
 <212> PRT
 <213> Penicillium coprobium PF1169

55 <400> 116

2009274832 06 Jun 2011

2009274832 06 Jun 2011

49

Pro Trp Glu Ser Pro Gly Ala Arg Arg Val Ser Val Asn Ser Phe Gly
1 5 10 15

5 Tyr Gly Gly Ser Asn Ala His Val Ile Ile Glu Asp
20 25

10 <210> 117
<211> 72
<212> PRT
<213> Penicillium coprobium PF1169

15 <400> 117

Lys Thr Leu Arg Glu Trp Met Thr Ala Glu Gly Lys Asp His Asn Leu
1 5 10 15

20 Ser Asp Ile Leu Thr Thr Leu Ala Thr Arg Arg Asp His His Asp Tyr
20 25 30

25 Arg Ala Ala Leu Val Val Asp Asp Asn Arg Asp Ala Glu Leu Ala Leu
35 40 45

Gln Ala Leu Glu His Gly Val Asp Gln Thr Phe Thr Thr Gln Ser Arg
50 55 60

30 Val Phe Gly Ala Asp Ile Ser Lys
65 70

35 <210> 118
<211> 80
<212> PRT
<213> Penicillium coprobium PF1169

40 <400> 118

Ser Asp Asp Tyr Arg Glu Val Asn Ser Gly Gln Asp Ile Asp Thr Tyr
1 5 10 15

45 Phe Ile Pro Gly Gly Asn Arg Ala Phe Thr Pro Gly Arg Ile Asn Tyr
20 25 30

50 Tyr Phe Lys Phe Ser Gly Pro Ser Val Ser Val Asp Thr Ala Cys Ser
35 40 45

55 Ser Ser Leu Ala Ala Ile His Val Ala Cys Asn Ser Leu Trp Arg Asn
50 55 60

2009274832 06 Jun 2011

50

Glu Ser Asp Ser Ala Val Ala Gly Gly Val Asn Ile Leu Thr Asn Pro
 65 70 75 80

5
 <210> 119
 <211> 56
 <212> PRT
 <213> Penicillium coprobium PF1169

10
 <400> 119

Leu Ser Ser Asp Gly Arg Cys His Thr Phe Asp Glu Lys Ala Asn Gly
 1 5 10 15

15
 Tyr Ala Arg Gly Glu Ala Val Gly Cys Leu Ile Leu Lys Pro Leu Ala
 20 25 30

20
 Lys Ala Leu His Asp Gln Asn Lys Ile Arg Ala Val Ile Arg Gly Thr
 35 40 45

25
 Gly Ser Asn Gln Gly Arg Ala Asn
 50 55

30
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 <213> Penicillium coprobium PF1169

35
 <400> 120

Asp Thr Ala Cys Ser Ser Ser Leu Tyr Ala Leu His Ser Ala Cys Leu
 1 5 10 15

40
 Ala Leu Asp Ser Arg Asp Cys Asp Gly Ala Val Val Ala Ala Ala Asn
 20 25 30

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

50
 Leu Ser Pro Asp Ser Met Cys His Thr Phe Asp Glu Ser Ala Asn
 50 55 60

55
 <210> 121
 <211> 28
 <212> PRT
 <213> Penicillium coprobium PF1169

<400> 121

51

Pro Trp Pro Thr Thr Gly Leu Arg Arg Ala Ser Val Asn Ser Phe Gly
1 5 10 15

5

Tyr Gly Gly Thr Asn Ala His Cys Val Leu Asp Asp
20 25

10

<210> 122
<211> 62
<212> PRT
<213> Penicillium coprobium PF1169

15

<400> 122

Ala Gly Ile Pro Leu Ala Asn Ile Met Gly Thr Lys Thr Ser Cys Phe
1 5 10 15

20

Val Gly Ser Phe Ser Ala Asp Tyr Thr Asp Leu Leu Leu Arg Asp Pro
20 25 30

25

Glu Cys Val Pro Met Tyr Gln Cys Thr Asn Ala Gly Gln Ser Arg Ala
35 40 45

30

Met Thr Ala Asn Arg Leu Ser Tyr Phe Leu Ile Lys Gly Pro
50 55 60

35

<210> 123
<211> 80
<212> PRT
<213> Penicillium coprobium PF1169

<400> 123

40

Arg Trp Glu Pro Tyr Tyr Arg Arg Asp Pro Arg Asn Glu Lys Phe Leu
1 5 10 15

45

Lys Gln Thr Thr Ser Arg Gly Tyr Phe Leu Asp His Leu Glu Asp Phe
20 25 30

50

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

Pro Gln Gln Arg Val Ser Leu Glu Val Ala Ser Glu Ala Leu Glu Asp
50 55 60

55

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

2009274832 06 Jun 2011

2009274832 06 Jun 2011

52

5 <210> 124
 <211> 28
 <212> PRT
 <213> Penicillium coprobium PF1169

<400> 124

10 Pro Gly Arg Ile Asn Tyr Phe Phe Lys Phe Ser Gly Pro Ser Phe Ser
 1 5 10 15

15 Ile Asp Thr Ala Cys Ser Ser Ser Leu Ala Thr Ile
 20 25

20 <210> 125
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 <212> PRT
 <213> Penicillium coprobium PF1169

<400> 125

25 Ala Gly Ile Pro Leu Ala Asn Ile Met Gly Thr Lys Thr Ser Cys Phe
 1 5 10 15

30 Val Gly Ser Phe Ser Ala Asp Tyr Thr Asp Leu Leu Leu Arg Asp Pro
 20 25 30

35 Glu Cys Val Pro Met Tyr Gln Cys Thr Asn Ala Gly Gln Ser Arg Ala
 35 40 45

Met Thr Ala Asn Arg Leu Ser Tyr Phe Phe Asp Leu Lys Gly Pro Ser
 50 55 60

40

<210> 126
 <211> 52
 <212> PRT
 <213> Penicillium coprobium PF1169

45

<400> 126

50 Glu Leu Arg His Gly Lys Asn Ile Asp Lys Pro Glu Tyr Ser Gln Pro
 1 5 10 15

50

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

55

Gly Val Val Pro Lys Ala Val Val Gly His Ser Ser Gly Glu Ile Ala
 35 40 45

2009274832 06 Jun 2011

54

<211> 69
 <212> PRT
 <213> Penicillium coprobium PF1169

5 <400> 129

Gln Phe Phe His Ala His Gly Thr Gly Thr Gln Ala Gly Asp Pro Gln
 1 5 10 15

10 Glu Ala Glu Ala Val Ser Thr Ala Leu Phe Pro Asp Gly Ser Asn Ile
 20 25 30

15 Glu Thr Lys Leu Phe Val Gly Ser Ile Lys Thr Val Ile Gly His Thr
 35 40 45

20 Glu Gly Ser Ala Gly Leu Ala Ser Leu Ile Gly Ser Ser Leu Ala Met
 50 55 60

Lys His Gly Val Ile
 65
 25

<210> 130
 <211> 64
 <212> PRT
 30 <213> Penicillium coprobium PF1169

<400> 130

35 Ala Gly Ile Pro Leu Ala Asn Ile Met Gly Thr Lys Thr Ser Cys Phe
 1 5 10 15

40 Val Gly Ser Phe Ser Ala Asp Tyr Thr Asp Leu Leu Leu Arg Asp Pro
 20 25 30

Glu Cys Val Pro Met Tyr Gln Cys Thr Asn Ala Gly Gln Ser Arg Ala
 35 40 45

45 Met Thr Ala Asn Arg Leu Ser Tyr Phe Phe Asp Leu Lys Gly Pro Ser
 50 55 60

50 <210> 131
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 <212> PRT
 <213> Penicillium coprobium PF1169

55 <400> 131

Leu Asp Asp Leu Ala Phe Thr Val Asn Glu Arg Arg Ser Ile Phe Pro

2009274832 06 Jun 2011

55

1 5 10 15

5 Trp Lys Ala Ala Val Val Gly Asp Thr Met Glu Gly Leu Ala Ala Ser
20 25 30

10 Leu Ala Gln Asn Ile Lys Pro Arg Ser Val Leu Arg Met Pro Thr Leu
35 40 45

15 Gly Phe Val Phe Thr Gly Gln Gly Ala Gln Trp Pro Gly Met Gly Lys
50 55 60

65 Glu Leu Leu

20 <210> 132
<211> 21
<212> PRT
<213> Penicillium coprobium PF1169

25 <400> 132

Ala His Gly Thr Gly Thr Lys Val Gly Asp Pro Met Glu Val Glu Ala
1 5 10 15

30 Ile Ala Asp Val Phe
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35 <210> 133
<211> 71
<212> PRT
<213> Penicillium coprobium PF1169

40 <400> 133

Lys Gly Gly Met Leu Ala Val Gly Ala Ser Ala Ser Asp Ile Gln Gln
1 5 10 15

45 Ile Leu Asp Ala Met Arg Gly Asn Lys Ala Val Ile Ala Cys Val Asn
20 25 30

50 Ser Glu Ser Ser Val Thr Leu Ser Gly Asp Leu Asp Val Ile Ala Asn
35 40 45

55 Leu Gln Thr Ala Leu Asp Lys Glu Gly Ile Phe Thr Arg Lys Leu Lys
50 55 60

2009274832 06 Jun 2011

56

Val Asp Val Ala Tyr His Ser
65 70

5 <210> 134
<211> 75
<212> PRT
<213> Penicillium coprobium PF1169

10 <400> 134

Leu Glu Asn Leu Glu Thr Ala Leu Ala Arg Asn Ala Pro Ile Tyr Ala
1 5 10 15

15 Glu Val Thr Gly Tyr Ala Asn Tyr Ser Asp Ala Tyr Asp Ile Thr Ala
20 25 30

20 Pro Ala Asp Asp Leu Met Gly Arg Tyr Met Ser Ile Thr Lys Ala Ile
35 40 45

25 Glu Gln Ala Gln Leu Asn Ile Asn Glu Ile Asp Tyr Ile Asn Ala His
50 55 60

30 Gly Thr Ser Thr Gln Leu Asn Asp Leu Asn Glu
65 70 75

<210> 135
<211> 53
<212> PRT
35 <213> Penicillium coprobium PF1169

<400> 135

40 Met Ala Met Lys Lys Ala Leu Lys Gln Ala Gln Leu Arg Pro Ser Ala
1 5 10 15

45 Val Asp Tyr Val Asn Ala His Ala Thr Ser Thr Ile Val Gly Asp Ala
20 25 30

Ala Glu Asn Ala Ala Ile Lys Ala Leu Leu Leu Gly Ala Asp Gly Lys
35 40 45

50 Asp Lys Ala Ala Asp
50

55 <210> 136
<211> 38
<212> PRT

2009274832 06 Jun 2011

57

<213> Penicillium coprobium PF1169

<400> 136

5 Gly Thr Gly Asn Gly Ser Ala Met Ile Ser Asn Arg Ile Ser Trp Phe
1 5 10 15

10 Phe Asp Leu Lys Gly Pro Ser Leu Ser Leu Asp Thr Ala Cys Ser Ser
20 25 30

15 Ser Leu Val Ala Leu His
35

<210> 137

<211> 76

<212> PRT

20 <213> Penicillium coprobium PF1169

<400> 137

25 Gly Pro Ser Met Thr Ile Asp Thr Ala Cys Ser Ser Ser Leu Ile Ala
1 5 10 15

30 Leu His Gln Ala Val Gln Ser Leu Arg Ser Gly Glu Thr Asp Val Ala
20 25 30

35 Val Ala Ala Gly Thr Asn Leu Leu Leu Gly Pro Glu Gln Tyr Ile Ala
35 40 45

40 Glu Ser Lys Leu Lys Met Leu Ser Pro Asn Gly Arg Ser Arg Met Trp
50 55 60

40 Asp Lys Asp Ala Asp Gly Tyr Ala Arg Gly Asp Gly
65 70 75

<210> 138

45 <211> 85

<212> PRT

<213> Penicillium coprobium PF1169

<400> 138

50 Ile Gly Ser Ile Lys Pro Asn Ile Gly His Leu Glu Ala Gly Ala Gly
1 5 10 15

55 Val Met Gly Phe Ile Lys Ala Ile Leu Ser Ile Gln Lys Gly Val Leu
20 25 30

2009274832 06 Jun 2011

58

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

5 Thr Ala Gly Val Lys Val Val Gln Glu Ala Thr Pro Trp Pro Ser Ser
50 55 60

10 Asp Ser Ile Arg Arg Ala Gly Val Cys Ser Tyr Gly Tyr Gly Gly Thr
65 70 75 80

15 Val Ser His Ala Val
85

20 <210> 139
<211> 57
<212> PRT
<213> Penicillium coprobium PF1169

<400> 139

25 Asn Ala Ala Gly Ala His Phe Leu Thr Glu Asp Ile Gly Leu Phe Asp
1 5 10 15

30 Ala Pro Phe Phe Asn Ile Thr Leu Gln Glu Ala Gln Thr Met Asp Pro
20 25 30

35 Gln Gln Arg Ile Phe Leu Glu Cys Val Tyr Glu Ala Leu Glu Asn Gly
35 40 45

Gly Ile Pro Thr His Glu Ile Thr Gly
50 55

40 <210> 140
<211> 68
<212> PRT
<213> Penicillium coprobium PF1169

45 <400> 140

50 Leu Ser Ser Asp Gly Arg Cys His Thr Phe Asp Glu Lys Ala Asn Gly
1 5 10 15

Tyr Ala Arg Gly Glu Ala Val Gly Cys Leu Ile Leu Lys Pro Leu Ala
20 25 30

55 Lys Ala Leu His Asp Gln Asn Lys Ile Arg Ala Val Ile Arg Gly Thr
35 40 45

59

Gly Ser Asn Gln Asp Gly Arg Thr Ala Gly Ile Thr Val Pro Asn Gly
50 55 60

5

Ala Ala Gln Glu
65

10

<210> 141
<211> 37
<212> PRT
<213> Penicillium coprobium PF1169

15

<400> 141

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

20

Thr Val Val Val Lys Pro Leu Ser Thr Ala Ile Arg Asp Gly Asp Thr
20 25 30

25

Ile Arg Ala Val Ile
35

30

<210> 142
<211> 72
<212> PRT
<213> Penicillium coprobium PF1169

35

<400> 142

Gly Ile Pro Ile Asp Thr Leu Pro Gly Ser Asn Thr Ala Val Tyr Ser
1 5 10 15

40

Gly Ser Met Thr Asn Asp Tyr Glu Leu Leu Ser Thr Arg Asp Ile Tyr
20 25 30

45

Asp Met Pro His Asn Ser Ala Thr Gly Asn Gly Arg Thr Met Leu Ala
35 40 45

50

Asn Arg Leu Ser Trp Phe Phe Asp Leu Gln Gly Pro Ser Ile Met Met
50 55 60

55

Asp Thr Ala Cys Ser Ser Ser Leu
65 70

<210> 143

2009274832 06 Jun 2011

60

<211> 83
 <212> PRT
 <213> Penicillium coprobium PF1169

5 <400> 143

Ala Gln Gln Ser Leu Ile Leu Ala Thr Tyr Ala Arg Ala Gly Leu Ser
 1 5 10 15

10

Pro Gln Asn Asn Pro Glu Asp Arg Cys Gln Tyr Phe Glu Ala His Gly
 20 25 30

15

Thr Gly Thr Gln Ala Gly Asp Pro Gln Glu Ala Ala Ala Ile Asn Ser
 35 40 45

20

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

25

Val Gly Ser Ile Lys Thr Ile Ile Gly His Thr Glu Ala Thr Ala Gly
 65 70 75 80

Leu Ala Gly

30

<210> 144
 <211> 69
 <212> PRT
 <213> Penicillium coprobium PF1169

35

<400> 144

Pro Leu Trp Arg Lys Ile Glu Thr Ala Pro Leu Asn Thr Gly Leu Thr
 1 5 10 15

40

His Asp Val Glu Lys His Thr Leu Leu Gly Gln Arg Ile Pro Val Ala
 20 25 30

45

Gly Thr Asp Thr Phe Val Tyr Thr Thr Arg Leu Asp Asn Glu Thr Lys
 35 40 45

50

Pro Phe Pro Gly Ser His Pro Leu His Gly Thr Glu Ile Val Pro Ala
 50 55 60

55

Ala Gly Leu Ile Asn
 65

2009274832 06 Jun 2011

2009274832 06 Jun 2011

61

<210> 145
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<400> 145

Ala Gly Ile Pro Leu Ala Asn Ile Met Gly Thr Lys Thr Ser Cys Phe
 1 5 10 15

10

Val Gly Ser Phe Ser Ala Asp Tyr Thr Asp Leu Leu Leu Arg Asp Pro
 20 25 30

15

Glu Cys Val Pro Met Tyr Gln Cys Thr Asn Ala Gly Gln Ser Arg Ala
 35 40 45

20

Met Thr Ala Asn Arg Leu Ser Tyr Phe Phe Asp Leu Lys Gly Pro Ser
 50 55 60

25

<210> 146
 <211> 81
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30

<400> 146

Gly Tyr Gly Arg Gly Glu Gly Val Ala Ser Val Val Leu Lys Arg Leu
 1 5 10 15

35

Gln Asp Ala Ile Asn Asp Gly Asp Pro Ile Glu Cys Val Ile Arg Ala
 20 25 30

40

Ser Gly Ala Asn Ser Asp Gly Arg Thr Met Gly Ile Thr Met Pro Asn
 35 40 45

45

Pro Lys Ala Gln Gln Ser Leu Ile Leu Ala Thr Tyr Ala Arg Ala Gly
 50 55 60

Leu Ser Pro Gln Asn Asn Pro Glu Asp Arg Cys Gln Tyr Phe Glu Ala
 65 70 75 80

50

His

55

<210> 147
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2009274832 06 Jun 2011

62

<213> Penicillium coprobium PF1169

<400> 147

5 Gly Thr Gly Asn Gly Ser Ala Met Ile Ser Asn Arg Ile Ser Trp Phe
1 5 10 15

10 Phe Asp Leu Lys Gly Pro Ser Leu Ser Leu Asp Thr Ala Cys Ser Ser
20 25 30

15 Ser Leu Val Ala Leu His
35

<210> 148

<211> 53

<212> PRT

20 <213> Penicillium coprobium PF1169

<400> 148

25 Glu Ala Thr Ser Met Asp Ala Gln Gln Arg Lys Leu Leu Glu Val Thr
1 5 10 15

30 Tyr Glu Ala Leu Glu Asn Ala Gly Val Pro Leu Glu Thr Ile Gln Gly
20 25 30

Ser Asn Thr Gly Val Tyr Val Gly Asn Phe Thr Asn Asp Phe Leu Asn
35 40 45

35 Met Gln Tyr Lys Asp
50

40 <210> 149

<211> 82

<212> PRT

<213> Penicillium coprobium PF1169

45 <400> 149

Gly Ser Leu Ile Asp Ile Glu Glu Pro Ile Ile Pro Leu Ser Thr Met
1 5 10 15

50 Arg Tyr Ile Gln Gly Ala Asp Ile Val Arg Ile Ser Asp Gly Ile Ala
20 25 30

55 Arg Thr Ser Arg Phe Arg Ser Leu Pro Arg Thr Lys Leu Arg Pro Val
35 40 45

2009274832 06 Jun 2011

63

Ser Asp Gly Pro Arg Leu Leu Pro Arg Pro Glu Gly Thr Tyr Leu Ile
 50 55 60

5 Thr Gly Gly Leu Gly Ile Leu Gly Leu Glu Val Ala Asp Phe Leu Val
 65 70 75 80

10 Glu Lys

15 <210> 150
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20 <400> 150
 Gln Leu Gly Thr Thr Cys Val Gln Met Ala Leu Thr Lys Tyr Trp Thr
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25 Ser Leu Gly Val Thr Pro Ser Phe Val Met Gly His Ser Leu Gly Glu
 20 25 30

30 Phe Ala Ala Leu Asn Ala Ala Gly Val Leu Thr Ile Ser Asp Thr Ile
 35 40 45

35 Tyr Leu Ala Gly Arg Arg Ala Gln Leu Leu Thr Glu Gln Ile Lys Val
 50 55 60

Gly
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40 <210> 151
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 <222> (45)..(45)
 50 <223> Xaa can be any naturally occurring amino acid

<400> 151

55 Gly Pro Arg Leu Leu Pro Arg Pro Glu Gly Thr Tyr Leu Ile Thr Gly
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2009274832

65

Ser Leu Arg Asn Gly Glu
35

5

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<213> Penicillium coprobium PF1169

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<400> 154

15

Gly Ser Gly Leu Thr Val Leu Ala Asn Arg Ile Thr His Cys Phe Asp
1 5 10 15

Leu Arg Gly Pro Ser His Val Val Asp Thr Ala Cys Ser Ser Ser Leu
20 25 30

20

Tyr Ala Leu His Ser Ala Cys Leu Ala Leu Asp Ser Arg Asp Cys Asp
35 40 45

25

Gly Ala Val Val Ala Ala Ala Asn Leu Ile Gln Ser Pro Glu Gln Gln
50 55 60

30

Met Ile Ala Val Lys Ala Gly Ile Leu Ser
65 70

35

<210> 155
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<213> Penicillium coprobium PF1169

<400> 155

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Gln Leu Gly Thr Thr Cys Val Gln Met Ala Leu Thr Lys Tyr Trp Thr
1 5 10 15

45

Ser Leu Gly Val Thr Pro Ser Phe Val Met Gly His Ser Leu Gly Glu
20 25 30

50

Phe Ala Ala Leu Asn Ala Ala Gly Val Leu Thr Ile Ser Asp Thr Ile
35 40 45

Tyr Leu Ala Gly Arg Arg Ala Gln Leu
50 55

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<210> 156
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2009274832 06 Jun 2011

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<212> PRT

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<400> 156

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His Leu Asn Leu Met Gly Pro Ser Thr Ala Val Asp Ala Ala Cys Ala
1 5 10 15

10

Ser Ser Leu Val Ala Ile His His Gly Val Gln Ala Ile Lys Leu Gly
20 25 30

15

Glu Ser Arg Val Ala Ile Val Gly Gly Val Asn Ala Leu Cys Gly Pro
35 40 45

20

Gly Leu Thr Arg Val Leu Asp Lys Ala Gly Ser Ile Ser Ser Asp Gly
50 55 60

Ser Cys Lys Ser Phe Asp Asp Asp
65 70

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<210> 157

<211> 81

<212> PRT

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<400> 157

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Leu Lys Gly Thr Gly Gly Gln Met Leu Gln Asn Val Val Leu Arg Val
1 5 10 15

Pro Val Ala Ile Asn Ala Pro Arg Ser Val Gln Val Val Val Gln Gln
20 25 30

40

Asp Gln Val Lys Val Val Ser Arg Leu Ile Pro Ser Glu Ala Ser Val
35 40 45

45

Leu Asp Asp Asp Ala Ser Trp Val Thr His Thr Thr Ala Tyr Trp Asp
50 55 60

50

Arg Arg Val Leu Gly Ser Glu Asp Arg Ile Asp Leu Ala Ala Val Lys
65 70 75 80

Ser

55

<210> 158

2009274832 06 Jun 2011

2009274832 06 Jun 2011

67

<211> 82
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 <213> Penicillium coprobium PF1169

5 <400> 158

Ile Met Gly Thr Lys Thr Ser Cys Phe Val Gly Ser Phe Ser Ala Asp
 1 5 10 15

10 Tyr Thr Asp Leu Leu Leu Arg Asp Pro Glu Cys Val Pro Met Tyr Gln
 20 25 30

15 Cys Thr Asn Ala Gly Gln Ser Arg Ala Met Thr Ala Asn Arg Leu Ser
 35 40 45

20 Tyr Phe Phe Asp Leu Lys Gly Pro Ser Val Thr Val Asp Thr Ala Cys
 50 55 60

25 Ser Gly Ser Leu Val Ala Leu His Leu Ala Cys Gln Ser Leu Arg Thr
 65 70 75 80

Gly Asp

30

<210> 159
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 <213> Penicillium coprobium PF1169

35

<400> 159

Gly Ser Gly Leu Thr Val Leu Ala Asn Arg Ile Thr His Cys Phe Asp
 1 5 10 15

40

Leu Arg Gly Pro Ser His Val Val Asp Thr Ala Cys Ser Ser Ser Leu
 20 25 30

45

Tyr Ala Leu His Ser Ala Cys Phe Gly Pro Leu Asn Ser Arg Asp Cys
 35 40 45

50 Asp Gly Ala Val Val Ala Ala Ala Asn Leu Ile Gln Ser Pro Glu Gln
 50 55 60

55 Gln Met Ile Ala Val Lys Arg Asp Ser Ile Ala
 65 70 75

2009274832 06 Jun 2011

68

<210> 160
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 <213> Penicillium coprobium PF1169

5

<400> 160

Pro Trp Pro Thr Thr Gly Leu Arg Arg Ala Ser Val Asn Ser Phe Gly
 1 5 10 15

10

Tyr Gly Gly Thr Asn Ala His Cys Val Leu Asp Asp
 20 25

15

<210> 161
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<400> 161

Gln Leu Gly Thr Thr Cys Val Gln Met Ala Leu Thr Lys Tyr Trp Thr
 1 5 10 15

25

Ser Leu Gly Val Thr Pro Ser Phe Val Met Gly His Ser Leu Gly Glu
 20 25 30

30

Phe Ala Ala Leu Asn Ala Ala Gly Val Leu Thr Ile Ser Asp Thr Ile
 35 40 45

35 Tyr Leu Ala Gly Arg Arg Ala Gln Leu Leu Thr Glu Gln Ile Glu Gly
 50 55 60

40

<210> 162
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 <212> PRT
 <213> Penicillium coprobium PF1169

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<400> 162

Ile Ala Pro Asn Ile His Phe Lys Met Pro Asn Pro Gln Ile Pro Phe
 1 5 10 15

50 Asn Glu Ala Asn Leu His Val Pro Leu Glu Pro Thr Pro Trp Pro Ala
 20 25 30

55 Gly Arg Pro Glu Arg Ile Ser Val Asn Ser Phe Gly Ile Gly Gly Ser
 35 40 45

2009274832 06 Jun 2011

69

Asn Ala His Ala Ile Leu Glu Ser Ala Ser Thr Val
 50 55 60

5 <210> 163
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 <212> PRT
 <213> Penicillium coprobium PF1169

10 <400> 163

Gly Leu Val Asn Ile Leu Arg Ser Trp Gly Ile Glu Pro Ser Thr Val
 1 5 10 15

15 Val Gly His Ser Ser Gly Glu Ile Val Ala Ala Tyr Thr Ala Arg Ala
 20 25 30

20 Ile Ser

25 <210> 164
 <211> 51
 <212> PRT
 <213> Penicillium coprobium PF1169

30 <400> 164

Pro Trp Pro Ser Glu Gly Leu Arg Arg Ile Ser Val Asn Ser Phe Gly
 1 5 10 15

35 Phe Gly Gly Ser Asn Thr His Val Ile Leu Asp Asp Ala Leu His Tyr
 20 25 30

40 Met Gln Gln Arg Gly Leu Thr Gly Asn His Cys Thr Ala Arg Leu Pro
 35 40 45

Gly Ile Leu
 50

45

<210> 165
 <211> 71
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<220>
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 55 <222> (5)..(5)
 <223> Xaa can be any naturally occurring amino acid

2009274832 06 Jun 2011

70

<400> 165

Ile Gly His Thr Xaa Gly Ser Ala Gly Leu Ala Ser Leu Ile Gly Ser
 1 5 10 15

5

Ser Leu Ala Met Lys His Gly Val Ile Pro Pro Asn Leu His Phe Gly
 20 25 30

10

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

15

Thr Glu Pro Val Pro Trp Pro Asn Ser Thr Ser Ser Gln Val Lys Arg
 50 55 60

20

Ala Ser Ile Asn Ser Phe Gly
 65 70

<210> 166

<211> 74

25

<212> PRT

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<400> 166

30

Gly Ser Asn Thr Ala Val Tyr Ser Gly Ser Met Thr Asn Asp Tyr Glu
 1 5 10 15

35

Leu Leu Ser Thr Arg Asp Ile Tyr Asp Met Pro His Asn Ser Ala Thr
 20 25 30

40

Gly Asn Gly Arg Thr Met Leu Ala Asn Arg Leu Ser Trp Phe Phe Asp
 35 40 45

45

Leu Gln Gly Pro Ser Ile Met Met Asp Thr Ala Cys Ser Ser Ser Leu
 50 55 60

Thr Ala Val His Leu Ala Ala Gln Ser Leu
 65 70

50

<210> 167

<211> 85

<212> PRT

<213> Penicillium coprobium PF1169

55

<400> 167

Asp Ala Gln Phe Phe Gly Thr Lys Pro Val Glu Ala Asn Ser Ile Asp

2009274832 06 Jun 2011

71

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5	Pro	Gln	Gln	Arg	Leu	Leu	Leu	Glu	Thr	Val	Tyr	Glu	Gly	Leu	Glu	Thr
				20					25					30		
10	Ser	Gly	Ile	Pro	Met	Glu	Arg	Leu	Gln	Gly	Ser	Asn	Thr	Ala	Val	Tyr
			35					40					45			
15	Val	Gly	Leu	Met	Thr	Asn	Asp	Tyr	Ala	Asp	Met	Leu	Gly	Arg	Asp	Met
		50					55					60				
	Gln	Asn	Phe	Pro	Thr	Tyr	Phe	Ala	Ser	Gly	Thr	Ala	Arg	Ser	Ile	Leu
	65					70					75				80	
20	Ser	Asn	Arg	Val	Ser											
					85											
25	<210>	168														
	<211>	60														
	<212>	PRT														
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30	<400>	168														
	Val	Val	Ala	Cys	Val	Asn	Ser	Pro	Ala	Ser	Thr	Thr	Leu	Ser	Gly	Asp
	1				5					10					15	
35	Val	Asp	Tyr	Ile	Asn	Gln	Leu	Glu	Ala	Arg	Leu	Gln	Gln	Asp	Gly	His
				20					25					30		
40	Phe	Ala	Arg	Lys	Leu	Arg	Ile	Asp	Thr	Ala	Tyr	His	Ser	Pro	His	Met
			35					40					45			
45	Glu	Glu	Leu	Val	Gly	Val	Val	Gly	Asp	Ala	Ile	Ser				
		50					55					60				
50	<210>	169														
	<211>	56														
	<212>	PRT														
	<213>	Penicillium coprobium PF1169														
	<400>	169														
55	Phe	Tyr	Gly	Met	Thr	Ser	Asp	Asp	Tyr	Arg	Glu	Val	Asn	Ser	Gly	Gln
	1				5					10					15	

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Asp Ile Asp Thr Tyr Phe Ile Pro Gly Gly Asn Arg Ala Phe Thr Pro
      20                      25                      30

5  Gly Arg Ile Asn Tyr Tyr Phe Lys Phe Ser Gly Pro Ser Val Ser Val
      35                      40                      45

      Asp Thr Ala Cys Ser Ser Ser Leu
      50                      55

10  <210> 170
      <211> 53
      <212> PRT
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      <400> 170

20  Val Ala Ile Val Gly Gly Val Asn Ala Leu Cys Gly Pro Gly Leu Thr
      1                      5                      10                      15

      Arg Val Leu Asp Lys Ala Gly Ala Ile Ser Ser Asp Gly Ser Cys Lys
      20                      25                      30

      Ser Phe Asp Asp Asp Ala His Gly Tyr Ala Arg Gly Glu Gly Ala Gly
      35                      40                      45

30  Ala Leu Val Thr Lys
      50

35  <210> 171
      <211> 40
      <212> PRT
      <213> Penicillium coprobium PF1169

40  <400> 171

      Gln Leu Gly Thr Thr Cys Val Gln Met Ala Leu Thr Lys Tyr Trp Thr
      1                      5                      10                      15

45  Ser Leu Gly Val Thr Pro Ser Phe Val Met Gly His Ser Leu Gly Glu
      20                      25                      30

50  Phe Ala Ala Leu Asn Ala Ala Gly
      35                      40

55  <210> 172
      <211> 69
      <212> PRT

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2009274832 06 Jun 2011

73

<213> Penicillium coprobium PF1169

<400> 172

5 Arg Glu Trp Met Thr Ala Glu Gly Lys Asp His Asn Leu Ser Asp Ile
 1 5 10 15

10 Leu Thr Thr Leu Ala Thr Arg Arg Asp His His Asp Tyr Arg Ala Ala
 20 25 30

15 Leu Val Val Asp Asp Asn Arg Asp Ala Glu Leu Ala Leu Gln Ala Leu
 35 40 45

Glu His Gly Val Asp Gln Thr Phe Thr Thr Gln Ser Arg Val Phe Gly
 50 55 60

20 Ala Asp Ile Ser Lys
 65

25 <210> 173
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 <212> PRT
 <213> Penicillium coprobium PF1169

30 <400> 173

Pro Trp Pro Ser Glu Gly Leu Arg Arg Ile Ser Val Asn Ser Phe Gly
 1 5 10 15

35 Phe Gly Gly Ser Asn Thr His Val Ile Leu Asp Asp Ala Leu His Tyr
 20 25 30

40 Met Gln Gln Arg Gly Leu Thr Gly Asn His Cys Thr Ala Arg Leu Pro
 35 40 45

45 Gly Ile Leu
 50

50 <210> 174
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 <213> Penicillium coprobium PF1169

<400> 174

55 Phe Val Glu Met His Gly Thr Gly Thr Lys Ala Gly Asp Pro Val Glu
 1 5 10 15

2009274832 06 Jun 2011

74

Ala Ala Ala Val His Ala Ala Leu Gly Lys Asn Arg Thr Leu Arg Asn
 20 25 30

5 Pro Leu Tyr Ile Gly Ser Val Lys Ser Asn Ile Gly His Leu Glu Gly
 35 40 45

10 Ala Ser Gly Ile Val Ala Val Ile Lys Ala Ala Met Met Leu Asp Arg
 50 55 60

15 Asp Leu Met Leu Pro Asn Ala
 65 70

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 20 <212> PRT
 <213> Penicillium coprobium PF1169

<400> 175

25 Leu Ala Ile Val Gly Met Ala Cys Arg Leu Pro Gly Gln Ile Thr Thr
 1 5 10 15

30 Pro Gln Glu Leu Trp Glu Leu Cys Ser Arg Gly Arg Ser Ala Trp Ser
 20 25 30

Glu Ile Pro Pro Glu Arg Phe Asn Pro
 35 40

35

<210> 176
 <211> 64
 <212> PRT
 40 <213> Penicillium coprobium PF1169

<400> 176

45 Gln Leu Gly Thr Thr Cys Val Gln Met Ala Leu Thr Lys Tyr Trp Thr
 1 5 10 15

Ser Leu Gly Val Thr Pro Ser Phe Val Met Gly His Ser Leu Gly Glu
 20 25 30

50

Phe Ala Ala Leu Asn Ala Ala Gly Val Leu Thr Ile Ser Asp Thr Ile
 35 40 45

55 Tyr Leu Ala Gly Arg Arg Ala Gln Leu Leu Thr Glu Gln Ile Glu Gly
 50 55 60

2009274832 06 Jun 2011

75

5 <210> 177
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 <213> *Penicillium coprobium* PF1169

 <400> 177
 10 Gly Ala Ser Val Tyr Val Leu Ala Leu Asp Ile Thr Lys Pro Asp Ala
 1 5 10 15

 Val Glu Gln Leu Ser Thr Ala Leu Asp Arg Leu Ala Leu Pro Ser Val
 15 20 25 30

 Gln Gly Val Val His Ala Ala Gly Val Leu Asp Asn Glu Leu Val Met
 20 35 40 45

 Gln Thr Thr Gln Glu Ala Phe Asn Arg Val Leu Ala Pro Lys Ile Ala
 50 55 60
 25 Gly Ala Leu Ala Leu His Glu Pro Phe Pro
 65 70
 30 <210> 178
 <211> 72
 <212> PRT
 <213> *Penicillium coprobium* PF1169
 35 <400> 178

 Gly Leu Val Asn Ile Leu Arg Ser Trp Gly Ile Glu Pro Ser Thr Val
 1 5 10 15
 40 Val Gly His Ser Ser Gly Glu Ile Val Ala Ala Tyr Thr Ala Arg Ala
 20 25 30
 45 Ile Ser Met Arg Thr Ala Ile Ile Leu Ala Tyr Tyr Arg Gly Lys Val
 35 40 45

 Ala Gln Pro Leu Glu Gly Leu Gly Ala Met Val Ala Val Gly Leu Ser
 50 50 55 60

 Pro Asp Glu Val Ala Gln Tyr Met
 55 65 70

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76

<211> 70
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5 <400> 179

Gly Arg Phe Leu Ser Ser Asp Gly Arg Cys His Thr Phe Asp Glu Lys
 1 5 10 15

10 Ala Asn Gly Tyr Ala Arg Gly Glu Ala Val Gly Cys Leu Ile Leu Lys
 20 25 30

15 Pro Leu Ala Lys Ala Leu His Asp Gln Asn Lys Ile Arg Ala Val Ile
 35 40 45

20 Arg Gly Thr Gly Ser Asn Gln Asp Gly Arg Thr Ala Gly Ile Thr Val
 50 55 60

25 Pro Asn Gly Ala Ala Gln
 65 70

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 30 <213> Penicillium coprobium PF1169

<400> 180

35 Ser Ser Phe Leu Thr Ser Thr Val Gln Gln Ile Val Glu Glu Thr Ile
 1 5 10 15

40 Gln Gly Gly Thr Gly Gln Val Val Met Glu Ser Asp Leu Met Gln Thr
 20 25 30

Glu Phe Leu Glu Ala Ala Asn Gly His Arg Met Asn Asp Cys Gly Val
 35 40 45

45 Val Thr Ser
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55 <400> 181

Leu Leu Gly Leu Arg Leu Lys Trp Lys Glu Tyr His Gln Asp Phe Asn

2009274832 06 Jun 2011

2009274832 06 Jun 2011

77

1 5 10 15

5 Ala Ala His Arg Val Leu Pro Leu Pro Ser Tyr Lys Trp Asp Leu Lys
20 25 30

10 Asn Tyr Trp Ile Pro Tyr Thr Asn Asn Phe Cys Leu Leu Lys Gly Ala
35 40 45

15 Pro Ala Ala Pro Val Ala Glu Ala Thr Pro Ile Ser Val Phe Leu Ser
50 55 60

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<213> Penicillium coprobium PF1169

20 <400> 182

25 Ser Phe Arg Arg Gln Glu Asp Thr Trp Lys Val Leu Ser Asn Ala Thr
1 5 10 15

Ser Thr Leu Tyr Leu Ala Gly Ile Glu Ile
20 25

30 <210> 183
<211> 65
<212> PRT
<213> Penicillium coprobium PF1169

35 <400> 183

40 Ala Gly Gly Asn Thr Thr Val Ala Leu Glu Asp Ala Pro Ile Arg Thr
1 5 10 15

Arg Ser Gly Ser Asp Pro Arg Ser Leu His Pro Ile Ala Ile Ser Ala
20 25 30

45 Lys Ser Lys Val Ser Leu Arg Gly Asn Leu Glu Asn Leu Leu Ala Tyr
35 40 45

50 Leu Asp Thr His Pro Asp Val Ser Leu Ser Asp Leu Ser Tyr Thr Thr
50 55 60

55 Thr
65

78

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<400> 184

Phe Asp Ala Ala Phe Phe Asn Met Ser Pro Arg Glu Ala Gln Gln Thr
 1 5 10 15

10

Asp Pro Met Gln Arg Leu Ala Ile Val Thr Ala Tyr Glu Ala Leu Glu
 20 25 30

15

Arg Ala Gly Tyr Val Ala Asn Arg Thr Ala Ala Thr Asn Leu His Arg
 35 40 45

20

Ile Gly Thr Phe Tyr Gly Gln Ala Ser Asp Asp Tyr Arg Glu Val Asn
 50 55 60

25

Thr Ala Gln Glu Ile Ser Thr Tyr Phe Ile Pro Gly Gly Cys Arg Ala
 65 70 75 80

30

Phe Gly Pro Gly Arg Ile Asn Tyr Phe Phe Lys Phe Leu Gly Pro Ala
 85 90 95

35

<210> 185
 <211> 58
 <212> PRT
 <213> Penicillium coprobium PF1169

<400> 185

40

Phe Leu Gln Ile Ser Gly Pro Ser Phe Ser Ile Asp Thr Ala Cys Ser
 1 5 10 15

45

Ser Ser Leu Ala Thr Ile Gln Val Cys Thr His Leu Phe His Val His
 20 25 30

50

Leu Asn Arg Gln Leu Thr Ile Ala Ala Cys Thr Ser Leu Trp Asn Gly
 35 40 45

Glu Thr Asp Thr Val Val Ala Gly Gly Met
 50 55

55

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2009274832 06 Jun 2011

2009274832 06 Jun 2011

79

<213> Penicillium coprobium PF1169

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5 Val Tyr Ser Gly Ser Met Thr Asn Asp Tyr Glu Leu Leu Ser Thr Arg
1 5 10 15

10 Asp Ile Tyr Asp Met Pro His Asn Ser Ala Thr Gly Asn Gly Arg Thr
20 25 30

15 Met Leu Ala Asn Arg Leu Ser Trp Phe Phe Asp Leu Gln Gly Pro Ser
35 40 45

20 Ile Met Met Asp Thr Ala Cys Ser Ser Ser Leu
50 55

<210> 187
<211> 31
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<213> Penicillium coprobium PF1169

25 <400> 187

30 Leu Phe Leu Phe Pro Asp Gly Ser Gly Ser Ala Thr Ser Tyr Ala Thr
1 5 10 15

Ile Pro Gly Ile Ser Pro Asp Val Cys Val Tyr Gly Leu Asn Cys
20 25 30

35 <210> 188
<211> 26
<212> PRT
<213> Penicillium coprobium PF1169

40 <400> 188

45 Ala Lys His Pro Pro Ala Thr Ser Ile Leu Leu Gln Gly Asn Pro Lys
1 5 10 15

Thr Ala Thr Gln Ser Phe Ile Phe Val Pro
20 25

50 <210> 189
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<212> PRT
<213> Penicillium coprobium PF1169

55 <400> 189

2009274832 06 Jun 2011

80

Gly Asn Gly Ser Ala Met Ile Ser Asn Arg Ile Ser Trp Phe Phe Asp
 1 5 10 15

5 Leu Lys Gly Pro Ser Leu Ser Leu Asp Thr Ala Cys Ser Ser Ser Leu
 20 25 30

10 Val Ala Leu His Leu Ala
 35

15 <210> 190
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 <213> Penicillium coprobium PF1169
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20 Ala Ile His His Gly Val Gln Ala Ile Lys Leu Gly Glu Ser Arg Val
 1 5 10 15

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

30 Val Leu Asp Lys Ala Gly Ala Ile Ser Ser Asp Gly Ser Cys Lys Ser
 35 40 45

Phe Asp Asp Asp Ala His Gly Tyr Ala Arg Gly Glu Gly Ala Gly Ala
 50 55 60

35 Leu Val Leu Lys Ser Leu His Gln Ala Leu Leu Asp
 65 70 75

40 <210> 191
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 <213> Penicillium coprobium PF1169

45 <400> 191

Val Trp Ile Glu Ile Gly Pro His Pro Val Cys Leu Gly Phe Val Lys
 1 5 10 15

50 Ala Thr Leu Glu Ser Val Ala Val Ala Val Pro Ser Leu Arg Arg Gly
 20 25 30

55 Glu Asn Ala Trp Cys Thr Leu Ala Gln Ser Leu Thr Thr Leu His Asn
 35 40 45

2009274832 06 Jun 2011

81

Ala Gly Val Pro Val Gly Trp Ser Glu Phe His Arg Pro Phe Glu Arg
50 55 60

5
Ala
65

10 <210> 192
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<212> PRT
<213> Penicillium coprobium PF1169

15 <400> 192

Thr Ser Asp Asp Tyr Arg Glu Val Asn Ser Gly Gln Asp Ile Asp Thr
1 5 10 15

20 Tyr Phe Ile Pro Gly Gly Asn Arg Ala Phe Thr Pro Gly Arg Ile Asn
20 25 30

25 Tyr Tyr Phe Lys Phe Ser Gly Pro Ser Val Ser Val Asp Thr Ala Cys
35 40 45

30 Ser Ser Ser Leu Ala
50

35 <210> 193
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<212> PRT
<213> Penicillium coprobium PF1169

<400> 193

40 Val Asp Thr Ala Cys Ser Ser Ser Leu Tyr Ala Leu His Ser Ala Cys
1 5 10 15

45 Phe Gly Pro Leu Asn Ser Arg Asp Cys Asp Gly Ala Val Val Ala Ala
20 25 30

50 Ala Asn Leu Ile Gln Ser Pro Glu
35 40

55 <210> 194
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<213> Penicillium coprobium PF1169

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82

Met Leu Ala Val Gly Ala Ser Ala Ser Asp Ile Gln Gln Ile Leu Asp
 1 5 10 15
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 Ala Met Arg Gly Asn Lys Ala Val Ile Ala Cys Val Asn Ser Glu Ser
 20 25 30
 10 Ser Val Thr Leu Ser Gly Asp Leu Asp Val Ile Ala Asn Leu Gln Thr
 35 40 45
 15 Ala Leu Asp Lys Glu Gly Ile Phe Thr Arg Lys Leu Lys Val Asp Val
 50 55 60
 Ala Tyr His Ser
 65
 20
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 <211> 62
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 25 <213> Penicillium coprobium PF1169
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 30 Phe Leu Asp Asp Leu Ala Phe Thr Val Asn Glu Arg Arg Ser Ile Phe
 1 5 10 15
 Pro Trp Lys Ala Ala Val Val Gly Asp Thr Met Glu Gly Leu Ala Ala
 20 25 30
 35 Ser Leu Ala Gln Asn Ile Lys Pro Arg Ser Val Leu Arg Met Pro Thr
 35 40 45
 40 Leu Gly Phe Val Phe Thr Gly Gln Gly Ala Gln Trp Pro Gly
 50 55 60
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 <211> 76
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 Gly Pro Ser Met Thr Ile Asp Thr Ala Cys Ser Ser Ser Leu Ile Ala
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 55 Leu His Gln Ala Val Gln Ser Leu Arg Ser Gly Glu Thr Asp Val Ala
 20 25 30

2009274832 06 Jun 2011

83

Val Ala Ala Gly Thr Asn Leu Leu Leu Gly Pro Glu Gln Tyr Ile Ala
35 40 45

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Glu Ser Lys Leu Lys Met Leu Ser Pro Asn Gly Arg Ser Arg Met Trp
50 55 60

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Asp Lys Asp Ala Asp Gly Tyr Ala Arg Gly Asp Gly
65 70 75

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<400> 197

Ser Val Pro Ile Glu Glu His Ser Pro Val Val Thr Gln Leu Gly Thr
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25

Thr Cys Val Gln Met Ala Leu Thr Lys Tyr Trp Thr Ser Leu Gly Val
20 25 30

30

Thr Pro Ser Phe Val Met Gly His Ser Leu Gly Glu Phe Ala Ala Leu
35 40 45

35

Asn Ala Ala Gly Val Leu Thr Ile Ser Asp Thr Ile Tyr Leu Ala Gly
50 55 60

40

Arg Arg Ala Gln Leu Leu Thr Glu Gln Ile Glu Gly Gly Thr His
65 70 75

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<211> 40

<212> PRT

45 <213> Penicillium coprobium PF1169

<400> 198

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

55

Gln Asp Leu Met Leu Ala Gly Gly

2009274832 06 Jun 2011

84

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40

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<210> 199
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<400> 199

Gly Arg Phe Leu Ser Ser Asp Gly Arg Cys His Thr Phe Asp Glu Lys
 1 5 10 15

15

Ala Asn Gly Tyr Ala Arg Gly Glu Ala Val Gly Cys Leu Ile Leu Lys
 20 25 30

20

Pro Leu Ala Lys Ala Leu His Asp Gln Asn Lys Ile Arg Ala Val Ile
 35 40 45

25

Arg Gly Thr Gly Ser Asn Gln Asp Gly Arg Thr Ala Gly Ile Thr Val
 50 55 60

30

Pro Asn Gly Ala Ala Gln
 65 70

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<400> 200

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Leu Ser Val Lys Arg Val Gly Ile His Asp Asp Phe Phe Glu Leu Gly
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45

Gly His Ser Leu Leu Ala Val Lys Leu Val Asn His Leu Lys Lys Val
 20 25 30

Phe Gly Thr Glu Leu Ser Val Ala Leu Leu Ala Gln Tyr Ser Thr Val
 35 40 45

50

Glu Ser Leu Gly Glu Ile Ile Arg Glu Asn Lys Glu Ile Lys Pro Ser
 50 55 60

55

Ile Val Ile Glu Leu Arg Ser Gly Thr Tyr Glu Gln Pro Leu Trp Leu
 65 70 75 80

2009274832 06 Jun 2011

2009274832 06 Jun 2011

85

Phe His Pro Ile Gly Gly Ser Thr Phe Cys Tyr Met Glu Leu Ser Arg
 85 90 95

5 His Leu Asn Pro Asn Arg Thr Leu Arg Ala Ile Gln Ser Pro Gly Leu
 100 105 110

10 Ile Glu Ala Asp Ala Ala Glu Val Ala Ile Glu Glu Met Ala Thr Leu
 115 120 125

15 Tyr Ile Ala Glu Met Gln Lys Met Gln Pro Gln Gly Pro Tyr Phe Leu
 130 135 140

20 Gly Gly Trp Cys Phe Gly Gly Ala Ile Ala Tyr Glu Ile Ser Arg Gln
 145 150 155 160

Leu Arg Gln Met Gly Gln Gln Val Thr Gly Ile Val Met Ile Asp Thr
 165 170 175

25 Arg Ala Pro Ile Pro Glu Asn Val Pro Glu Asp Ala Asp Asp Ala Met
 180 185 190

30 Leu Leu Ser Trp Phe Ala Arg Asp Leu Ala Val Pro Tyr Gly Lys Lys
 195 200 205

35 Leu Thr Ile Ser Ala Gln Tyr Leu Arg Glu Leu Ser Pro Asp His Met
 210 215 220

Phe Asp His Val Leu Lys Glu Ala Lys Ala Ile Asn Val Ile Pro Leu
 225 230 235 240

40 Asp Ala Asn Pro Ser Asp Phe Arg Leu Tyr Phe Asp Thr Tyr Leu Ala
 245 250 255

45 Asn Gly Val Ala Leu Gln Thr Tyr Phe Pro Glu Pro Glu Asp Phe Pro
 260 265 270

50 Ile Leu Leu Val Lys Ala Lys Asp Glu Ser Glu Asp
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 55 <212> PRT
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86

<400> 201

Pro Met Asn Lys Asp Lys Val Tyr Trp Ser Ala Ile Ile Arg Thr Leu
 1 5 10 15

5

Val Ala Lys Glu Met Arg Val Glu Pro Glu Thr Ile Asp Pro Glu Gln
 20 25 30

10

Lys Phe Thr Thr Tyr Gly Leu Asp Ser Ile Val Ala Leu Ser Val Ser
 35 40 45

15

Gly Asp Leu Glu Asp Leu Thr Lys Leu Glu Leu Glu Pro Thr Leu Leu
 50 55 60

20

Trp Asp Tyr Pro Thr Ile Asn Ala Leu
 65 70

25

<210> 202
 <211> 63
 <212> PRT
 <213> Penicillium coprobium PF1169

<400> 202

30

Gly Ser Leu Ile Asp Ile Glu Glu Pro Ile Ile Pro Leu Ser Thr Met
 1 5 10 15

35

Arg Tyr Ile Gln Gly Ala Asp Ile Val Arg Ile Ser Asp Gly Ile Ala
 20 25 30

40

Arg Thr Ser Arg Phe Arg Ser Leu Pro Arg Thr Lys Leu Arg Pro Val
 35 40 45

45

Ser Asp Gly Pro Arg Leu Leu Pro Arg Pro Glu Gly Thr Tyr Leu
 50 55 60

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<210> 203
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 <212> PRT
 <213> Penicillium coprobium PF1169

<400> 203

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

Arg Gly Lys Glu Ile Gly Cys Phe Val Gly Val Phe Gly Glu Asp Trp

2009274832 06 Jun 2011

2009274832 06 Jun 2011

87

				20					25					30			
5	Leu	Glu	Met	Ser	His	Lys	Asp	Pro	Gln	His	Leu	Asn	Gln	Met	Phe	Pro	
			35					40					45				
10	Ile	Ala	Thr	Gly	Gly	Phe	Ala	Leu	Ala	Asn	Gln	Val	Ser	Tyr	Arg	Phe	
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15	Asp	Leu	Thr	Gly	Pro												
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	1				5					10					15		
30	Asp	Pro	Met	Gln	Arg	Leu	Ala	Ile	Val	Thr	Ala	Tyr	Glu	Ala	Leu	Glu	
				20					25					30			
35	Arg	Ala	Gly	Tyr	Val	Ala	Asn	Arg	Thr	Ala	Ala	Thr	Asn	Leu	His	Arg	
			35					40					45				
40	Ile	Gly	Thr	Phe	Tyr	Gly	Gln	Ala	Ser	Asp	Asp	Tyr	Arg	Glu	Val	Asn	
		50					55					60					
45	Thr	Ala	Gln	Glu	Ile	Ser	Thr	Tyr	Phe	Ile	Pro	Gly	Gly	Cys	Arg	Ala	
	65					70					75					80	
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55	<210>	205															
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	1				5					10					15		

2009274832 06 Jun 2011

88

Ser Ser Leu Ala Thr Ile Gln Val Cys Thr His Leu Phe His Val His
20 25 30

5 Leu Asn Arg Gln Leu Thr Ile Ala Ala Cys Thr Ser Leu Trp Asn Gly
35 40 45

10 Glu Thr Asp Thr Val Val Ala Gly Gly Met
50 55

15 <210> 206
<211> 52
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<213> Penicillium coprobium PF1169

<400> 206

20 Glu Leu Arg His Gly Lys Asn Ile Asp Lys Pro Glu Tyr Ser Gln Pro
1 5 10 15

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

30 Gly Val Val Pro Lys Ala Val Val Gly His Ser Ser Gly Glu Ile Ala
35 40 45

Ala Ala Tyr Val
50

35 <210> 207
<211> 59
<212> PRT
<213> Penicillium coprobium PF1169

40 <400> 207

45 Val Tyr Ser Gly Ser Met Thr Asn Asp Tyr Glu Leu Leu Ser Thr Arg
1 5 10 15

Asp Ile Tyr Asp Met Pro His Asn Ser Ala Thr Gly Asn Gly Arg Thr
20 25 30

50 Met Leu Ala Asn Arg Leu Ser Trp Phe Phe Asp Leu Gln Gly Pro Ser
35 40 45

55 Ile Met Met Asp Thr Ala Cys Ser Ser Ser Leu
50 55

2009274832 06 Jun 2011

89

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 <211> 28
 <212> PRT
 5 <213> Penicillium coprobium PF1169

 <400> 208
 10 Pro Trp Pro Thr Thr Gly Leu Arg Arg Ala Ser Val Asn Ser Phe Gly
 1 5 10 15

 Tyr Gly Gly Thr Asn Ala His Cys Val Leu Asp Asp
 20 25
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 <210> 209
 <211> 71
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 20 <213> Penicillium coprobium PF1169

 <400> 209
 25 Lys Ala Ser Leu Ser Leu Gln His Gly Met Ile Ala Pro Asn Leu Leu
 1 5 10 15

 Met Gln His Leu Asn Pro Lys Ile Lys Pro Phe Ala Ala Lys Leu Ser
 20 25 30
 30

 Val Pro Thr Glu Cys Ile Pro Trp Pro Ala Val Pro Asp Gly Cys Pro
 35 40 45
 35

 Arg Arg Ala Ser Val Asn Ser Phe Gly Phe Gly Gly Ala Asn Val His
 50 55 60

 40 Val Val Leu Glu Ser Tyr Thr
 65 70

 <210> 210
 45 <211> 80
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 <213> Penicillium coprobium PF1169

 <400> 210
 50 Leu Lys Gly Thr Gly Gly Gln Met Leu Gln Asn Val Val Leu Arg Val
 1 5 10 15

 55 Pro Val Ala Ile Asn Ala Pro Arg Ser Val Gln Val Val Val Gln Gln
 20 25 30

2009274832 06 Jun 2011

90

Asp Gln Val Lys Val Val Ser Arg Leu Ile Pro Ser Glu Ala Ser Val
 35 40 45

5
 Leu Asp Asp Asp Ala Ser Trp Val Thr His Thr Thr Ala Tyr Trp Asp
 50 55 60

10
 Arg Arg Val Leu Gly Ser Glu Asp Arg Ile Asp Leu Ala Ala Val Lys
 65 70 75 80

15
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 <212> PRT
 <213> *Penicillium coprobium* PF1169

20
 Gly Asn Gly Ser Ala Met Ile Ser Asn Arg Ile Ser Trp Phe Phe Asp
 1 5 10 15

25
 Leu Lys Gly Pro Ser Leu Ser Leu Asp Thr Ala Cys Ser Ser Ser Leu
 20 25 30

30
 Val Ala Leu His Leu Ala
 35

35
 <210> 212
 <211> 76
 <212> PRT
 <213> *Penicillium coprobium* PF1169

<400> 212

40
 Ala Ile His His Gly Val Gln Ala Ile Lys Leu Gly Glu Ser Arg Val
 1 5 10 15

45
 Ala Ile Val Gly Gly Val Asn Ala Leu Cys Gly Pro Gly Leu Thr Arg
 20 25 30

50
 Val Leu Asp Lys Ala Gly Ala Ile Ser Ser Asp Gly Ser Cys Lys Ser
 35 40 45

Phe Asp Asp Asp Ala His Gly Tyr Ala Arg Gly Glu Gly Ala Gly Ala
 50 55 60

55
 Leu Val Leu Lys Ser Leu His Gln Ala Leu Leu Asp
 65 70 75

2009274832 06 Jun 2011

91

5 <210> 213
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 <212> PRT
 <213> Penicillium coprobium PF1169

 <400> 213
 10 Arg Glu Trp Met Thr Ala Glu Gly Lys Asp His Asn Leu Ser Asp Ile
 1 5 10 15

 15 Leu Thr Thr Leu Ala Thr Arg Arg Asp His His Asp Tyr Arg Ala Ala
 20 25 30

 20 Leu Val Val Asp Asp Asn Arg Asp Ala Glu Leu Ala Leu Gln Ala Leu
 35 40 45

 Glu His Gly Val Asp Gln Thr Phe Thr Thr Gln Ser Arg Val Phe Gly
 50 55 60
 25 Ala Asp Ile Ser Lys
 65
 30 <210> 214
 <211> 53
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 <213> Penicillium coprobium PF1169
 35 <400> 214

 Thr Ser Asp Asp Tyr Arg Glu Val Asn Ser Gly Gln Asp Ile Asp Thr
 1 5 10 15
 40 Tyr Phe Ile Pro Gly Gly Asn Arg Ala Phe Thr Pro Gly Arg Ile Asn
 20 25 30
 45 Tyr Tyr Phe Lys Phe Ser Gly Pro Ser Val Ser Val Asp Thr Ala Cys
 35 40 45

 50 Ser Ser Ser Leu Ala
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 55 <210> 215
 <211> 63
 <212> PRT
 <213> Penicillium coprobium PF1169

92

<400> 215

Ala Gly Ile Pro Leu Ala Asn Ile Met Gly Thr Lys Thr Ser Cys Phe
 1 5 10 15

5

Val Gly Ser Phe Ser Ala Asp Tyr Thr Asp Leu Leu Leu Arg Asp Pro
 20 25 30

10

Glu Cys Val Pro Met Tyr Gln Cys Thr Asn Ala Gly Gln Ser Arg Ala
 35 40 45

15

Met Thr Ala Asn Arg Leu Ser Tyr Phe Phe Asp Leu Lys Gly Pro
 50 55 60

20

<210> 216

<211> 68

<212> PRT

<213> Penicillium coprobium PF1169

25

<400> 216

Met Leu Ala Val Gly Ala Ser Ala Ser Asp Ile Gln Gln Ile Leu Asp
 1 5 10 15

30

Ala Met Arg Gly Asn Lys Ala Val Ile Ala Cys Val Asn Ser Glu Ser
 20 25 30

35

Ser Val Thr Leu Ser Gly Asp Leu Asp Val Ile Ala Asn Leu Gln Thr
 35 40 45

40

Ala Leu Asp Lys Glu Gly Ile Phe Thr Arg Lys Leu Lys Val Asp Val
 50 55 60

Ala Tyr His Ser
 65

45

<210> 217

<211> 39

<212> PRT

<213> Penicillium coprobium PF1169

50

<400> 217

Asn Ala Ala Gly Ala His Phe Leu Thr Glu Asp Ile Gly Leu Phe Asp
 1 5 10 15

55

Ala Pro Phe Phe Asn Ile Thr Leu Gln Glu Ala Gln Thr Met Asp Pro

2009274832 06 Jun 2011

2009274832 06 Jun 2011

93

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20	Leu His Gln Ala Val Gln Ser Leu Arg Ser Gly Glu Thr Asp Val Ala 20 25 30		
25	Val Ala Ala Gly Thr Asn Leu Leu Leu Gly Pro Glu Gln Tyr Ile Ala 35 40 45		
30	Glu Ser Lys Leu Lys Met Leu Ser Pro Asn Gly Arg Ser Arg Met Trp 50 55 60		
35	Asp Lys Asp Ala Asp Gly Tyr Ala Arg Gly Asp Gly 65 70 75		
40	<210> 219 <211> 61 <212> PRT <213> Penicillium coprobium PF1169 <400> 219		
45	Gly Leu Val Asn Ile Leu Arg Ser Trp Gly Ile Glu Pro Ser Thr Val 1 5 10 15		
50	Val Gly His Ser Ser Gly Glu Ile Val Ala Ala Tyr Thr Ala Arg Ala 20 25 30		
55	Ile Ser Met Arg Thr Ala Ile Ile Leu Ala Tyr Tyr Arg Gly Lys Val 35 40 45		
	Ala Gln Pro Leu Glu Gly Leu Gly Ala Met Val Ala Val 50 55 60		

2009274832 06 Jun 2011

94

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 1 5 10 15
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 20 25 30
 Thr Pro Ser Phe Val Met Gly His Ser Leu Gly Glu Phe Ala Ala Leu
 35 40 45
 20 Asn Ala Ala Gly Val Leu Thr Ile Ser Asp Thr Ile Tyr Leu Ala Gly
 50 55 60
 Arg Arg Ala Gln Leu Leu Thr Glu Gln Ile Glu Gly Gly Thr His
 25 65 70 75
 <210> 221
 <211> 81
 30 <212> PRT
 <213> Penicillium coprobium PF1169
 <400> 221
 35 Val Tyr Thr Gly Arg Ile Ser Leu Lys Asp Leu Gly Met Arg Cys Leu
 1 5 10 15
 Pro Leu Cys Leu Phe Leu Phe Leu Trp Thr Ile Tyr Phe Asn Thr Ala
 40 20 25 30
 Tyr Ser Tyr Gln Asp Ile Lys Asp Asp Cys Lys Leu Asn Val Asn Ser
 35 40 45
 45 Ser Tyr Val Leu Ala Gly Ser His Val Arg Gly Met Leu Leu Leu Gln
 50 55 60
 50 Ala Ile Ala Val Val Leu Val Ile Pro Trp Ile Leu Tyr Thr Ser Ala
 65 70 75 80
 55 Ser

2009274832 06 Jun 2011

95

<210> 222
 <211> 82
 <212> PRT
 5 <213> *Penicillium coprobium* PF1169

 <400> 222
 10 Arg His Phe Gly Leu Trp Asp Glu Pro Arg Glu Leu Glu Asp Val Glu
 1 5 10 15

 Phe Leu Leu Lys Ala Asp Val Arg Asn Asn Ser Ala Trp Asn His Arg
 20 25 30
 15

 Tyr Met Leu Arg Phe Gly Pro Arg Asp Thr Ser Leu Pro Asp Ala Gly
 35 40 45
 20
 Met Val Asn Ala Gly Asp Leu Ser Thr Ala Pro Ala Glu Lys Gly Arg
 50 55 60

 25 Leu Ser Val Val Asp Glu Asp Met Val Asp Gly Glu Leu Lys Phe Ala
 65 70 75 80

 Gln Glu
 30

 <210> 223
 <211> 35
 35 <212> PRT
 <213> *Penicillium coprobium* PF1169

 <400> 223
 40 Ile Met Arg Gly Ala Gly Cys Ala Ile Asn Asp Leu Trp Asp Arg Asn
 1 5 10 15

 Leu Asp Pro His Val Glu Arg Thr Lys Phe Arg Pro Ile Ala Arg Gly
 20 25 30
 45

 Ala Leu Ser
 35
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 <210> 224
 <211> 86
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 55 <213> *Penicillium coprobium* PF1169

 <400> 224

2009274832 06 Jun 2011

96

Phe Pro Thr Phe Pro Pro Lys Glu Ala Asp Phe Leu Met Glu Met Phe
 1 5 10 15
 5
 Ala Gln Asp Ser Lys Asn Tyr His Val Trp Thr Tyr Arg His Trp Leu
 20 25 30
 10 Val Arg His Phe Gly Leu Trp Asp Glu Pro Arg Glu Leu Glu Asp Val
 35 40 45
 15 Glu Phe Leu Leu Lys Ala Asp Val Arg Asn Asn Ser Ala Trp Asn His
 50 55 60
 20 Arg Tyr Met Leu Arg Phe Gly Pro Arg Asp Thr Ser Leu Pro Asp Ala
 65 70 75 80
 Gly Met Val Asn Ala Gly
 85
 25
 <210> 225
 <211> 82
 <212> PRT
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 30
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 Asn His Arg Tyr Met Leu Arg Phe Gly Pro Arg Asp Thr Ser Leu Pro
 1 5 10 15
 35 Asp Ala Gly Met Val Asn Ala Gly Asp Leu Ser Thr Ala Pro Ala Glu
 20 25 30
 40 Lys Gly Arg Leu Ser Val Val Asp Glu Asp Met Val Asp Gly Glu Leu
 35 40 45
 45 Lys Phe Ala Gln Glu Ala Ile Leu Arg Ala Pro Glu Asn Arg Ser Pro
 50 55 60
 50 Trp Trp Tyr Ala Arg Gly Val Leu Arg Ala Ala Gly Arg Gly Leu Gly
 65 70 75 80
 Glu Trp
 55
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<211> 45
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5 <400> 226

Arg Pro Thr Ser Arg Lys Leu Gly Val Tyr Pro Gln Tyr Ile Leu Gly
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10

Ala Ser Ser Ala Leu Thr Ile Leu Pro Ala Trp Ala Ser Val Tyr Thr
 20 25 30

15

Gly Arg Ile Ser Leu Lys Asp Leu Gly Met Arg Cys Leu
 35 40 45

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30

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40

<400> 228
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20

45

<210> 229
 <211> 20
 <212> DNA
 <213> Artificial Sequence

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<220>
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<400> 229
 cgcaagactt gaggaacaag

20

55

<210> 230
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 <212> DNA

2009274832 06 Jun 2011

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10 <210> 231
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15 <220>
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 cgctttttacg gcaatcatct 20

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30 <400> 232
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35 <210> 233
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 40 <223> a primer sequence for PCR
 <400> 233
 cagacgctgc ataggatcag 20

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 55 ttactagcct ctggggtgga 20

2009274832 06 Jun 2011

2009274832 06 Jun 2011

99

5 <210> 235
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 10 <400> 235
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 15 <210> 236
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 <212> DNA
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 20 <400> 236
 atgcggcctt tttcaacat 19

 25 <210> 237
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 30 <400> 237
 cgacgtaagg agctgtgagc 20
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 40 <210> 238
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 45 <400> 238
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 50 <210> 239
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 55 <400> 239

2009274832 06 Jun 2011

100

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10 <220>
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<400> 240
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15 <210> 241
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20 <220>
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<400> 241
 25 agagcatagc ccggttgta 20

<210> 242
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 30 <212> DNA
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35 <400> 242
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40 <210> 243
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45 <220>
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<400> 243
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50 <210> 244
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 55 <213> Artificial Sequence

<220>

2009274832 06 Jun 2011

101

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15 <220>
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20 <210> 246
 <211> 20
 <212> DNA
 <213> Artificial Sequence

25 <220>
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 <400> 246
 ccaggaagac acttggaagg 20

30 <210> 247
 <211> 20
 <212> DNA
 <213> Artificial Sequence

35 <220>
 <223> a primer sequence for PCR
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 agagcatagc ccggttgta 20

45 <210> 248
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 <213> Artificial Sequence

50 <220>
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 <400> 248
 ccaccttcga tttgctcagt 20

55 <210> 249
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 <212> DNA

2009274832 06 Jun 2011

102

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 <400> 249
 cgcaagactt gaggaacaag 20

10 <210> 250
 <211> 20
 <212> DNA
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15 <220>
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 <400> 250
 tccctctacg cagaagaacc 20

20 <210> 251
 <211> 20
 <212> DNA
 <213> Artificial Sequence

25 <220>
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 <400> 251
 agagcatagc ccggttgta 20

30 <210> 252
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35 <220>
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 <400> 252
 ccaccttcga ttgctcagt 20

40 <210> 253
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 gaggcgctgg ttagagaat 20

50 <210> 254
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55 <220>
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 <400> 254
 gaggcgctgg ttagagaat 20

2009274832 06 Jun 2011

103

5 <210> 254
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 10 tgtctccgc tttgtctctt 20

15 <210> 255
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 <212> DNA
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 20 <400> 255
 agacgtggag ttctctctga 20

25 <210> 256
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 <213> Artificial Sequence
 30 <220>
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 <400> 256
 35 caaacttcag ttcgccatca 20

40 <210> 257
 <211> 20
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 <213> Artificial Sequence
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 <223> a primer sequence for PCR
 45 <400> 257
 caactttccc acccaaagaa 20

50 <210> 258
 <211> 20
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 <213> Artificial Sequence
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 55 <223> a primer sequence for PCR
 <400> 258

2009274832 06 Jun 2011

104

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2009274832 06 Jun 2011

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2009274832 06 Jun 2011

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2009274832 06 Jun 2011

107

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2009274832 06 Jun 2011

108

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2009274832 06 Jun 2011

109

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2009274832 06 Jun 2011

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2009274832 06 Jun 2011

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2009274832 06 Jun 2011

112

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2009274832 06 Jun 2011

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2009274832 06 Jun 2011

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2009274832 06 Jun 2011

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2009274832 06 Jun 2011

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2009274832 06 Jun 2011

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2009274832 06 Jun 2011

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2009274832 06 Jun 2011

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2009274832 06 Jun 2011

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2009274832 06 Jun 2011

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2009274832 06 Jun 2011

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2009274832 06 Jun 2011

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2009274832 06 Jun 2011

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2009274832 06 Jun 2011

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2009274832 06 Jun 2011

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2009274832 06 Jun 2011

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 45 cagatccatc agcccttggg gtctgcattg atattccac gttctttttc gcaaatggca 37680
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 50 ctcatatgca gttgtcgcaa aaggtagcgg tcgtgggtat cctactcttg ggaagtttgt 37800
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 cgtaggcaagt atctgccgga tcatcgact tcagaatatt accgacggga cagatacgac 37920
 55 gtgggctatc gcccagttct ttatttgggt gtccgtggaa ccatttggtg ggattatttg 37980

2009274832 06 Jun 2011

128

cgcatacctc ccaacatttg ggcctttctt tcggcaatgg cggatccatcg ctcggacgcg 38040
 ctcatacaact gatggcagta ccgatccaag ctctgagcta ccatctgaga caacgacctg 38100
 5 gctccgaaga tcccgaacca aaaaacctgc caaggactca atattcagta tcaatgattt 38160
 ttgctgtgtc gatgaggtcc aactaatgaa cgatatcaat gccactcggg cgctggggga 38220
 cgaggctgcg agtgaccatc aggacgtgga gggaggctgt atcacagtcc aaaaagatgt 38280
 10 ggaagtgaca tgggccaagt acaagtcagg aaaaaaaaaat gatctggcct tcaagtatca 38340
 taaaggggct tgatcagctt tgcaaataatt tcgacttgac acggactata ttgctgtttt 38400
 15 gtgtatatatt aataaaaaata gacgccactg gcaatttgta attgataaag gtaagtctta 38460
 ttccgtaatc cataccccgt actctataca aagtactctg tgctccgtac ggagtacacg 38520
 gaaacaaacg gggatatagt cgtggcacct ttcccggtgtt ggcggacttg cccgtaacgt 38580
 20 aaacactccg cagatccctt ccaacacagt acataatcct gcagcgaaga gcgatctgat 38640
 agacgctatg tgccgtcgtg acttggttatg ccaattaacg gtggcagaat tgtggagcaa 38700
 25 tctagcagag gaaagtttcg atgtgcatgc cgagccctaa aaagtcccag tgcggagaat 38760
 gtagtaatcg actggacatt ccatgtactt tgcacgctat aacatatttc tatgccatat 38820
 acccctctgg taatcatgta gatcctcttg ctactgcgt tggctccttt gtatcgtact 38880
 30 ttccgcgtcg cagcattata agaggataga gagaccgcat gagagaatac acaagagaaa 38940
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 35 attgcaaa 39008

<210> 267
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 40 <212> PRT
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 <400> 267

45 Met Glu His Glu Thr Asp Leu Val Ser Phe Ala Phe Ser Gly Pro Ala
 1 5 10 15

50 Phe Asp Gln Ser Lys Pro Ile Tyr Ile Asp Ala Arg Asn Pro Ser Arg
 20 25 30

55 Ala Phe Asn Ala Ile Gln Phe Arg Arg Leu Val Arg Ser Leu Ile Ala
 35 40 45

Gly Leu Lys Ala Arg Gly Val Glu Arg Gly Asp Cys Val Leu Val Gln

2009274832 06 Jun 2011

129

	50		55		60
5	Leu 65	Glu Asn Ser Val	Leu 70	His Ser Ala	Leu 75 Phe Phe Ala Ile Val Gly 80
10	Ala Gly Gly Val Tyr 85	Met Gly Phe Asp 90	Val Ala Ser Arg Pro His 95	Glu	
15	Val Ala His 100	Leu Leu Arg Val Ala 105	Glu Pro Arg Leu Ile 110	Ile Thr Ala	
20	Pro Ser Ala 115	Leu Thr Arg Val 120	Leu Glu Val Cys Asn 125	Asn Gln Gly Met	
25	Ser Ser Asn Gln Val 130	Leu Leu Met Asp Glu Lys 135	Ser Ile Glu Ser Val 140		
30	Val Gln Phe Ala His 145	Gly Gln Ala Glu Gln Thr 150	Glu Asp Leu Asp Thr 155		
35	Gln Thr Val Asp 165	Gln Pro Ile Arg Leu Glu 170	Ser Leu Leu Gln Tyr Gly 175		
40	Glu Leu Asp Trp Leu Arg Phe 180	Glu Asp Ser Glu Glu Ser 185	Lys Ile Thr 190		
45	Pro Ala Ala Met Phe Leu Thr 195	Ser Gly Thr Ser Gly 200	Leu Pro Lys Ala 205		
50	Ala Ile Arg Thr His His 210	Thr Ile Ile Ser His 215	His Leu Ser Val Tyr 220		
55	Tyr Glu Val Pro Tyr 225	Pro Val Val Arg Leu Met 230	Ala Leu Pro Leu Tyr 235		
	His Ser Phe Gly 245	Asp Phe Trp Gly Asn 250	Ile Phe Pro Ile Arg Tyr Gly 255		
	Gln Pro Leu Tyr Ile Ile 260	Pro Arg Phe Glu Ile Thr 265	Ala Leu Leu Asp 270		
	Gly Ile Arg Gln His His 275	Ile Thr Glu Thr Tyr Met 280	Val Pro Ala Met 285		

2009274832 06 Jun 2011

130

5 Ile His Ile Leu Asn Arg Ser Ser Leu Asn Val Ala Glu Ser Leu Ser
 290 295 300

10 Ser Leu Arg Tyr Ile Gly Ile Ser Gly Ala Pro Ile Asp Gly Tyr Ser
 305 310 315 320

15 Met Gln Gln Phe Gln Ser Leu Leu Ser Pro Asp Ala Ile Ala Gly Asn
 325 330 335

20 Leu Trp Gly Met Ser Glu Val Gly Val Val Phe Gln Asn Arg Tyr Gly
 340 345 350

25 Ile Gln Pro Gln Phe Gly Ser Val Gly Thr Leu Leu Pro Arg Tyr Glu
 355 360 365

30 Leu Arg Phe Val Asn Pro Asp Thr Gly Glu Asp Val Ala Gly Thr Pro
 370 375 380

35 Asp Ser Pro Gly Glu Leu Tyr Val Arg Gly Pro Gly Leu Leu Leu Ala
 385 390 395 400

40 Tyr Lys Gly Arg Thr Asp Ala Lys Asp Glu Gln Gly Trp Phe Arg Thr
 405 410 415

45 Gly Asp Met Phe His Val Glu Asp Gly Asn Tyr His Val Ile Gly Arg
 420 425 430

50 Thr Lys Asp Leu Ile Lys Val Arg Gly Gln Val Thr Gln Tyr Ser Val
 435 440 445

55 Ala Pro Ala Glu Ile Glu Gly Ile Leu Arg Lys Asp Pro Ser Ile Lys
 450 455 460

60 Asp Ala Ala Val Ile Gly Val Met Leu Pro Asp Gly Ser Ser Glu Val
 465 470 475 480

65 Pro Arg Ala Tyr Val Val Arg Asn Asp Thr Ser Pro Glu Thr Thr Ala
 485 490 495

70 Asp Gln Val Ala Gly Leu Ile Gln Ser Gln Leu Ala Ser Tyr Lys Ala
 500 505 510

2009274832 06 Jun 2011

131

Leu Asp Gly Gly Val Val Phe Val Asp Asp Ile Pro Arg Ile Gly Ile
 515 520 525

5 Gly Lys His His Arg Ala Lys Leu Ser Gln Leu Asp His Gln Arg Glu
 530 535 540

10 Thr Ile Ala Ser Ile Leu Ala Glu Pro Val Ala Val
 545 550 555

<210> 268
 <211> 2447
 <212> PRT
 <213> Penicillium coprobium PF1169

<400> 268

20 Met Lys Ala Thr Glu Pro Val Ala Ile Ile Gly Thr Gly Cys Arg Phe
 1 5 10 15

25 Pro Gly Gly Ala Ser Ser Pro Ser Lys Leu Trp Glu Leu Leu Gln Ser
 20 25 30

30 Pro Arg Asp Ile Ala Arg Lys Val Pro Ala Asp Arg Phe Asn Ile Asp
 35 40 45

35 Ala Phe Tyr His Pro Asp Gly Asp His His Gly Thr Thr Asn Val Lys
 50 55 60

Glu Ser Tyr Phe Leu Asp Glu Asp Ile Lys Ala Phe Asp Ala Ala Phe
 65 70 75 80

40 Phe Asn Ile Ser Pro Thr Glu Ala Val Ala Met Asp Pro Gln Gln Arg
 85 90 95

45 Leu Leu Leu Glu Thr Val Tyr Glu Ser Leu Asp Ala Ala Gly Leu Arg
 100 105 110

50 Met Asp Ala Leu Gln Arg Ser Lys Thr Gly Val Phe Cys Gly Thr Leu
 115 120 125

Arg Asn Asp Tyr Asn Gln Ile Gln Ala Met Asp Pro Gln Ala Phe Pro
 130 135 140

55 Ala Tyr Val Val Thr Gly Asn Ser Pro Ser Ile Met Ala Asn Arg Ile

2009274832 06 Jun 2011

132

	145		150		155		160
5	Ser Tyr Tyr Phe Asp Trp Gln Gly Pro Ser Met Ala Val Asp Thr Gly	165	170	175			
10	Cys Ser Ser Ser Leu Leu Ala Val His Leu Gly Val Glu Ala Leu Gln	180	185	190			
15	Asn Asp Asp Cys Ser Met Ala Val Ala Val Gly Ser Asn Leu Ile Leu	195	200	205			
20	Ser Pro Asn Ala Tyr Ile Ala Asp Ser Lys Thr Arg Met Leu Ser Pro	210	215	220			
25	Thr Gly Arg Ser Arg Met Trp Asp Ser Lys Ala Asp Gly Tyr Gly Arg	225	230	235			
30	Gly Glu Gly Val Ala Ser Val Val Leu Lys Arg Leu Gln Asp Ala Ile	245	250	255			
35	Asn Asp Gly Asp Pro Ile Glu Cys Val Ile Arg Ala Ser Gly Ala Asn	260	265	270			
40	Ser Asp Gly Arg Thr Met Gly Ile Thr Met Pro Asn Pro Lys Ala Gln	275	280	285			
45	Gln Ser Leu Ile Leu Ala Thr Tyr Ala Arg Ala Gly Leu Ser Pro Gln	290	295	300			
50	Asn Asn Pro Glu Asp Arg Cys Gln Tyr Phe Glu Ala His Gly Thr Gly	305	310	315			
55	Thr Gln Ala Gly Asp Pro Gln Glu Ala Ala Ala Ile Asn Ser Ser Phe	325	330	335			
	Phe Gly Pro Glu Ser Val Pro Asp Ser Thr Asp Arg Leu Tyr Val Gly	340	345	350			
	Ser Ile Lys Thr Ile Ile Gly His Thr Glu Ala Thr Ala Gly Leu Ala	355	360	365			
	Gly Leu Ile Lys Ala Ser Leu Ser Leu Gln His Gly Met Ile Ala Pro	370	375	380			

2009274832 06 Jun 2011

133

5 Asn Leu Leu Met Gln His Leu Asn Pro Lys Ile Lys Pro Phe Ala Ala
 385 390 395 400

Lys Leu Ser Val Pro Thr Glu Cys Ile Pro Trp Pro Ala Val Pro Asp
 405 410 415

10 Gly Cys Pro Arg Arg Ala Ser Val Asn Ser Phe Gly Phe Gly Gly Ala
 420 425 430

15 Asn Val His Val Val Leu Glu Ser Tyr Thr Arg Ser Glu Leu Ser Pro
 435 440 445

20 Ser Asn Asn Ile Pro Ser Ser Leu Pro Phe Val Phe Ser Ala Ala Ser
 450 455 460

25 Glu Arg Thr Leu Thr Cys Val Met Glu Ser Tyr Ala Thr Phe Leu Gln
 465 470 475 480

Glu His Ala Thr Val Ser Leu Val Gly Leu Ala Leu Ser Leu Trp Asp
 485 490 495

30 Arg Arg Ser Thr His Arg His Arg Leu Thr Leu Met Ala His Ser Ile
 500 505 510

35 Gln Glu Leu Lys Asp Gln Ile Asn Thr Glu Ile Ser Arg Arg Val Thr
 515 520 525

40 Gly Lys Pro Ala Ser Val Val Ser Arg Ser Asn Thr Arg Pro Arg Arg
 530 535 540

45 Val Met Gly Ile Phe Thr Gly Gln Gly Val Gln Trp Pro Gln Met Gly
 545 550 555 560

Leu Asp Leu Ile Glu Ala Ser Pro Ser Ile Arg Lys Trp Ile Met Asn
 565 570 575

50 Leu Glu Glu Ala Leu Asp Glu Leu Pro Leu Asp Leu Arg Pro Gln Phe
 580 585 590

55 Ser Leu Leu Asp Glu Leu Ser Gln Pro Ala Ser Ser Ser Arg Val Asn
 595 600 605

2009274832 06 Jun 2011

134

Glu Gly Leu Leu Ser Leu Pro Leu Arg Thr Ala Leu Gln Ile Met Gln
 610 615 620

5

Val Asn Met Leu Arg Ala Val Gly Ile Glu Leu Thr Ile Val Val Gly
 625 630 635 640

10

His Ser Ser Gly Glu Ile Val Ala Ala Tyr Ala Ala Gly Val Leu Thr
 645 650 655

15

Ala Ser Asp Ala Ile Arg Ile Ala Tyr Leu Arg Gly Met Thr Ile Asp
 660 665 670

20

Lys Ser Arg Asp Pro Thr Gly Arg Met Met Ala Val Asn Leu Thr Trp
 675 680 685

25

Gln Gln Ala Gln Asn Ile Cys Ala Leu Glu Ala Tyr Ser Gly Arg Ile
 690 695 700

30

Ser Val Ala Ala Ala Asn Ser Pro Ser Ser Val Thr Leu Ser Gly Asp
 705 710 715 720

35

Ala Glu Cys Leu Arg Glu Leu Glu Trp Leu Leu Lys Ser Leu Gly Leu
 725 730 735

40

Thr Pro Arg Met Leu Arg Val Asp Thr Ala Tyr His Ser Pro His Met
 740 745 750

45

Lys Pro Cys Ala Asp Pro Tyr Arg Asp Ala Met Lys Ala Tyr Pro Val
 755 760 765

50

Ala Leu Ser Ala Ser Ala Ser Arg Trp Tyr Ser Ser Val Tyr Pro Gly
 770 775 780

55

Glu Val Met Thr Gly Tyr Asp Gln Gln Glu Leu Thr Gly Glu Tyr Trp
 785 790 795 800

Val Glu Asn Met Leu Arg Pro Val Gln Phe Ser Gln Ala Leu Glu Ala
 805 810 815

Ala Ala Arg Asp Ala Gly Pro Pro Asp Leu Ile Ile Glu Ile Gly Pro
 820 825 830

2009274832 06 Jun 2011

135

	His	Pro	Thr	Leu	Arg	Gly	Pro	Val	Leu	Gln	Thr	Leu	Ser	Lys	Met	His
				835				840					845			
5	Ser	Ala	His	Ser	Ala	Ile	Pro	Tyr	Leu	Ala	Leu	Ala	Glu	Arg	Gly	Lys
		850					855					860				
10	Pro	Gly	Leu	Asp	Thr	Trp	Ala	Thr	Ala	Leu	Gly	Ser	Ser	Trp	Ala	His
	865					870					875					880
15	Leu	Gly	Pro	Asn	Val	Val	Arg	Leu	Thr	Asp	Tyr	Val	Ser	Leu	Phe	Asp
					885					890					895	
20	Pro	Asn	His	Trp	Pro	Val	Leu	Val	Glu	Ser	Leu	Pro	Phe	Tyr	Pro	Phe
				900					905						910	
25	Asp	His	Thr	Gln	Thr	Tyr	Trp	Thr	Gln	Ser	Arg	Met	Ser	Ser	Asn	His
		915						920					925			
30	Asn	His	Arg	Ala	Thr	Ser	Pro	Asn	Ala	Leu	Leu	Gly	Ser	Leu	Ser	Pro
		930					935					940				
35	Glu	Thr	Gly	Ala	Glu	Lys	Phe	Arg	Trp	Arg	Asn	Tyr	Leu	Arg	Pro	Glu
	945					950					955					960
40	Glu	Leu	Pro	Trp	Leu	Ala	Asp	Arg	Arg	Ala	Asp	Ser	Gly	Ser	Val	Phe
					965					970					975	
45	Pro	Glu	Thr	Gly	Tyr	Ile	Ser	Met	Ala	Leu	Glu	Ala	Gly	Met	Ile	Met
				980					985					990		
50	Ala	Gln	Thr	Gln	Gly	Leu	Arg	Leu	Leu	Asn	Val	Lys	Asp	Leu	Thr	Ile
		995						1000					1005			
55	His	Thr	Gln	Leu	Pro	Ile	Gln	Asn	Asp	Pro	Ile	Gly	Thr	Glu	Val	
		1010					1015					1020				
60	Leu	Val	Thr	Val	Gly	Ser	Ile	His	Ser	His	Asp	Gly	Ala	Ile	Thr	
		1025					1030					1035				
65	Ala	Trp	Phe	Cys	Cys	Glu	Ala	Val	Val	Ser	Gly	Glu	Leu	Val	Gln	
		1040					1045					1050				
70	Cys	Ala	Thr	Ala	Lys	Met	Ile	Met	His	Pro	Gly	Asp	Ser	Asp	Arg	

2009274832 06 Jun 2011

136

	1055		1060		1065
5	Ala Leu 1070	Leu Pro Pro	Gln Gly 1075	Gln Leu Pro	Gln Ala 1080
10	Val Asp 1085	Ser Thr Glu Phe	Tyr 1090	Asp Ser Leu Arg	Arg Ala 1095
15	His Cys 1100	Thr Gly Pro Phe	Ser 1105	Thr Leu Thr Gly	Leu Arg 1110
20	Arg Asp 1115	Leu Ala Thr Gly	Ser 1120	Val Pro Val Pro	Ser Asn 1125
25	Asp Glu 1130	Pro Met Ala Leu	His 1135	Pro Ala Ile Leu	Asp Leu 1140
30	Gln Thr 1145	Met Ile Ala Ala	Ile 1150	Gly Gly Leu Glu	Glu Thr 1155
35	Thr Gly 1160	Pro Phe Leu Ser	Arg 1165	Asn Val Asp Ser	Thr Trp 1170
40	Pro Val 1175	Leu Cys Ala Ser	Asp 1180	Trp Gln Gly Lys	Glu Leu 1185
45	Ala Ser 1190	Tyr Leu Thr Cys	Val 1195	Asn Gly Asp Arg	Ile Arg 1200
50	Ile Asp 1205	Ile Phe Thr Met	Asn 1210	Gly Glu Lys Ala	Val Gln 1215
55	Gly Val 1220	Ser Leu Ile Cys	Gln 1225	Pro Ser Gly Thr	Ala Pro 1230
	Leu Gln 1235	Val Leu Ser Gln	Thr 1240	Ala Trp Gly Pro	Leu Glu 1245
	Leu Lys 1250	Lys Gly Ser Arg	Lys 1255	Leu Pro Ala Thr	Met Leu 1260
	His Ser 1265	Leu Arg Glu Glu	Leu 1270	Ala Leu Leu Tyr	Leu Lys 1275

2009274832 06 Jun 2011

137

5	Arg	Asn	Gly	Leu	Thr	Asp	Leu	Glu	Arg	Ser	Gly	Leu	Asp	Phe	Asp
	1280						1285					1290			
10	Gly	Ala	Arg	Leu	Leu	Ala	Trp	Met	Asn	Gln	Cys	Ile	Ala	Asn	Ala
	1295						1300					1305			
15	Ser	Gln	Glu	Pro	Asp	Pro	Val	Gly	Glu	Ser	Glu	Cys	Leu	Asp	Gln
	1310						1315					1320			
20	Lys	Ile	Glu	Asp	Phe	Thr	Ala	Gly	Val	Ser	Pro	Ser	Leu	Leu	Asn
	1325						1330					1335			
25	Asp	Pro	Gly	Leu	Thr	Ala	Ile	Ala	Ala	Val	Gly	Gln	Arg	Leu	Pro
	1340						1345					1350			
30	Arg	Val	Leu	Arg	Asp	Ser	Gly	Leu	Gln	Ile	Glu	Ala	Trp	Pro	Ala
	1355						1360					1365			
35	Ile	Asp	Glu	Glu	Ser	Gln	Tyr	Leu	Lys	Glu	Asp	Leu	Gln	Val	Leu
	1370						1375					1380			
40	Asp	Leu	Glu	Asp	Glu	Leu	Val	Ser	Val	Val	Ser	Gln	Ala	Cys	Phe
	1385						1390					1395			
45	Arg	Phe	Pro	Gln	Met	Asn	Ile	Leu	Gln	Ile	Gly	Gln	Phe	Gly	Gly
	1400						1405					1410			
50	His	Val	His	Ser	Gly	Leu	Lys	Lys	Met	Gly	Arg	Thr	Tyr	Arg	Ser
	1415						1420					1425			
55	Phe	Thr	Tyr	Ala	Gly	Leu	Ser	Val	Ser	Gly	Leu	Gln	Ala	Ile	Glu
	1430						1435					1440			
	Glu	Asp	Leu	Glu	Gln	Pro	Gly	Glu	Val	Ser	His	Lys	Thr	Leu	Asp
	1445						1450					1455			
	Ile	Asn	Glu	Asp	Pro	Val	Glu	Gln	Gly	Cys	Arg	Glu	Gln	Phe	Tyr
	1460						1465					1470			
	Asp	Met	Val	Leu	Ile	Thr	Ala	Ala	Val	Phe	Leu	Gln	Glu	Val	Ala
	1475						1480					1485			

Val Ala His Val Arg Arg Leu Leu Lys Pro Gly Gly Phe Leu Val
 1490 1495 1500
 5
 Leu Leu Val Arg Thr Asn Pro Ser Thr Thr Tyr Leu Asn Leu Leu
 1505 1510 1515
 10 Phe Gly Pro Pro Met Arg Cys Thr Glu Thr Gly Lys Gly Tyr Cys
 1520 1525 1530
 15 Ser Gly Glu Pro Ile Thr Thr Arg Arg Asp Trp Val Glu Leu Leu
 1535 1540 1545
 20 Ser Asn Gly Gly Phe Tyr Gly Leu Asp Ser Phe Asp Ala Ser Gln
 1550 1555 1560
 Glu Ser Glu Ser Leu Gly Asp Phe Ser Leu Leu Leu Cys Arg Thr
 1565 1570 1575
 25 Pro Asp Ser Pro Ala Glu Pro Gln Ser Arg Gly Asp Leu Leu Leu
 1580 1585 1590
 30 Leu Gly Gly Asp Ala Glu Glu Ala Asp Cys Leu Thr Ser Glu Leu
 1595 1600 1605
 35 Phe Glu Leu Val Gln Asp Asp Phe Val Lys Val Ala His Ala Pro
 1610 1615 1620
 40 Asp Leu Asp Leu Ile Glu Asp Arg Asp Leu Ser Lys Leu Thr Val
 1625 1630 1635
 Leu Tyr Leu Val Asp Asp Arg Asp Leu Thr Asn Ala Thr Leu Ser
 1640 1645 1650
 45 Glu Leu Cys Arg Leu Met Thr Val Ser Lys Arg Met Leu Val Val
 1655 1660 1665
 50 Thr Cys Glu Lys Val Asp His Pro Asp Ala Gly Leu Val Lys Gly
 1670 1675 1680
 55 Leu Leu Ser Thr Phe Leu Ala Ser Glu Arg Ser Ser Ser Leu Leu
 1685 1690 1695

2009274832 06 Jun 2011

139

Gln Leu Leu His Ile Thr Asp Pro Val Gly Val Thr Thr Glu Ile
 1700 1705 1710

5 Leu Ala Thr Ala Leu Gly His Phe Val Gln Ala Ser Ala Ala Gln
 1715 1720 1725

10 Glu Asn Pro His Ser Cys Gly Leu Thr Asn Ile Glu Pro Glu Ile
 1730 1735 1740

15 Gln Tyr Asp Gly Ser Met Phe Arg Val Pro Arg Gln Tyr His Asp
 1745 1750 1755

20 His Ala Thr Gly Leu Arg His Leu Ala Arg Arg Gln Lys Val Thr
 1760 1765 1770

Asp Cys Val Asp Leu Asp Lys Gly Val Val Gln Ile Leu Pro Ala
 1775 1780 1785

25 Thr Thr Asp Lys Thr Cys Glu Gly Phe Arg Leu Leu Ser Met Ala
 1790 1795 1800

30 Asp Pro Pro Ile Thr Ala Ser Tyr Gly Pro Thr Leu His Leu Arg
 1805 1810 1815

35 Val Arg His Ser Ser Ile Ala Ala Val Arg Val Ala Gly Ala Ile
 1820 1825 1830

Phe Leu Arg Leu Val Ile Gly Leu Asp Val Lys Ser Asn Lys Arg
 1835 1840 1845

40 Met Ile Ala Leu Ser Ser His Ile Ala Ser His Val Ile Val Pro
 1850 1855 1860

45 Asp Ser Trp Ala Trp Ser Val Pro Asp Thr Val Leu Glu Ala His
 1865 1870 1875

50 Glu Gln Ser Tyr Leu Arg Ala Thr Ala Ala Ala Leu Leu Ala Gly
 1880 1885 1890

55 Tyr Leu Val Glu Gln Val Pro Gln Ser Gly Thr Leu Val Val His
 1895 1900 1905

Glu Ala Asp Gly Val Leu Gln Ser Val Phe His Gln Met Leu Thr

2009274832 06 Jun 2011

	1910						1915						1920					
5	Arg	Arg	Asp	Gly	Lys	Val	Ile	Phe	Ser	Thr	Ser	Lys	Ser	Asn	Pro			
	1925						1930					1935						
10	Asp	Lys	Glu	Arg	Pro	Met	Leu	Leu	Leu	His	Glu	His	Ser	Thr	Ala			
	1940						1945					1950						
15	Arg	Gln	Leu	Ser	Gln	Val	Leu	Pro	Ser	Asp	Val	Ser	Ala	Ile	Ala			
	1955						1960					1965						
20	Ile	Leu	His	Arg	Arg	Gly	Gln	Gly	Val	Tyr	Asp	Arg	Met	Leu	Ser			
	1970						1975					1980						
25	Leu	Leu	Pro	Asp	Asn	Ala	Thr	Arg	Ile	His	Leu	Gln	Asp	Phe	Tyr			
	1985						1990					1995						
30	Leu	Thr	Ser	Ala	Ser	Thr	Gly	Pro	Ile	Asn	Ala	Asp	Asp	Ser	Ser			
	2000						2005					2010						
35	Leu	Ile	Ala	Lys	Ala	Phe	Leu	Thr	Ala	Cys	Leu	Val	Ala	Tyr	Thr			
	2015						2020					2025						
40	Gly	Arg	Glu	Gly	Leu	Pro	Pro	Asn	Ser	Val	Asp	Ser	Leu	Pro	Ile			
	2030						2035					2040						
45	Ser	Arg	Ile	Ser	Glu	Tyr	Pro	Ile	Leu	Asp	Ser	Gln	Asp	Ala	Val			
	2045						2050					2055						
50	Val	Asp	Trp	Asp	Ser	Thr	Thr	Pro	Val	Leu	Ala	Gln	Ile	Pro	Thr			
	2060						2065					2070						
55	Ala	Gly	Ser	Gln	Val	Gln	Leu	Ser	Glu	Lys	Lys	Thr	Tyr	Ile	Leu			
	2075						2080					2085						
60	Val	Gly	Leu	Gly	Ser	Glu	Leu	Ala	His	Ala	Ile	Cys	Leu	Trp	Leu			
	2090						2095					2100						
65	Ala	Thr	His	Gly	Ala	Lys	Trp	Ile	Leu	Leu	Ala	Gly	Ser	Arg	Leu			
	2105						2110					2115						
70	Asp	Ser	Asp	Ala	Trp	Trp	Leu	Glu	Glu	Val	Ser	Arg	Arg	Gly	Thr			
	2120						2125					2130						

2009274832 06 Jun 2011

141

5	Arg 2135	Ile	Ala	Val	Ser	Lys	Ile 2140	Asn	Leu	Ile	Asp	Gly 2145	Ile	Ser	Ala
10	Thr 2150	Ser	Leu	His	Gln	Thr	Ile 2155	Pro	Tyr	Ala	Phe	Pro 2160	Pro	Val	Val
15	Gly 2165	Gly	Val	Leu	Ile	Gln	Pro 2170	Pro	Pro	Leu	Pro	Asp 2175	Cys	Ser	Leu
20	Ser 2180	Gln	Leu	Thr	Ile	Asp	Ser 2185	Leu	Arg	Asn	His	Leu 2190	His	Pro	Val
25	Leu 2195	Lys	Gly	Leu	Gln	Gln	Leu 2200	Asp	Glu	Leu	Tyr	Lys 2205	Thr	Pro	Thr
30	Leu 2210	Asp	Phe	Trp	Val	Leu	Ile 2215	Gly	Ser	Ile	Ala	Gly 2220	Val	Leu	Gly
35	His 2225	Ala	Asp	Gln	Ala	Met	Thr 2230	Ala	Ala	Met	Ser	Glu 2235	Lys	Met	Ala
40	Leu 2240	Leu	Val	Arg	His	Arg	Arg 2245	Ala	Gln	Gly	Arg	Pro 2250	Ala	Ser	Leu
45	Val 2255	His	Leu	Gly	Glu	Ile	His 2260	Gly	Ile	Ser	Ser	Pro 2265	Ser	Pro	Ser
50	Gln 2270	Pro	Leu	Trp	Cys	Gly	Pro 2275	Val	Ala	Val	Ser	Gln 2280	Arg	Asp	Val
55	Asp 2285	Glu	Ile	Leu	Ala	Glu	Ala 2290	Ile	Leu	Cys	Gly	Arg 2295	Ser	Asp	Ser
	Asn 2300	Ser	Asn	Ala	Glu	Leu	Ile 2305	Gly	Gly	Leu	Arg	His 2310	Gln	Ser	Leu
	Lys 2315	Cys	Gly	Tyr	Gly	Glu	Cys 2320	Pro	Ile	Pro	Lys	Leu 2325	Trp	Pro	Phe
	Tyr 2330	Ser	Tyr	Thr	Ala	Thr	Ala 2335	Ser	Gln	Asp	Gln	Ile 2340	Leu	Ala	Leu

2009274832 06 Jun 2011

142

Ile Glu Thr Arg Ser Thr Lys Asp Leu Val Thr Ala Ala Thr Ser
 2345 2350 2355

5

Leu Glu Glu Lys Ala Glu Ala Val Val Arg Pro Leu Met Glu Lys
 2360 2365 2370

10

Ile Arg Ala Ser Leu Asn Leu Ala Glu Asp Ala Pro Leu Ser Ala
 2375 2380 2385

15

Asp Thr Leu Ile Pro Glu Leu Gly Ile Asp Ser Leu Ile Ala Ile
 2390 2395 2400

20

Gly Leu Ser Gln Trp Phe Thr Lys Glu Leu Ser Val Asp Ile Gly
 2405 2410 2415

Val Ile Leu Ile Leu Ser Gly Val Ser Val Gly Glu Leu Ala His
 2420 2425 2430

25

Ala Ala Ala Ser Lys Leu Cys Asn Val Ser Val Gly Lys Pro
 2435 2440 2445

30

<210> 269
 <211> 509
 <212> PRT
 <213> Penicillium coprobium PF1169

35

<400> 269

Met Asp Asn Met Asp Asn Met Asn Asn Thr Pro Leu Gly Phe Asn Trp
 1 5 10 15

40

Ala Trp Ala Val Ile Ile Ser Phe Leu Gly Leu Leu Thr Phe Ser Phe
 20 25 30

45

Val Ser Pro His Leu Phe Pro Ser Arg Leu Thr Val Ile Asn Gly Gly
 35 40 45

50

Arg Ala Trp Asp Ile Phe Arg Thr Lys Ala Lys Lys Arg Phe Arg Ser
 50 55 60

55

Asp Ala Ala Arg Leu Ile Lys Asn Gly Phe Glu Glu Ser Pro Asp Ala
 65 70 75 80

Phe Arg Ile Ile Thr Asp Asn Gly Pro Leu Leu Val Leu Ser Pro Gln

2009274832 06 Jun 2011

143

85

90

95

5 Tyr Ala Arg Glu Val Arg Ser Asp Asp Arg Leu Ser Leu Asp His Phe
100 105 110

10 Ile Ala Ser Glu Phe His Pro Asn Ile Pro Gly Phe Glu Pro Phe Lys
115 120 125

15 Leu Ile Leu Asp Pro Lys Asn Pro Leu Asn Thr Ile Leu Lys Ser Asn
130 135 140

20 Leu Thr Gln Ala Leu Glu Asp Leu Ser Ala Glu Val Thr Glu Ala Leu
145 150 155 160

25 Ser Ala Thr Cys Thr Asp Asp Pro Glu Trp His Glu Val Ser Val Ser
165 170 175

30 Gln Thr Ala Leu Lys Ile Ile Ala Gln Met Ala Ser Lys Ala Phe Ile
180 185 190

35 Gly Gln Glu Arg Cys Arg Asp Ala Lys Trp His Asn Ile Ile Ile Thr
195 200 205

40 Tyr Thr His Asn Val Tyr Gly Ala Ala Gln Ala Leu His Phe Trp Pro
210 215 220

45 Ser Phe Leu Arg Pro Ile Val Ala Gln Phe Leu Pro Ala Cys Arg Thr
225 230 235 240

50 Leu Gln Ala Gln Ile Ala Glu Ala Arg Glu Ile Leu Glu Pro Leu Val
245 250 255

55 Ala Gln Arg Arg Ala Glu Arg Ala Thr Arg Ala Ala Gln Glu Lys Pro
260 265 270

His Pro Ser Gly Gly Asp Ile Ile Asp Trp Leu Glu Gln Phe Tyr Gly
275 280 285

Asp Gln Pro Tyr Asp Pro Val Ala Ala Gln Leu Leu Leu Ser Phe Ala
290 295 300

Ala Ile His Gly Thr Ser Asn Leu Leu Ala Gln Ala Leu Ile Asp Leu
305 310 315 320

144

5 Cys Gly Gln Pro Glu Leu Val Gln Asp Leu Arg Glu Glu Ala Val Ser
 325 330 335
 10 Val Leu Gly Lys Glu Gly Trp Thr Arg Ala Ala Leu Tyr Gln Leu Lys
 340 345 350
 15 Leu Met Asp Ser Ala Leu Lys Glu Ser Gln Arg Leu Ala Pro Asn Arg
 355 360 365
 20 Gly Leu Arg Ile His Arg Gly Thr Thr Leu Met Val Ser Ala His Asn
 385 390 395 400
 25 Met Trp Asp Pro Glu Ile Tyr Pro Asp Pro Arg Lys Tyr Asp Gly Tyr
 405 410 415
 30 Arg Phe His Lys Leu Arg Gln Thr Ser Gly Gln Glu Gly Gln His Gln
 420 425 430
 35 Leu Val Ser Ser Thr Pro Asp His Met Gly Phe Gly Tyr Gly Lys His
 435 440 445
 40 Ala Cys Pro Gly Arg Phe Phe Ala Ala Ala Gln Ile Lys Val Ala Leu
 450 455 460
 45 Cys Asn Ile Leu Leu Lys Tyr Asp Ile Glu Tyr Arg Gly Gly Lys Ser
 465 470 475 480
 50 Pro Gly Val Trp Gly Gln Gly Ile His Leu Phe Pro Asp Pro Thr Ser
 485 490 495
 55 Arg Ile His Val Arg Arg Arg Lys Glu Glu Ile Asn Leu
 500 505
 <210> 270
 <211> 505
 <212> PRT
 <213> Penicillium coprobium PF1169
 <400> 270

2009274832 06 Jun 2011

2009274832 06 Jun 2011

145

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

2009274832 06 Jun 2011

146

	225		230		235		240									
5	Arg	Glu	Ile	Arg	Thr	Ala	Arg	Thr	Leu	Ile	Asp	Ser	Ala	Leu	Glu	Lys
				245					250						255	
10	Ser	Arg	Lys	Asn	Pro	Gln	Thr	Glu	Lys	Phe	Ser	Ser	Val	Ala	Trp	Val
				260					265					270		
15	Asp	Ala	Phe	Ala	Lys	Gly	Asn	Lys	Tyr	Asn	Ala	Ala	Met	Val	Gln	Leu
		275						280					285			
20	Arg	Leu	Ala	Asn	Ala	Ser	Ile	His	Ser	Ser	Ala	Asp	Leu	Leu	Val	Lys
	290						295					300				
25	Ile	Leu	Ile	Asn	Leu	Cys	Glu	Gln	Pro	Glu	Leu	Ile	Arg	Asp	Leu	Arg
	305					310					315					320
30	Asp	Glu	Ile	Ile	Ser	Val	Leu	Gly	Glu	Asn	Gly	Trp	Arg	Ser	Ser	Thr
					325					330					335	
35	Leu	Asn	Gln	Leu	Lys	Leu	Leu	Asp	Ser	Val	Leu	Lys	Glu	Ser	Gln	Arg
				340					345					350		
40	Leu	His	Pro	Val	Thr	Thr	Gly	Ala	Phe	Ser	Arg	Phe	Thr	Arg	Gln	Asp
			355					360					365			
45	Ile	Lys	Leu	Thr	Asn	Gly	Thr	Glu	Ile	Pro	Ser	Gly	Thr	Pro	Ile	Met
	370						375					380				
50	Val	Thr	Asn	Asp	Val	Ala	Gly	Asp	Ala	Ser	Ile	Tyr	Asp	Asp	Pro	Asp
	385					390					395					400
55	Val	Phe	Asp	Gly	Tyr	Arg	Tyr	Phe	Arg	Met	Arg	Glu	Gly	Ala	Asp	Lys
					405					410					415	
60	Ala	Arg	Ala	Pro	Phe	Thr	Thr	Thr	Gly	Gln	Asn	His	Leu	Gly	Phe	Gly
				420					425					430		
65	Tyr	Gly	Lys	Tyr	Ala	Cys	Pro	Gly	Arg	Phe	Phe	Ala	Ala	Thr	Glu	Ile
			435					440					445			
70	Lys	Ile	Ala	Leu	Cys	His	Met	Leu	Leu	Lys	Tyr	Glu	Trp	Arg	Leu	Val
	450						455					460				

147

5 Lys Asp Arg Pro His Gly Ile Val Thr Ser Gly Phe Ala Ala Phe Arg
 465 470 475 480
 10 Asp Pro Arg Ala Ser Ile Glu Val Arg Arg Arg Ala Val Ala Gly Glu
 485 490 495
 15 Glu Leu Glu Val Leu Thr Gly Lys Lys
 500 505
 20 <210> 271
 <211> 241
 <212> PRT
 <213> Penicillium coprobium PF1169
 25 Met Asp Gly Trp Ser Asp Ile Ser Ser Ala Pro Ala Gly Tyr Lys Asp
 1 5 10 15
 30 Val Val Trp Ile Ala Asp Arg Ala Leu Leu Ala Gln Gly Leu Gly Trp
 20 25 30
 35 Ser Ile Asn Tyr Leu Ala Met Ile Tyr Gln Ser Arg Lys Asp Arg Thr
 35 40 45
 40 Tyr Gly Met Ala Ile Leu Pro Leu Cys Cys Asn Phe Ala Trp Glu Phe
 50 55 60
 45 Val Tyr Thr Val Ile Tyr Pro Ser Gln Asn Pro Phe Glu Arg Ala Val
 65 70 75 80
 50 Leu Thr Thr Trp Met Val Leu Asn Leu Tyr Leu Met Tyr Thr Thr Ile
 85 90 95
 55 Lys Phe Ala Pro Asn Glu Trp Gln His Ala Pro Leu Val Gln Arg Ile
 100 105 110
 60 Leu Pro Val Ile Phe Pro Val Ala Ile Ala Ala Phe Thr Ala Gly His
 115 120 125
 65 Leu Ala Leu Ala Ala Thr Val Gly Val Ala Lys Ala Val Asn Trp Ser
 130 135 140

2009274832 06 Jun 2011

2009274832 06 Jun 2011

148

Ala Phe Leu Cys Phe Glu Leu Leu Thr Ala Gly Ala Val Cys Gln Leu
145 150 155 160

5 Met Ser Arg Gly Ser Ser Arg Gly Ala Ser Tyr Thr Ile Trp Val Ser
165 170 175

10 Arg Phe Leu Gly Ser Tyr Ile Gly Ser Ile Phe Met His Val Arg Glu
180 185 190

15 Thr His Trp Pro Gln Glu Phe Asp Trp Ile Ser Tyr Pro Phe Val Ala
195 200 205

20 Trp His Gly Ile Met Cys Phe Ser Leu Asp Ile Ser Tyr Val Gly Leu
210 215 220

25 Leu Trp Tyr Ile Arg Arg Gln Glu Arg Gln Gly Gln Leu Lys Lys Ala
225 230 235 240

30 <210> 272
<211> 464
<212> PRT
<213> Penicillium coprobium PF1169

35 <400> 272
Met Lys Val Ile Ile Val Gly Gly Ser Ile Ala Gly Leu Ala Leu Ala
1 5 10 15

40 His Cys Leu Asp Lys Ala Asn Ile Asp Tyr Val Ile Leu Glu Lys Lys
20 25 30

45 Lys Glu Ile Ala Pro Gln Glu Gly Ala Ser Ile Gly Ile Met Pro Asn
35 40 45

50 Gly Gly Arg Ile Leu Glu Gln Leu Gly Leu Tyr Asp Gln Ile Glu Glu
50 55 60

55 Leu Ile Glu Pro Leu Val Arg Ala His Val Thr Tyr Pro Asp Gly Phe
65 70 75 80

Asn Tyr Thr Ser Arg Tyr Pro Ala Leu Ile Gln Gln Arg Phe Gly Tyr
85 90 95

2009274832 06 Jun 2011

149

5 Pro Leu Ala Phe Leu Asp Arg Gln Lys Leu Leu Gln Ile Leu Ala Thr
 100 105 110

10 Gln Pro Val Gln Ser Ser Arg Val Lys Leu Asp His Lys Val Glu Ser
 115 120 125

15 Ile Glu Val Ser Pro Cys Gly Val Thr Val Ile Thr Ser Asn Gly His
 130 135 140

20 Thr Tyr Gln Gly Asp Leu Val Val Gly Ala Asp Gly Val His Ser Arg
 145 150 155 160

25 Val Arg Ala Glu Met Trp Arg Leu Ala Asp Ala Ser Gln Gly Asn Val
 165 170 175

30 Cys Gly Asn Gly Asp Lys Ala Phe Thr Ile Asn Tyr Ala Cys Ile Phe
 180 185 190

35 Gly Ile Ser Ser His Val Asp Gln Leu Asp Pro Gly Glu Gln Ile Thr
 195 200 205

40 Cys Tyr Asn Asp Gly Trp Ser Ile Leu Ser Val Ile Gly Gln Asn Gly
 210 215 220

45 Arg Ile Tyr Trp Phe Leu Phe Ile Lys Leu Glu Lys Glu Phe Val Tyr
 225 230 235 240

50 Asp Gly Ser His Lys Thr Gln Leu His Phe Ser Arg Glu Asp Ala Arg
 245 250 255

55 Ala His Cys Glu Arg Leu Ala Gln Glu Pro Leu Trp Lys Asp Val Thr
 260 265 270

60 Phe Gly Gln Val Trp Ala Arg Cys Glu Val Phe Gln Met Thr Pro Leu
 275 280 285

65 Glu Glu Gly Val Leu Gly Lys Trp His Trp Arg Asn Ile Ile Cys Ile
 290 295 300

70 Gly Asp Ser Met His Lys Phe Ala Pro His Ile Gly Gln Gly Ala Asn
 305 310 315 320

2009274832 06 Jun 2011

150

Cys Ala Ile Glu Asp Ala Ala Gln Leu Ser Asn Ser Leu His Thr Trp
 325 330 335

5
 Leu Ser Gly Ser Gly Lys Glu His Gln Leu Lys Thr Asp Asp Leu Thr
 340 345 350

10
 Glu Ile Leu Ala Gln Phe Ala Gln Thr Arg Leu Gln Arg Leu Gly Pro
 355 360 365

15
 Thr Ala Met Ala Ala Arg Ser Ala Met Arg Leu His Ala Arg Glu Gly
 370 375 380

20
 Leu Lys Asn Trp Ile Leu Gly Arg Tyr Phe Leu Pro Tyr Ala Gly Asp
 385 390 395 400

25
 Lys Pro Ala Asp Trp Ala Ser Arg Gly Ile Ala Gly Gly Asn Thr Leu
 405 410 415

30
 Asp Phe Val Glu Pro Pro Thr Arg Ala Gly Pro Gly Trp Ile Gln Phe
 420 425 430

35
 Ser Gln Ser Gly Lys Arg Thr Ser Phe Pro Met Ala Val Ala Gly Leu
 435 440 445

40
 Cys Leu Val Ser Ile Val Ala Arg Ile Met Tyr Leu Lys Leu Val Ala
 450 455 460

<210> 273
 <211> 317
 <212> PRT
 <213> *Penicillium coprobium* PF1169
 <400> 273

45
 Met Ala Gly Ser Gln Ser Thr Ala Gln Leu Ala Arg Leu Leu Ile Asp
 1 5 10 15

50
 Ile Ser Arg Phe Asp Lys Tyr Asn Cys Leu Phe Ala Ile Phe Pro Gly
 20 25 30

55
 Val Trp Ser Ile Phe Leu Ala Ala Ala Ser Arg His Ala Asp Gly Asp
 35 40 45

Pro Val Pro Leu Asp Phe Val Leu Gly Arg Ala Gly Leu Ala Phe Met

2009274832 06 Jun 2011

151

	50	55	60
5	Tyr Thr Tyr Met Leu Ser Gly Ala Gly Met Val Trp Asn Asp Trp Ile 65 70 75 80		
10	Asp Arg Asp Ile Asp Ala Gln Val Ala Arg Thr Lys Asn Arg Pro Leu 85 90 95		
15	Ala Ser Gly Arg Leu Ser Thr Arg Ala Ala Leu Ile Trp Met Leu Val 100 105 110		
20	Gln Tyr Ala Ala Ser Val Trp Leu Met Asp Arg Met Val Ser Gly Gln 115 120 125		
25	Asp Val Trp Thr Tyr Met Leu Pro Leu Thr Thr Gly Ile Ile Leu Tyr 130 135 140		
30	Pro Phe Gly Lys Arg Pro Thr Ser Arg Lys Leu Gly Val Tyr Pro Gln 145 150 155 160		
35	Tyr Ile Leu Gly Ala Ser Ser Ala Leu Thr Ile Leu Pro Ala Trp Ala 165 170 175		
40	Ser Val Tyr Thr Gly Arg Ile Ser Leu Lys Asp Leu Gly Met Arg Cys 180 185 190		
45	Leu Pro Leu Cys Leu Phe Leu Phe Leu Trp Thr Ile Tyr Phe Asn Thr 195 200 205		
50	Ala Tyr Ser Tyr Gln Asp Ile Lys Asp Asp Cys Lys Leu Asn Val Asn 210 215 220		
55	Ser Ser Tyr Val Leu Ala Gly Ser His Val Arg Gly Met Leu Leu Leu 225 230 235 240		
60	Gln Ala Ile Ala Val Val Leu Val Ile Pro Trp Ile Leu Tyr Thr Ser 245 250 255		
65	Ala Ser Thr Trp Leu Trp Val Ser Trp Leu Gly Val Trp Thr Ala Ser 260 265 270		
70	Leu Gly Glu Gln Leu Tyr Leu Phe Asp Val Lys Asp Pro Ser Ser Gly 275 280 285		

2009274832 06 Jun 2011

152

5 Gly Lys Val His Arg Arg Asn Phe Ala Leu Gly Ile Trp Asn Val Leu
 290 295 300

10 Ala Cys Phe Val Glu Leu Leu Tyr Ala Ser Gly Ser Leu
 305 310 315

<210> 274
 <211> 522
 <212> PRT
 <213> Penicillium coprobium PF1169

15 <400> 274

20 Met Ser Thr Gln Glu Val Cys Leu Pro Val Ser Gln Arg Asp Gln Val
 1 5 10 15

Lys Glu Gly Pro Val Arg Leu His Gly Leu Cys Glu Asp Gly Met Cys
 20 25 30

25 Asp Ala Arg Arg Thr Gly Asp Arg Ser Ala Tyr Pro Leu Ser Ser Leu
 35 40 45

30 Asp His Asn Pro Leu Gly Met Asn Val Thr Phe Leu Leu Phe Phe Gln
 50 55 60

35 Thr Thr Gln Pro Glu Lys Ser Ile Gly Val Leu Glu Asn Gly Ile Glu
 65 70 75 80

40 Leu Leu Leu Lys Val His Pro Phe Leu Ala Gly Asp Val Thr Arg Arg
 85 90 95

Thr Glu Ser Ser Gln Thr Lys Tyr Thr Trp Gln Ile Glu Pro Glu Ala
 100 105 110

45 Ser Glu Ser Leu Val Gln Phe Pro Ile Leu Arg Ile Arg His Tyr Gln
 115 120 125

50 Ala Glu Ser Phe Lys Glu Ile Gln Ser Lys Cys Leu Leu Thr Gly Thr
 130 135 140

55 Glu Glu Gln Glu Ile Ile Ser Arg Leu Ala Pro Leu Pro Ile Asp Met
 145 150 155 160

2009274832 06 Jun 2011

153

	Asp	Ile	Ser	Leu	Pro	Arg	Arg	Pro	Ile	Leu	Arg	Phe	Gln	Ala	Asn	Val	
					165					170					175		
5	Met	Arg	Asp	Gly	Ile	Ile	Leu	Ala	Met	Thr	Phe	His	His	Ser	Ala	Met	
				180					185					190			
10	Asp	Gly	Ala	Gly	Ala	Ala	Arg	Val	Leu	Gly	Leu	Leu	Ala	Asp	Cys	Cys	
			195					200					205				
15	Arg	Asp	Pro	Thr	Ala	Met	Ser	Ser	Ala	Ser	Val	Ser	Pro	Asp	Arg	Gln	
		210					215					220					
20	Leu	Arg	Ser	Glu	Ile	Glu	Arg	Leu	Val	Pro	Glu	Ser	Ser	Ser	Gly	Leu	
	225				230						235					240	
25	Ser	Arg	Met	Asp	Phe	Ser	Lys	His	Tyr	Cys	Gly	Leu	Gly	Asp	Trp	Ala	
					245					250					255		
30	Ala	Leu	Leu	Ala	Lys	Asn	Trp	Ser	Gly	Phe	Val	Arg	Ala	Arg	Ala	Thr	
				260					265					270			
35	Glu	Leu	Val	Thr	Trp	Arg	Leu	Lys	Ile	Pro	Gly	Pro	Lys	Ile	Glu	Tyr	
			275					280					285				
40	Leu	Lys	Glu	Ala	Cys	Asn	Thr	Leu	Ile	Lys	Gly	Gln	Thr	Ser	Phe	Gln	
	290					295						300					
45	Ala	Asp	Gly	Arg	Pro	Ser	Pro	Gly	Phe	Leu	Ser	Ser	Asn	Asp	Ile	Val	
	305					310					315					320	
50	Ser	Ala	Leu	Leu	Ala	Met	Ile	Leu	Arg	Gln	Ala	Gly	Gln	Leu	Ala	Gly	
					325					330					335		
55	Lys	Ser	Thr	Glu	Leu	Ser	Ile	Ala	Val	Asp	Met	Arg	Gly	Asn	Phe	Lys	
				340					345					350			
60	Thr	Pro	Ala	Phe	Asp	Asp	Tyr	Leu	Gly	Asn	Met	Val	Leu	Leu	Thr	Tyr	
		355						360					365				
65	Thr	Pro	Ile	Gln	Ala	Gly	Arg	Asn	Glu	Ala	Leu	Val	Asp	Gly	Thr	Asp	
		370					375					380					
70	Pro	Ser	Val	Glu	Leu	Arg	Gln	Glu	Cys	Leu	Glu	Asp	Leu	Thr	Gln	Ile	

2009274832 06 Jun 2011

154

	385		390		395		400									
5	Ala	Ala	Arg	Ile	Arg	Gln	Ser	Leu	Leu	Ala	Val	Asp	Ala	Glu	Tyr	Ile
					405					410					415	
10	Gln	Asp	Ala	Leu	Ser	His	Leu	His	Ser	Gln	Pro	Asp	Trp	Ala	Asp	Ile
				420					425					430		
15	Gly	Phe	Arg	Gly	Val	Pro	Ile	Pro	Leu	Ser	Ser	Phe	Arg	Asn	Phe	Glu
			435					440					445			
20	Ile	Phe	Gly	Leu	Asp	Phe	Gly	Glu	Ser	Leu	Gly	Ala	Gln	Pro	Arg	Gly
	450						455					460				
25	Phe	Gln	Leu	His	Leu	Pro	Val	Leu	Gly	Gly	Met	Cys	Phe	Ile	Leu	Pro
	465					470					475					480
30	Lys	Gly	Gln	Asp	Asp	Val	Ala	Ser	Thr	Glu	Pro	Trp	Asp	Leu	His	Leu
				485						490					495	
35	Thr	Leu	Asn	Arg	Asp	Asp	Gln	Leu	Leu	Leu	Ala	Lys	Asp	Pro	Leu	Phe
			500						505					510		
40	Cys	Trp	Ala	Ile	Gly	Ala	Gln	Ala	Lys	Glu						
		515					520									
45	<210>	275														
	<211>	434														
	<212>	PRT														
	<213>	Penicillium coprobium PF1169														
50	<400>	275														
55	Met	Asp	Ser	Leu	Leu	Thr	Ser	Pro	Leu	Trp	Leu	Lys	Ile	Ala	His	Glu
	1				5					10					15	
60	Leu	Ala	Leu	Tyr	Leu	Ser	Phe	Ile	Val	Pro	Thr	Ala	Phe	Leu	Ile	Ile
				20					25					30		
65	Thr	Thr	Gln	Lys	Ser	Ser	Ile	Ile	Arg	Trp	Ala	Trp	Thr	Pro	Cys	Leu
			35					40					45			
70	Leu	Tyr	Ile	Leu	Tyr	Gln	Phe	Ser	Leu	Arg	Val	Pro	Ser	Leu	Ser	Thr
	50						55					60				

2009274832 06 Jun 2011

155

	Ser	Gln	Phe	Leu	Lys	Gly	Val	Ala	Ala	Gly	Gln	Ala	Thr	Val	Ala	Ala	65	70	75	80
5	Leu	Gln	Cys	Leu	Asn	Leu	Leu	Leu	Ile	Thr	Lys	Leu	Asp	Gln	Thr	Asp	85	90	95	
10	Leu	Leu	Arg	Ala	Asn	Leu	Tyr	Ser	Pro	Ser	Ala	Gly	Leu	Leu	Ser	Arg	100	105	110	
15	Leu	Ala	Gln	Ser	Cys	Ala	Leu	Leu	Val	Asn	Phe	Arg	Gly	Ile	Gly	Thr	115	120	125	
20	Ile	Trp	Glu	Val	Arg	Asn	Ile	Pro	Gln	His	Ala	Ala	Phe	Val	Gln	Pro	130	135	140	
25	Lys	Gly	Lys	Asp	Gln	Ser	Met	Ser	Arg	Lys	Arg	Phe	Val	Leu	Arg	Glu	145	150	155	160
30	Ile	Ala	Ile	Ile	Val	Trp	Gln	Tyr	Leu	Leu	Leu	Asp	Phe	Ile	Tyr	Glu	165	170	175	
35	Ser	Thr	Lys	Gly	Thr	Ser	Ala	Glu	Asp	Leu	Met	Arg	Leu	Phe	Gly	Pro	180	185	190	
40	Gly	Met	Glu	Ile	Lys	Tyr	Leu	Asp	Ala	Thr	Phe	Glu	Gln	Trp	Met	Gly	195	200	205	
45	Arg	Leu	Ser	Val	Gly	Ile	Phe	Ser	Trp	Leu	Val	Pro	Ser	Arg	Val	Cys	210	215	220	
50	Leu	Asn	Ile	Thr	Ser	Arg	Leu	Tyr	Phe	Leu	Ile	Leu	Val	Val	Leu	Gly	225	230	235	240
55	Ile	Ser	Ser	Pro	Glu	Ser	Cys	Arg	Pro	Gly	Phe	Gly	Arg	Val	Arg	Asp	245	250	255	
	Val	Cys	Thr	Ile	Arg	Gly	Val	Trp	Gly	Lys	Phe	Trp	His	Gln	Ser	Phe	260	265	270	
	Arg	Trp	Pro	Leu	Thr	Ser	Val	Gly	Asn	Tyr	Ile	Ala	Arg	Asp	Val	Leu	275	280	285	

2009274832 06 Jun 2011

156

Gly Leu Ala His Pro Ser Leu Leu Glu Arg Tyr Thr Asn Ile Phe Phe
 290 295 300

5 Thr Phe Phe Thr Ser Gly Val Leu His Leu Val Cys Asp Ala Ile Leu
 305 310 315 320

10 Gly Val Pro Pro Ser Ala Ser Gly Ala Met Gln Phe Phe Cys Ser Phe
 325 330 335

15 Pro Leu Ala Ile Met Ile Glu Asp Gly Val Gln Glu Ile Trp Arg Arg
 340 345 350

20 Ala Thr Gly Gln Thr Lys Asp Ser Asp Arg Ala Val Pro Phe Trp Gln
 355 360 365

Arg Leu Val Gly Tyr Leu Trp Val Ala Val Trp Met Cys Val Thr Ser
 370 375 380

25 Pro Phe Tyr Leu Tyr Pro Ala Ala Arg Gln His Ala Glu Lys Asn Trp
 385 390 395 400

30 Ile Val Pro Phe Ser Ile Val Glu Glu Ile Gly Leu Gly Thr Ala Gln
 405 410 415

35 Lys Ile Leu Leu Gly Tyr Gly Leu Phe Val Tyr Trp Ala Val Gly Gly
 420 425 430

Glu Ile

40 <210> 276
 <211> 1299
 <212> PRT
 <213> Penicillium coprobium PF1169

45 <400> 276

50 Met Leu Tyr Arg Ala Lys Leu Val Asp Asp His Gln Ile His Thr Ala
 1 5 10 15

Ser Leu His Asn Pro Ile Pro Trp Gln Leu His Thr Tyr Val Trp Pro
 20 25 30

55 Phe Leu Ile Ile Trp Pro Val Phe Phe Ala Phe Tyr Leu Ser Pro Glu
 35 40 45

2009274832 06 Jun 2011

157

5 Arg Tyr Asp Thr Tyr Ile Gln Gly Gln Glu Trp Thr Phe Val Phe Ala
 50 55 60

10 Gly Ser Ile Ile Thr Val Gln Ser Leu Phe Trp Leu Met Thr Lys Trp
 65 70 75 80

15 Asn Ile Asp Ile Asn Thr Leu Phe Thr Thr Thr Arg Ser Lys Ser Ile
 85 90 95

20 Ala Glu Ile Cys Asn Leu Ile Arg Glu His Ile Gly Pro Lys Lys Thr
 115 120 125

25 Leu Ser Phe Leu Phe Gln Lys Arg Arg Phe Leu Phe Tyr Pro Glu Thr
 130 135 140

30 Arg Ser Phe Ala Pro Leu Ser Tyr Ala Leu Asp Ala Glu Pro Lys Pro
 145 150 155 160

35 Ala Leu Lys Thr Phe Gln Gln Ser Glu Gly Phe Thr Ser Lys Ala Glu
 165 170 175

40 Ile Glu Arg Val Gln Asn His Tyr Gly Asp Asn Thr Phe Asp Ile Pro
 180 185 190

45 Val Pro Gly Phe Ile Glu Leu Phe Gln Glu His Ala Val Ala Pro Phe
 195 200 205

50 Phe Val Phe Gln Ile Phe Cys Val Gly Leu Trp Met Leu Asp Glu Tyr
 210 215 220

55 Trp Tyr Tyr Ser Leu Phe Thr Leu Phe Met Leu Val Met Phe Glu Ser
 225 230 235 240

Thr Val Val Trp Gln Arg Gln Arg Thr Leu Ser Glu Phe Arg Gly Met
 245 250 255

Ser Ile Lys Pro Tyr Asp Val Trp Val Tyr Arg Glu Arg Lys Trp Gln
 260 265 270

2009274832 06 Jun 2011

158

Glu Ile Thr Ser Asp Lys Leu Leu Pro Gly Asp Leu Met Ser Val Asn
 275 280 285

5

Arg Thr Lys Glu Asp Ser Gly Val Ala Cys Asp Ile Leu Leu Val Glu
 290 295 300

10

Gly Ser Val Ile Val Asn Glu Ala Met Leu Ser Gly Glu Ser Thr Pro
 305 310 315 320

15

Leu Leu Lys Asp Ser Ile Gln Leu Arg Pro Gly Asp Asp Leu Ile Glu
 325 330 335

20

Pro Asp Gly Leu Asp Lys Leu Ser Phe Val His Gly Gly Thr Lys Val
 340 345 350

25

Leu Gln Val Thr His Pro Asn Leu Thr Gly Asp Ala Gly Leu Lys Asn
 355 360 365

30

Leu Ala Ser Asn Val Thr Met Pro Pro Asp Asn Gly Ala Leu Gly Val
 370 375 380

35

Val Val Lys Thr Gly Phe Glu Thr Ser Gln Gly Ser Leu Val Arg Thr
 385 390 395 400

40

Met Ile Tyr Ser Thr Glu Arg Val Ser Ala Asn Asn Val Glu Ala Leu
 405 410 415

45

Leu Phe Ile Leu Phe Leu Leu Ile Phe Ala Ile Ala Ala Ser Trp Tyr
 420 425 430

50

Val Trp Gln Glu Gly Val Ile Arg Asp Arg Lys Arg Ser Lys Leu Leu
 435 440 445

55

Leu Asp Cys Val Leu Ile Ile Thr Ser Val Val Pro Pro Glu Leu Pro
 450 455 460

Met Glu Leu Ser Leu Ala Val Asn Thr Ser Leu Ala Ala Leu Ser Lys
 465 470 475 480

Tyr Ala Ile Phe Cys Thr Glu Pro Phe Arg Ile Pro Phe Ala Gly Arg
 485 490 495

2009274832 06 Jun 2011

159

	Val	Asp	Ile	Ala	Cys	Phe	Asp	Lys	Thr	Gly	Thr	Leu	Thr	Gly	Glu	Asp	
				500					505					510			
5	Leu	Val	Val	Asp	Gly	Ile	Ala	Gly	Leu	Thr	Leu	Gly	Glu	Ala	Gly	Ser	
			515					520					525				
10	Lys	Val	Glu	Ala	Asp	Gly	Ala	His	Thr	Glu	Leu	Ala	Asn	Ser	Ser	Ala	
		530					535					540					
15	Ala	Gly	Pro	Asp	Thr	Thr	Leu	Val	Leu	Ala	Ser	Ala	His	Ala	Leu	Val	
	545					550					555					560	
20	Lys	Leu	Asp	Glu	Gly	Glu	Val	Val	Gly	Asp	Pro	Met	Glu	Lys	Ala	Thr	
					565					570					575		
25	Leu	Glu	Trp	Leu	Gly	Trp	Thr	Leu	Gly	Lys	Asn	Asp	Thr	Leu	Ser	Ser	
				580					585					590			
30	Lys	Gly	Asn	Ala	Pro	Val	Val	Ser	Gly	Arg	Ser	Val	Glu	Ser	Val	Gln	
			595					600					605				
35	Ile	Lys	Arg	Arg	Phe	Gln	Phe	Ser	Ser	Ala	Leu	Lys	Arg	Gln	Ser	Thr	
		610					615					620					
40	Ile	Ala	Thr	Ile	Thr	Thr	Asn	Asp	Arg	Asn	Ala	Ser	Lys	Lys	Thr	Lys	
	625					630					635					640	
45	Ser	Thr	Phe	Val	Gly	Val	Lys	Gly	Ala	Pro	Glu	Thr	Ile	Asn	Thr	Met	
					645					650					655		
50	Leu	Val	Asn	Thr	Pro	Pro	Asn	Tyr	Glu	Glu	Thr	Tyr	Lys	His	Phe	Thr	
				660					665					670			
55	Arg	Asn	Gly	Ala	Arg	Val	Leu	Ala	Leu	Ala	Tyr	Lys	Tyr	Leu	Ser	Ser	
			675				680					685					
60	Glu	Thr	Glu	Leu	Ser	Gln	Ser	Arg	Val	Asn	Asn	Tyr	Val	Arg	Glu	Glu	
		690					695					700					
65	Ile	Glu	Ser	Glu	Leu	Ile	Phe	Ala	Gly	Phe	Leu	Val	Leu	Gln	Cys	Pro	
	705					710					715					720	
70	Leu	Lys	Asp	Asp	Ala	Ile	Lys	Ser	Val	Gln	Met	Leu	Asn	Glu	Ser	Ser	

2009274832 06 Jun 2011

160

725

730

735

5	His Arg Val Val Met Ile Thr Gly Asp Asn Pro Leu Thr Ala Val His	740	745	750
10	Val Ala Arg Lys Val Glu Ile Val Asp Arg Glu Val Leu Ile Leu Asp	755	760	765
15	Ala Pro Glu His Asp Asn Ser Gly Thr Lys Ile Val Trp Arg Thr Ile	770	775	780
20	Asp Asp Lys Leu Asn Leu Glu Val Asp Pro Thr Lys Pro Leu Asp Pro	785	790	795
25	Glu Ile Leu Lys Thr Lys Asp Ile Cys Ile Thr Gly Tyr Ala Leu Ala	805	810	815
30	Lys Phe Lys Gly Gln Lys Ala Leu Pro Asp Leu Leu Arg His Thr Trp	820	825	830
35	Val Tyr Ala Arg Val Ser Pro Lys Gln Lys Glu Glu Ile Leu Leu Gly	835	840	845
40	Leu Lys Asp Ala Gly Tyr Thr Thr Leu Met Cys Gly Asp Gly Thr Asn	850	855	860
45	Asp Val Gly Ala Leu Lys Gln Ala His Val Gly Val Ala Leu Leu Asn	865	870	875
50	Gly Ser Gln Glu Asp Leu Thr Lys Ile Ala Glu His Tyr Arg Asn Thr	885	890	895
55	Lys Met Lys Glu Leu Tyr Glu Lys Gln Val Ser Met Met Gln Arg Phe	900	905	910
	Asn Gln Pro Ala Pro Pro Val Pro Val Leu Ile Ala His Leu Tyr Pro	915	920	925
	Pro Gly Pro Thr Asn Pro His Tyr Glu Lys Ala Met Glu Arg Glu Ser	930	935	940
	Gln Arg Lys Gly Ala Ala Ile Thr Ala Pro Gly Ser Thr Pro Glu Ala	945	950	955
				960

2009274832 06 Jun 2011

161

5 Ile Pro Thr Ile Thr Ser Pro Gly Ala Gln Ala Leu Gln Gln Ser Asn
 965 970 975

10 Leu Asn Pro Gln Gln Gln Lys Lys Gln Gln Ala Gln Ala Ala Ala Ala
 980 985 990

15 Gly Leu Ala Asp Lys Leu Thr Ser Ser Met Met Glu Gln Glu Leu Asp
 995 1000 1005

20 Asp Ser Glu Pro Pro Thr Ile Lys Leu Gly Asp Ala Ser Val Ala
 1010 1015 1020

25 Ala Pro Phe Thr Ser Lys Leu Ala Asn Val Ile Ala Ile Pro Asn
 1025 1030 1035

30 Ile Ile Arg Gln Gly Arg Cys Thr Leu Val Ala Thr Ile Gln Met
 1040 1045 1050

35 Tyr Lys Ile Leu Ala Leu Asn Cys Leu Ile Ser Ala Tyr Ser Leu
 1055 1060 1065

40 Ser Val Ile Tyr Leu Asp Gly Ile Lys Phe Gly Asp Gly Gln Val
 1070 1075 1080

45 Thr Ile Ser Gly Met Leu Met Ser Val Cys Phe Leu Ser Ile Ser
 1085 1090 1095

50 Arg Ala Lys Ser Val Glu Gly Leu Ser Lys Glu Arg Pro Gln Pro
 1100 1105 1110

55 Asn Ile Phe Asn Val Tyr Ile Ile Gly Ser Val Leu Gly Gln Phe
 1115 1120 1125

60 Ala Ile His Ile Ala Thr Leu Ile Tyr Leu Ser Asn Tyr Val Tyr
 1130 1135 1140

65 Lys His Glu Pro Arg Asp Ser Asp Ile Asp Leu Glu Gly Glu Phe
 1145 1150 1155

70 Glu Pro Ser Leu Leu Asn Ser Ala Ile Tyr Leu Leu Gln Leu Ile
 1160 1165 1170

2009274832 06 Jun 2011

162

Gln Gln Ile Ser Thr Phe Ser Ile Asn Tyr Gln Gly Arg Pro Phe
 1175 1180 1185
 5 Arg Glu Ser Ile Arg Glu Asn Lys Gly Met Tyr Trp Gly Leu Ile
 1190 1195 1200
 10 Ala Ala Ser Gly Val Ala Phe Ser Cys Ala Thr Glu Phe Ile Pro
 1205 1210 1215
 15 Glu Leu Asn Glu Lys Leu Arg Leu Val Pro Phe Thr Asn Glu Phe
 1220 1225 1230
 20 Lys Val Thr Leu Thr Val Leu Met Ile Phe Asp Tyr Gly Gly Cys
 1235 1240 1245
 Trp Leu Ile Glu Asn Val Leu Lys His Leu Phe Ser Asp Phe Arg
 1250 1255 1260
 25 Pro Lys Asp Ile Ala Ile Arg Arg Pro Asp Gln Leu Lys Arg Glu
 1265 1270 1275
 30 Ala Glu Arg Lys Leu Gln Glu Gln Val Asp Ala Glu Ala Gln Lys
 1280 1285 1290
 35 Glu Leu Gln Arg Lys Val
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 40 <210> 277
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 <220>
 <223> a primer sequence for PCR
 45 <400> 277
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 50 <210> 278
 <211> 26
 <212> DNA
 <213> Artificial Sequence
 55 <220>
 <223> a primer sequence for PCR

2009274832 06 Jun 2011

163

<400> 278
 atttaaataag ttagacaata gatatca 26

5 <210> 279
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10 <220>
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<400> 279
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15 <210> 280
 <211> 40
 <212> DNA
 20 <213> Artificial Sequence

<220>
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25 <400> 280
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<210> 281
 30 <211> 40
 <212> DNA
 <213> Artificial Sequence

<220>
 35 <223> a primer sequence for PCR

<400> 281
 tggaagctgg gtagtcaaag tcccccatct ttagcgtggt 40

40 <210> 282
 <211> 40
 <212> DNA
 <213> Artificial Sequence

45 <220>
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<400> 282
 50 ggtacctctg ccatgaagtc ctgatcatt gaccgataat 40

<210> 283
 <211> 40
 55 <212> DNA
 <213> Artificial Sequence

2009274832 06 Jun 2011

164

<220>
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 5 attatcggtc aatgatcgag gacttcacgg cagaggtacc 40

<210> 284
 <211> 40
 10 <212> DNA
 <213> Artificial Sequence

<220>
 <223> a primer sequence for PCR
 15 <400> 284
 gagaggtgct agcagttggt ctagttcgcg agtgatgaca 40

20 <210> 285
 <211> 40
 <212> DNA
 <213> Artificial Sequence

25 <220>
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 <400> 285
 30 tgtcatcact cgcgaactag aacaactgct agcacctctc 40

<210> 286
 <211> 40
 <212> DNA
 35 <213> Artificial Sequence

<220>
 <223> a primer sequence for PCR
 40 <400> 286
 cagtgcctacc tcacggcagt ctggtgagtc cgtccaagta 40

<210> 287
 45 <211> 40
 <212> DNA
 <213> Artificial Sequence

<220>
 50 <223> a primer sequence for PCR
 <400> 287
 tacttggacg gactcaccag actggcatga ggtagcactg 40

55 <210> 288
 <211> 40

2009274832 06 Jun 2011

165

<212> DNA
 <213> Artificial Sequence
 <220>
 5 <223> a primer sequence for PCR
 <400> 288
 agtaaaagcgc gaaaatgctc cggttgtgac tggatgcaac 40
 10
 <210> 289
 <211> 40
 <212> DNA
 <213> Artificial Sequence
 15
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 25 <212> DNA
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 35 <210> 291
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 <210> 292
 <211> 38
 <212> DNA
 50 <213> Artificial Sequence
 <220>
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 55 <400> 292
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