

**(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**

(71) Applicant(s)
INSTITUTE FOR RESEARCH IN BIOMEDICINE

(72) Inventor(s)
Lanzavecchia, Antonio

(74) Agent / Attorney
E F WELLINGTON & CO, 312 St. Kilda Road Southbank, Melbourne, VIC, 3006

(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)

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.

(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

(71) Applicant (for all designated States except US): INSTITUTE FOR RESEARCH IN BIOMEDICINE [CH/CH]; Via Vela 6, CH-5400 Bellinzona (CH).

(72) Inventor; and

(75) Inventor/Applicant (for US only): LANZAVECCHIA, Antonio [CH/CH]; Institute For Research In Biomedicine, Via Vela 6, CH-5400 Bellinzona (CH).

(74) Agent: SMAGGASGALE, Gillian, Helen; W.P. Thompson & Co., 55 Drury Lane, London WC2B 5SQ (GB).

(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,

(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



WO 2010/043977 A3

(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.

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.

5

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 20 dengue infection or passively transferred from mother form infectious immune complexes that 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 antibody neutralizes dengue virus infection.

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.

15 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 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.

20 In a further aspect, the invention comprises an epitope which specifically binds to an 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 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.

30 Figure 2. Anti-dengue virus antibodies that have a CH2 L4A and L5A substitution (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.

25 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 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.

30 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 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.

5 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, 169-10 171, 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, 15 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 5 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 10 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 15 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 20 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 25 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 30 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.

25 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 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 10 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 an exemplary embodiment, a cocktail of the invention 15 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 5 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, 10 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 15 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].

20 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 25 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, 30 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

5 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.

10 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 15 (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* 20 *al.* (1982) *Immunol. Rev.* 62:119-158.

25 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.

30 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--CH₂--CH₂)_n O--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.

5 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.

10 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 15 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 20 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.

25 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 30 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 5 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 10 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 15 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 20 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 25 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 30 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 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 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 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).

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 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; 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.* 5 heavy and light chain genes) that encode an antibody of interest, comprising the steps of: (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

5 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

10 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

15 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.

20 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.

25 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.

30 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.

5 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.

10 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 SynagisTM and HerceptinTM, 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.

25 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.

30 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.*, HerceptinTM 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; RituxanTM 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 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 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 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 thereof, and a pharmaceutically acceptable diluent or carrier.

15 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 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 fragment thereof, and a pharmaceutically acceptable diluent or carrier.

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 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 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) 15 is greater than the sum of the individual therapeutic effects observed with the respective individual therapies.

20

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 5 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).

10 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±2mg/ml NaCl is typical.

15 Compositions may comprise a sugar alcohol (*e.g.* mannitol) or a disaccharide (*e.g.* sucrose or trehalose) *e.g.* at around 15-30mg/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.

20 **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.

25 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.

30 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 5 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, 10 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 15 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 20 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.

25 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 30 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 5 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.

10 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 15 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

20 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.

25 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 30-30 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 5 [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 10 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 15 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 20 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 25 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			
		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)

- [1] Lefranc *et al.* (2003) *Dev Comp Immunol.* 27(1):55-77.
- [2] Lefranc *et al.* (1997) *Immunology Today*, 18:509.
- [3] Lefranc (1999) *The Immunologist*, 7:132-136.
- [4] US 4,676,980
- [5] US 4,831,175
- [6] US 5,595,721
- [7] WO00/52031
- [8] WO00/52473
- [9] US 4,766,106
- [10] US 4,179,337
- [11] US 4,495,285
- [12] US 4,609,546
- [13] Knauf *et al.* (1988) *J. Bio. Chem.* 263:15064-15070
- [14] Gabizon *et al.* (1982) *Cancer Research* 42:4734
- [15] Cafiso (1981) *Biochem Biophys Acta* 649:129
- [16] Szoka (1980) *Ann. Rev. Biophys. Eng.* 9:467
- [17] Poznansky *et al.* (1980) *Drug Delivery Systems* (R.L. Juliano, ed., Oxford, N.Y.) pp. 253-315
- [18] Poznansky (1984) *Pharm Revs* 36:277
- [19] Kohler, G. and Milstein, C., 1975, *Nature* 256:495-497.
- [20] Kozbar *et al.* 1983, *Immunology Today* 4:72.
- [21] WO2004/076677
- [22] Chapter 4 of *Kuby Immunology* (4th edition, 2000; ASIN: 0716733315)
- [23] Jones *et al.* *Biotechnol Prog* 2003,19(1):163-8
- [24] Cho *et al.* *Cytotechnology* 2001,37:23-30
- [25] Cho *et al.* *Biotechnol Prog* 2003,19:229-32
- [26] US 5,807,715
- [27] US 6,300,104
- [28] Traggiai *et al.* (2004) *Nat Med* 10(8):871-875

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 (vii) 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

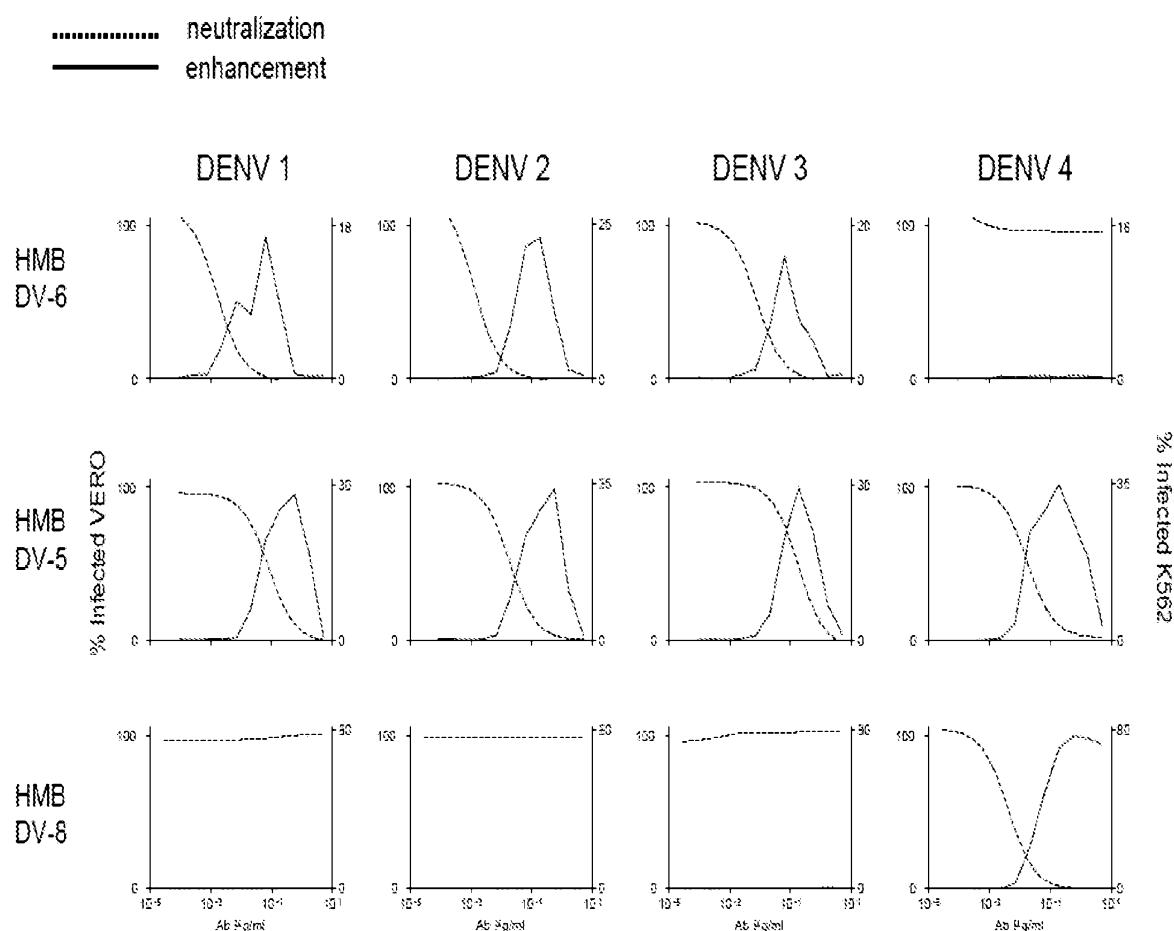
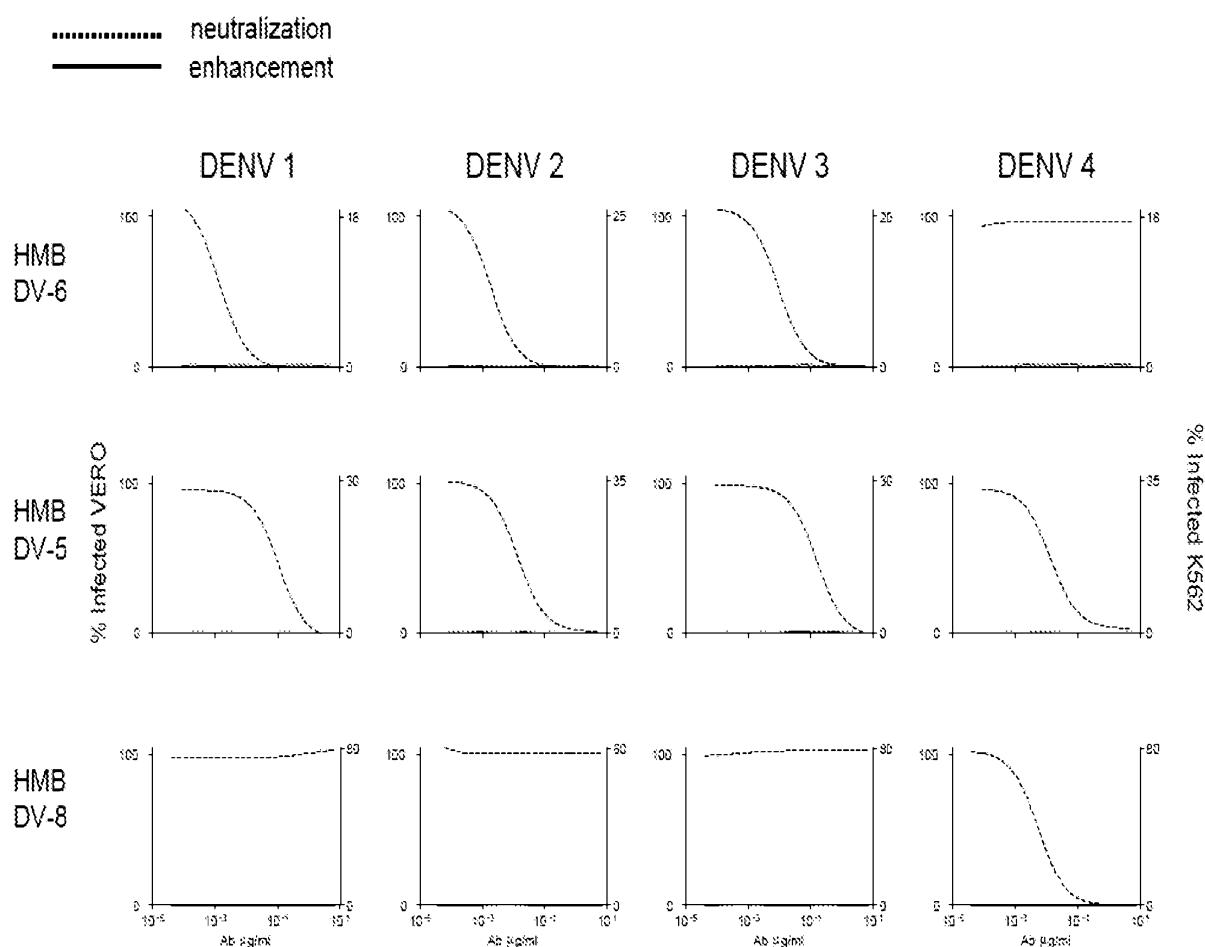


Figure 2. Neutralization and antibody-dependent enhancement of LALA-variant mAbs



SEQUENCE LISTING

110 Meiji Seika Kaisha Ltd.
5 <120> Pyripyropene A biosynthesis genes
<130> 181279
10 <150> JP2008-190862
<151> 2008-07-24
<150> JP2008-270294
<151> 2008-10-20
15 <150> JP2009-20591
<151> 2009-01-30
<160> 292
20 <170> PatentIn version 3.3
<210> 1
<211> 21
<212> DNA
25 <213> Filamentous fungi

<220>
<221> modified_base
30 <222> (6)..(6)
<223> I

<220>
<221> modified_base
35 <222> (9)..(9)
<223> I

<400> 1
40 gayccnmngt tyttyaayat g 21
<210> 2
<211> 20
<212> DNA
45 <213> Filamentous fungi

<220>
<221> modified_base
50 <222> (3)..(3)
<223> I

<220>
<221> modified_base
55 <222> (6)..(6)
<223> I

<220>
<221> modified_base
<222> (9)..(9)
<223> I

5 <400> 2
gtnccnngtnc crtgcatytc

10 <210> 3
<211> 500
<212> DNA
<213> Filamentous fungi

15 <400> 3
cattaccgag tgagggccct ctgggtccaa cctcccaaccc gtgtttatcc accttgttgc 60
ttcggcgggc ccgccttaac tggccgcccgg ggggcttacg ccccccggcc cgccgcggcc 120
20 gaagacaccc tcgaactctg tctgaagatt gtatgtctgatata aattatccaa 180
aactttcaac aacggatctc ttgggtccgg catcgatgaa gaacgcagcg aaatgcgata 240
25 cgtaatgtga attgcaaaatt cagtgaatca tcgagtcctt gaacgcacat tgcccccct 300
ggtattccgg ggggcatgcc tgtccgagcg tcattgctgc cctcaagccc ggcttgtgtg 360
ttggggcccg tcctccgatt ccgggggacg ggcccggaaag gcagcggcgg caccgcgtcc 420
30 ggtcctcgag cgtatgggc tttgtcaccc gctctgttagg cccggccggc gcttgccgat 480
caacccaaat ttttatccag 500

35 <210> 4
<211> 28
<212> PRT
<213> Penicillium coprobiuum PF1169

40 <400> 4
Gln Pro Trp Lys Asp Ser Ile Trp Ala Gly Asp Val Tyr Met Phe Glu
1 5 10 15

45 Gly Asp Asp Ile Val Ala Val Tyr Gly Gly Val Lys
20 25

50 <210> 5
<211> 36
<212> PRT
<213> Penicillium coprobiuum PF1169

55 <400> 5
His Asn Ser Ile Phe Gln Ala Leu Ala Arg Lys Ile Leu Asp Met Ala

2009274832 06 Jun 2011

3

1 5 10 15

5 Leu Pro Pro Gly Gly Gly Ala Pro Ala Pro Ala Pro Ala Ala Lys Arg
20 25 30

10 Pro Ala Pro Ile
35

15 <210> 6
<211> 70
<212> PRT
<213> *Penicillium coprobiuum* PF1169

<400> 6

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

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

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

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

40 <210> 7
<211> 74
<212> PRT
<213> *Penicillium coprobiuum* PF1169

45 <400> 7

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

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

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

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

5 Ser Pro Thr Gly Arg Ser Arg Met Trp Asp
65 70

10 <210> 8
<211> 51
<212> PRT
<213> Penicillium coprobiuum PF1169

15 <400> 8
Ser Ser Phe Leu Thr Ser Thr Val Gln Gln Ile Val Glu Glu Thr Ile
1 5 10 15

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

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

30 Val Thr Ser
50
35 <210> 9
<211> 79
<212> PRT
<213> Penicillium coprobiuum PF1169

40 Phe Asn Ala Ala His Arg Val Leu Pro Leu Pro Ser Tyr Lys Trp Asp
1 5 10 15

45 Leu Lys Asn Tyr Trp Ile Pro Tyr Thr Asn Asn Phe Cys Leu Leu Lys
20 25 30

50 Gly Ala Pro Ala Ala Pro Val Ala Glu Ala Thr Pro Ile Ser Val Phe
35 40 45

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

65 Ser Ala Phe Ile Val Ile Glu Asn Asp Ile Ala Asp Pro Asp Leu
65 70 75

2009274832 06 Jun 2011

C:\NRP\pmb\BDC\KLL\J667199_1.DOC-1/16/2011

5

<210> 10

<211> 84

<212> PRT

5 <213> **Penicillium coprobiuum PF1169**

<400> 10

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

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

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

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

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

30 Ser Ile Lys Thr

<210> 11

<211> 81

35 <212> PRT

<213> **Penicillium coprobiuum PF1169**

<400> 11

40 Gln Glu Ala Lys Ala Met Asp Pro Gln Gln Arg Met Leu Leu Glu Cys
1 5 10 15

45 Thr Tyr Glu Ala Leu Glu Asn Gly Gly Ile Ser Lys Glu Ser Leu Lys
20 25 30

50 Gly Gln Asn Val Gly Val Phe Val Gly Ser Ala Phe Pro Asp Tyr Glu
35 40 45

Met Tyr Asn Arg Arg Asp Leu Glu Thr Ala Pro Met His Gln Ser Thr
50 55 60

55 Gly Asn Ala Leu Ala Leu Gln Ser Asn Arg Ile Ser Tyr Tyr Phe Asp
65 70 75 80

2009274832 06 Jun 2011

C:\NRPon\bDCC\LLJ\67791.DOC (1462011)

6

Phe

5

<210> 12

<211> 66

<212> PRT

10 <213> *Penicillium coprobiuum* PF1169

<400> 12

15 Asn His Thr Gly Arg Ala Glu Gln Ser Lys Ile Ala Ile Ile Gly Leu
1 5 10 15

Ser Gly Arg Phe Pro Glu Ala Pro Asp Thr Glu Ala Phe Trp Asp Leu
20 25 30

20

Leu Lys Lys Gly Leu Asp Val His Arg Glu Val Pro Pro Glu Arg Trp
35 40 45

25

Asp Val Lys Ala His Val Asp Pro Glu Gly Lys Lys Arg Thr Pro Ala
50 55 60

30 Lys Leu
65

<210> 13

35 <211> 14

<212> PRT

<213> *Penicillium coprobiuum* PF1169

<400> 13

40

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

45 <210> 14

<211> 71

<212> PRT

<213> *Penicillium coprobiuum* PF1169

50 <400> 14

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

55

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

2009274832 06 Jun 2011

C:\NRP\bndcc\kll\3467299_1.DOC-1\6/2011

7

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

5

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

10

Thr Ala Arg Arg His His His
65 70

15

<210> 15
<211> 77
<212> PRT
<213> Penicillium coprobiuum PF1169

20

<400> 15

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

25

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

30

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

35

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

40

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

45

<210> 16
<211> 39
<212> PRT
<213> Penicillium coprobiuum PF1169

<400> 16

50

Pro Glu Tyr Ser Gln Pro Leu Cys Thr Ala Ile Gln Ile Ala Leu Val
1 5 10 15

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

55

Ser Ser Gly Glu Ile Ala Ala

2009274832 06 Jun 2011

C:\NRP\pmb\NDCC\ALL3\6729\1.DOC-1\062011

8

35

5 <210> 17
<211> 71
<212> PRT
<213> *Penicillium coprobiuum* PF1169

10 <400> 17
Arg Arg Thr Phe Leu Pro Trp Arg Leu Thr Ser Ser Ala Leu Ser Gly
1 5 10 15

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

20 Ser Lys Glu Pro Thr Val Gly Phe Val Phe Thr Gly Gln Gly Ala Gln
35 40 45

25 Trp His Gly Met Gly Lys Glu Leu Leu Ser Thr Tyr Pro Ile Phe Arg
50 55 60

30 Gln Thr Met Gln Asp Val Asp
65 70

35 <210> 18
<211> 75
<212> PRT
<213> *Penicillium coprobiuum* PF1169

40 <400> 18
Leu Arg Arg Leu Leu His Ala Lys Asn Asp Ser Leu Val Ala Ala Phe
1 5 10 15

45 Phe Gln Lys Thr Tyr Cys Ala Leu Arg Lys Glu Ile Thr Ser Leu Pro
20 25 30

50 Pro Ser Glu Arg Gln Val Phe Pro Arg Phe Thr Ser Ile Val Asp Leu
35 40 45

55 Leu Ala Arg Phe Lys Glu Phe Gly Pro Asn Pro Ala Leu Glu Ser Ala
50 55 60

55 Leu Thr Thr Ile Tyr Gln Leu Gly Cys Phe Ile
65 70 75

2009274832 06 Jun 2011

C:\NRForbNDCCXLL\J67294.DOC (106/211)

9

<210> 19
<211> 81
<212> PRT
<213> Penicillium coprobiun PF1169

5

<400> 19

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

Phe

30

<210> 20
<211> 38
<212> PRT
<213> Penicillium coprobiun PF1169

<400> 20

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

45 Ser Leu Arg Asn Gly Glu Ser Thr Glu Ala Leu Ile Ala Gly Cys His
20 25 30

Leu Asn Ile Val Pro Asp
35

50

<210> 21
<211> 75
<212> PRT
<213> Penicillium coprobiun PF1169

55

<400> 21

2009274832 06 Jun 2011

C:\WRP\onbind\CC\LL\3\m72\9\1.DOC-1467011

10

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

5 Thr Ala Thr Gln Ser Leu Phe Leu Phe Pro Asp Gly Ser Gly Ser Ala
20 25 30

10 Thr Ser Tyr Ala Thr Ile Pro Gly Ile Ser Pro Asp Val Cys Val Tyr
35 40 45

15 Gly Leu Asn Cys Pro Tyr Met Arg Thr Pro Glu Lys Leu Lys Phe Ser
50 55 60

Leu Asp Glu Leu Thr Ala Pro Tyr Val Ala Glu
65 70 75

20 <210> 22
<211> 38
<212> PRT
<213> Penicillium coprobiuum PF1169

25 <400> 22

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

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

35 Val Ala Leu His Leu Ala
35

40 <210> 23
<211> 57
<212> PRT
<213> Penicillium coprobiuum PF1169

45 <400> 23

Ala Ile Arg Asp Glu Val Arg Gln Leu Pro Thr Pro Leu Arg Ala Leu
1 5 10 15

50 Val Pro Ala Phe Glu Asn Val Leu Glu Leu Ala Asn Tyr Thr Asp Leu
20 25 30

55 Arg Lys Gly Pro Leu Ser Gly Ser Ile Asp Gly Val Leu Leu Cys Val
35 40 45

Val Gln Leu Ser Ser Leu Ile Gly Tyr
50 55

5
<210> 24
<211> 74
<212> PRT
<213> Penicillium coprobiun PF1169

10
<400> 24

Ala Val Ala Trp Asp Pro Gln Gln Arg Ile Leu Leu Glu Val Val Tyr
1 5 10 15

15

Glu Ala Leu Glu Ser Ala Gly Tyr Phe Arg Ala Gly Ile Lys Pro Glu
20 25 30

20
Leu Asp Asp Tyr Gly Cys Tyr Ile Gly Ala Val Met Asn Asn Tyr Tyr
35 40 45

25 Asp Asn Met Ser Cys Gln Pro Thr Thr Ala Tyr Ala Thr Val Gly Thr
50 55 60

30 Ser Arg Cys Phe Leu Ser Gly Cys Val Ser
65 70

35
<210> 25
<211> 52
<212> PRT
<213> Penicillium coprobiun PF1169

<400> 25

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

45 Ile Thr Val Pro His Ser Gly Ser Gln Val Lys Leu Tyr Gln Asn Val
20 25 30

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

55 His Gly Thr Gly
50

55
<210> 26
<211> 57

06 Jun 2011
2009274832

C:\WRP\onbind\CC\KLL\U667299_3.DOC (1/16/2011)

12

<212> PRT

<213> Penicillium coprobiuum PF1169

<400> 26

5

Trp Arg Ile Thr Val Ala Ile Val Gly Gly Val Asn Ala Leu Cys Gly
1 5 10 15

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

15 Gly Ser Cys Lys Ser Phe Asp Asp Asp Ala His Gly Tyr Ala Arg Gly
35 40 45

Glu Gly Ala Gly Ala Leu Val Leu Lys
50 55

20

<210> 27

<211> 78

<212> PRT

25 <213> Penicillium coprobiuum PF1169

<400> 27

30 Leu Ile Asp Asp Thr Thr Val Trp Ile Glu Ile Gly Pro His Pro Val
1 5 10 15

35 Cys Leu Gly Phe Val Lys Ala Thr Leu Glu Ser Val Ala Val Ala Val
20 25 30

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

40

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

45

His Arg Pro Phe Glu Arg Ala Leu Cys Leu Leu Asp Leu Pro
65 70 75

<210> 28

50 <211> 65

<212> PRT

<213> Penicillium coprobiuum PF1169

<400> 28

55

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

2009274832 06 Jun 2011

C:\NRP\bNDCC\KL\166724\1.DOC-1\162111

13

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

5

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

10

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

15

Ala
65

20

<210> 29
<211> 83
<212> PRT
<213> Penicillium coprobiuum PF1169

25

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

30

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

35

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

40

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

45

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

Pro Asp Asn

<210> 30
<211> 54
<212> PRT
<213> Penicillium coprobiuum PF1169

55

<400> 30

Gly Arg Phe Leu Ser Ser Asp Gly Arg Cys His Thr Phe Asp Glu Lys

2009274832 06 Jun 2011

C:\NRPort\NDCC\LLU\67299_1.DOC-1\162\111

14

1 5 10 15

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

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

15 Arg Gly Thr Gly Ser Asn
50

20 <210> 31
<211> 63
<212> PRT
<213> *Penicillium coprobiuum* PF1169
25 <400> 31

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

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

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

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

40 <210> 32
<211> 55
<212> PRT
<213> *Penicillium coprobiuum* PF1169

45 <400> 32
Lys Gln Thr Thr Ser Arg Gly Tyr Phe Leu Asp His Leu Glu Asp Phe
1 5 10 15

50 Asp Cys Gln Phe Phe Gly Ile Ser Pro Lys Glu Ala Glu Gln Met Asp
20 25 30

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

2009274832 06 Jun 2011

C:\WRP\mb\NDCC\KLC\U6672W_1.DOC-1\167211

15

Ala Gly Ile Pro Ala Lys Ser
50 55

5 <210> 33
<211> 38
<212> PRT
<213> *Penicillium coprobiuum* PF1169

10 <400> 33

Pro Val Gly Cys Arg Ala Phe Gly Pro Gly Arg Ile Asn Tyr Phe Phe
1 5 10 15

15 Lys Phe Ser Gly Pro Ser Phe Ser Ile Asp Thr Ala Cys Ser Ser Ser
20 25 30

20 Leu Ala Thr Ile Gln Val
35

25 <210> 34
<211> 18
<212> PRT
<213> *Penicillium coprobiuum* PF1169

30 <400> 34
Ala Cys Thr Ser Leu Trp Asn Gly Glu Thr Asp Thr Val Val Ala Gly
1 5 10 15

35 Gly Met

40 <210> 35
<211> 12
<212> PRT
<213> *Penicillium coprobiuum* PF1169

45 <400> 35
Thr Ala Gln Glu Ile Ser Thr Tyr Phe Ile Pro Gly
1 5 10

50 <210> 36
<211> 39
<212> PRT
<213> *Penicillium coprobiuum* PF1169

55 <400> 36

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

2009274832 06 Jun 2011

C:\NRPerl\NCBI\BLAST\BLAST99_1\DOC\1\062011

16

1 5 10 15

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

10 Ser Ser Gly Glu Ile Ala Ala
35

15 <210> 37
<211> 36
<212> PRT
<213> Penicillium coprobitum PF1169

20 <400> 37
Ile Ser Gln Pro Ala Cys Thr Ala Leu Gln Ile Ala Leu Val Asp Leu
1 5 10 15

25 Leu Ala Glu Trp Ser Ile Thr Pro Ser Val Val Val Gly His Ser Ser
20 25 30
Gly Glu Ile Ala
35

30 <210> 38
<211> 39
<212> PRT
<213> Penicillium coprobitum PF1169
35 <400> 38

40 Pro Glu Tyr Ser Gln Pro Leu Cys Thr Ala Ile Gln Ile Ala Leu Val
1 5 10 15

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

45 Ser Ser Gly Glu Ile Ala Ala
35

50 <210> 39
<211> 76
<212> PRT
<213> Penicillium coprobitum PF1169

55 <400> 39

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

2009274832 06 Jun 2011

C1NRF0nB0DCCkLLU667299_1.DOC (1/16/2011)

17

1 5 10 15

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

20 Ala Gln Thr Met Asp Pro Gln Gln Arg Ile Phe Leu
65 70 75

25 <210> 40
<211> 77
<212> PRT
<213> Penicillium coprobiuum PF1169

30 Ile Asn Glu Pro Arg Asp Arg Pro Gln Phe Phe His Ala His Gly Thr
1 5 10 15

35 Gly Thr Gln Ala Gly Asp Pro Gln Glu Ala Glu Ala Val Ser Thr Ala
20 25 30

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

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

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

60 <210> 41
<211> 43
<212> PRT
<213> Penicillium coprobiuum PF1169

65 <400> 41
70 Lys Leu Ala Phe Val Phe Thr Gly Gln Gly Gln Trp Ala Gly Met
75 10 15

2009274832 06 Jun 2011

C:\WRP\enb\NDCC\KL\U66729_1.DOC-1A62H11

18

Gly Arg Glu Leu Leu Ser Ile Ser Thr Phe Arg Glu Ser Met Ala Arg
20 25 30

5 Ser Gln Glu Ile Leu Ala Ser Leu Gly Cys Pro
35 40

10 <210> 42
<211> 71
<212> PRT
<213> *Penicillium coprobiuum* PF1169

15 <400> 42
Lys Ser Phe Leu Asp Asp Leu Ala Phe Thr Val Asn Glu Arg Arg Ser
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

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

Met Gly Lys Glu Leu Leu Gln
65 70

35 <210> 43
<211> 55
<212> PRT
<213> *Penicillium coprobiuum* PF1169
40 <400> 43

45 Ser Val Ala Cys Ile Asn Ser Pro Phe Asn Cys Thr Leu Ser Gly Pro
1 5 10 15

Glu Glu Asp Ile Asp Ala Val Lys Ala Gln Ala Asp Gln Asp Gly Leu
20 25 30

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

C:\VRP\bombCCNLL036672W_1.DOC (116/2111)

19

<210> 44

<211> 68

<212> PRT

5 <213> Penicillium coprobiun PF1169

<400> 44

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

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

15

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

20

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

25 Ala Tyr His Ser
65

30 <210> 45

<211> 38

<212> PRT

<213> Penicillium coprobiun PF1169

<400> 45

35

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

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

45 Val Ala Leu His Leu Ala
35

50 <210> 46

<211> 77

<212> PRT

<213> Penicillium coprobiun PF1169

<400> 46

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

2009274832 06 Jun 2011

C:\WRP\mrb\NDCTKLL\U667299_1.DOC (1062111)

20

Leu His Gln Ala Val Gln Ser Leu Arg Ser Gly Glu Thr Asp Val Ala
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

<210> 47

<211> 61

20 <212> PRT

<213> Penicillium coprobiuum PF1169

<400> 47

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

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

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

45

<210> 48

<211> 37

<212> PRT

<213> Penicillium coprobiuum PF1169

50

Gly Leu Phe Asp Ala Pro Phe Phe Asn Ile Thr Leu Gln Glu Ala Gln
1 5 10 15

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

Leu Glu Asn Gly Gly
35

5 <210> 49
<211> 70
<212> PRT
<213> Penicillium coprobiuum PF1169
<400> 49

10 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

35 <210> 50
<211> 37
<212> PRT
<213> Penicillium coprobiuum PF1169

40 <400> 50

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

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

55 Ile Arg Ala Val Ile
35

50 <210> 51
<211> 83
<212> PRT
<213> Penicillium coprobiuum PF1169
<400> 51

55 Trp Pro Arg Leu Pro Glu Arg Arg Arg Ile Ala Val Val Asn Asn Phe
1 5 10 15

2009274832 06 Jun 2011

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

5

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

10

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

15

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

Thr Thr Thr

20

<210> 52

<211> 59

25 <212> PRT

<213> Penicillium coprobiuum PF1169

<400> 52

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

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

40

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

45

<210> 53

<211> 67

<212> PRT

<213> Penicillium coprobiuum PF1169

50

<400> 53

Leu Ser Pro Gln Asn Asn Pro Glu Asp Arg Cys Gln Tyr Phe Glu Ala
1 5 10 15

55

His Gly Thr Gly Thr Gln Ala Gly Asp Pro Gln Glu Ala Ala Ala Ile

2009274832 06 Jun 2011

C:\NRP\panbNDCC\KCL\JUN\67294_1.DOC (146/203)

23

20

25

30

5 Asn Ser Ser Phe Phe Gly Pro Glu Ser Val Pro Asp Ser Thr Asp Arg
35 40 45

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

15 Ala Gly Leu
65

20 <210> 54
<211> 83
<212> PRT
<213> *Penicillium coprobiuum* PF1169

25 <400> 54

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

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

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

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

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

45 Ala His Gly

50 <210> 55
<211> 38
<212> PRT
<213> *Penicillium coprobiuum* PF1169

55 <400> 55

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

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

5 Ser Val Thr Leu Ser Gly
35

10 <210> 56
<211> 29
<212> PRT
<213> Penicillium coprobiuum PF1169

15 <400> 56
Ser Gly Cys Tyr Arg Glu Leu Ala Asp Cys Pro Gly Gln Arg Gly Ile
1 5 10 15

20 Phe Thr Arg Lys Leu Lys Val Asp Val Ala Tyr His Ser
20 25

25 <210> 57
<211> 38
<212> PRT
<213> Penicillium coprobiuum PF1169

30 <400> 57
Gly Asn Gly Ser Ala Met Ile Ser Asn Arg Ile Ser Trp Phe Phe Asp
1 5 10 15

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

40 Val Ala Leu His Leu Ala
35

45 <210> 58
<211> 59
<212> PRT
<213> Penicillium coprobiuum PF1169

<400> 58

50 Ile Ser Glu Cys Val Thr Val Tyr Trp Lys Ala Ile Lys Ser Ala Gln
1 5 10 15

55 Pro Asp Gly Pro Tyr Ala Leu Ala Gly Tyr Ser Tyr Gly Ser Met Leu
20 25 30

2009274832 06 Jun 2011

C:\NRP\enMDCC\KLLJ667294_1.DOC-1\162011

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
<211> 72
<212> PRT
<213> Penicillium coprobiuum PF1169

15 <400> 59
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

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

35 Ser Val Leu Lys Arg Gln Asp Ile
65 70

40 <210> 60
<211> 79
<212> PRT
<213> Penicillium coprobiuum PF1169

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

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

55 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

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

5

<210> 61
<211> 67
<212> PRT
<213> *Penicillium coprobiuum* PF1169

10

<400> 61

Phe Ile Glu Asp Ser Ile Ser Lys Glu His Lys Pro Thr Arg Val Pro
1 5 10 15

15

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
65

<210> 62
<211> 76
<212> PRT
<213> *Penicillium coprobiuum* PF1169

<400> 62

40

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

45

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

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

50

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

55

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

2009274832 06 Jun 2011

C:\NRPI\BNDCEK1\11667299_1.DOC 1/16/2011

28

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
65

20

<210> 67
<211> 76
<212> PRT
25 <213> Penicillium coprobiuum PF1169

<400> 67

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

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

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

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

55

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

50

<210> 68
<211> 71
<212> PRT
<213> Penicillium coprobiuum PF1169

<400> 68

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

2009274832 06 Jun 2011

C:\NRP\ondCC\KLJ667299_1.DOC (16/211)

30

1 5 10 15

5 Leu Leu Arg Asp Pro Glu Cys Val Pro Met Tyr Gln Cys Thr Asn Ala
20 25 30

10 Gly Gln Ser Arg Ala Met Thr Ala Asn Arg Leu Ser Tyr Phe Phe Asp
35 40 45

15 Leu Lys Gly Pro Ser Val Thr Val Asp Thr Ala Cys Ser Gly Ser Leu
50 55 60

20 Val Ala Leu His Leu Ala Cys Gln Ser Leu Arg Thr Gly Asp
65 70 75

25 <210> 71
<211> 67
<212> PRT
<213> Penicillium coprobiuum PF1169

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

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

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

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

55 Ser Pro Glu
65

70 <210> 72
<211> 79
<212> PRT
50 <213> Penicillium coprobiuum PF1169
75 <400> 72

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

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

20 <210> 73
<211> 40
<212> PRT
<213> *Penicillium coprobiuum* PF1169

<400> 73

25 Glu Ala Asn Leu His Val Pro Leu Glu Pro Thr Pro Trp Pro Ala Gly
1 5 10 15

30 Arg Pro Glu Arg Ile Ser Val Asn Ser Phe Gly Ile Gly Gly Ser Asn
20 25 30

35 Ala His Ala Ile Leu Glu Ser Ala
35 40

40 <210> 74
<211> 70
<212> PRT
<213> *Penicillium coprobiuum* PF1169

45 <220>
<221> misc_feature
<222> (5)..(5)
<223> Xaa can be any naturally occurring amino acid

<400> 74

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

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

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

35 40 45

5 Thr Glu Pro Val Pro Trp Pro Asn Ser Thr Ser Ser Gln Val Lys Arg
50 55 6010 Ala Ser Ile Asn Ser Phe
65 7015 <210> 75
<211> 45
<212> PRT
<213> *Penicillium coprobiuum* PF116920 <400> 75
Pro Val Cys Ser Gly Met Val Lys Ala Thr Phe Gly Pro Gln Ala Thr
1 5 10 1525 Thr Val Ala Ser Phe Arg Arg Gln Glu Asp Thr Trp Lys Val Leu Ser
20 25 30
Asn Ala Thr Ser Thr Leu Tyr Leu Ala Gly Ile Glu Ile
35 40 4530 <210> 76
<211> 19
<212> PRT
<213> *Penicillium coprobiuum* PF116935 <220>
<221> misc_feature
<222> (13)..(13)
40 <223> Xaa can be any naturally occurring amino acid
<400> 7645 Leu Leu Gly Leu Arg Leu Lys Trp Lys Glu Tyr His Xaa Asp Phe Asn
1 5 10 15

50 Ala Ala His

55 <210> 77
<211> 69
<212> PRT
<213> *Penicillium coprobiuum* PF1169
<400> 77

06 Jun 2011

2009274832

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

5

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

20

<210> 78

<211> 85

<212> PRT

25 <213> Penicillium coprobiuum PF1169

<400> 78

30 Asp Ala Gln Phe Phe Gly Thr Lys Pro Val Glu Ala Asn Ser Ile Asp
1 5 10 15

Pro Gln Gln Arg Leu Leu Leu Glu Thr Val Tyr Glu Gly Leu Glu Thr
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
85

<210> 79

<211> 28

55 <212> PRT

<213> Penicillium coprobiuum PF1169

2009274832 06 Jun 2011

C:\NRP\omb\NDC\XLL\0667399_1.DOC (146/2011)

34

<400> 79

Asp Pro Ala Tyr Phe Asp Ser Ser Phe Phe Asn Ile Thr Lys Thr Glu
1 5 10 15

5

Leu Leu Thr Leu Asp Pro Gln Gln Arg Leu Val Leu
20 25

10

<210> 80

<211> 51

<212> PRT

<213> *Penicillium coprobitum* PF1169

15

<400> 80

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

20

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

30

Glu Leu Val
50

<210> 81

35

<211> 24

<212> PRT

<213> *Penicillium coprobitum* PF1169

<400> 81

40

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

45

Gln Met Ala Leu Thr Lys Tyr Trp
20

<210> 82

50

<211> 59

<212> PRT

<213> *Penicillium coprobitum* PF1169

<400> 82

55

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

2009274832 06 Jun 2011

C:\WRP\bNDCCXL\U667299_1.DOC-1\N\2III

35

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

5

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

10

Ser Val Asp Thr Ala Cys Ser Ser Ser Leu Ala
50 55

15

<210> 83
<211> 16
<212> PRT
<213> Penicillium coprobiuum PF1169

20

<400> 83

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

25

<210> 84
<211> 53
<212> PRT
<213> Penicillium coprobiuum PF1169

30

<400> 84

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

35

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

40

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

45

Leu Val Thr Lys Lys
50

50

<210> 85
<211> 60
<212> PRT
<213> Penicillium coprobiuum PF1169

<400> 85

55

Ile Ala Ile Val Gly Ile Gly Gly Arg Phe Pro Gly Glu Ala Thr Asn
1 5 10 15

2009274832 06 Jun 2011

C:\NRPS\1b\NDCC\ALL\J67299_1.DOC-1\06\2011

36

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

5

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

10

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

15

<210> 86
<211> 53
<212> PRT
<213> Penicillium coprobiuum PF1169

20

<400> 86

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

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

Asn Ala Ala Gly Val
50

<210> 87

<211> 18

40

<212> PRT
<213> Penicillium coprobiuum PF1169

<400> 87

45

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

Thr Cys

50

<210> 88

<211> 62

55

<212> PRT
<213> Penicillium coprobiuum PF1169

<400> 88

Phe Leu Asp Asp Leu Ala Phe Thr Val Asn Glu Arg Arg Ser Ile Phe
1 5 10 15

5

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

10

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

15

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

<210> 89

20

<211> 51

<212> PRT

<213> Penicillium coprobiuum PF1169

<400> 89

25

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

30

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

35

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

Val Thr Ser

50

40

<210> 90

<211> 77

<212> PRT

45

<213> Penicillium coprobiuum PF1169

<400> 90

50

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

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

55

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

2009274832 06 Jun 2011

35 40 45

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

15 <210> 91
<211> 78
<212> PRT
<213> *Penicillium coprobiuum* PF1169

20 <220>
<221> misc_feature
<222> (4)..(4)
<223> Xaa can be any naturally occurring amino acid

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

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

50 <400> 92
<211> 79
<212> PRT
<213> *Penicillium coprobiuum* PF1169

55 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

2009274832 06 Jun 2011

C:\NR\Protein\CC\KL\U667291_1.DOC 1/16/2011

39

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

20 15 <210> 93
 <211> 68
 <212> PRT
 <213> Penicillium coprobitum PF1169

25 20 <400> 93
 Leu Ser Ser Asp Gly Arg Cys His Thr Phe Asp Glu Lys Ala Asn Gly
 1 5 10 15

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

35 30 Lys Ala Leu His Asp Gln Asn Lys Ile Arg Ala Val Ile Arg Gly Thr
 35 40 45

40 35 Gly Ser Asn Gln Asp Gly Arg Thr Ala Gly Ile Thr Val Pro Asn Gly
 50 55 60

45 40 Ala Ala Gln Glu
 65

50 45 <210> 94
 <211> 80
 <212> PRT
 <213> Penicillium coprobitum PF1169

55 50 Ser Pro Leu Phe Gly Leu Ala Arg Ile Ile Ala Ser Glu His Pro Asp
 1 5 10 15

60 55 Leu Gly Ser Leu Ile Asp Ile Glu Glu Pro Ile Ile Pro Leu Ser Thr
 20 25 30

65 55 Met Arg Tyr Ile Gln Gly Ala Asp Ile Val Arg Ile Ser Asp Gly Ile

40

5	35	40	45
	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
15	<210> 95		
	<211> 75		
	<212> PRT		
	<213> <i>Penicillium coprobiuum</i> PF1169		
	<400> 95		
20	Asn Arg Ile Ser Tyr Tyr Phe Asp Trp Gln Gly Pro Ser Met Ala Val		
	1	5	10
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	Leu Ser Pro Thr Gly Arg Ser Arg Met Trp Asp		
	65	70	75
	<210> 96		
	<211> 81		
	<212> PRT		
	<213> <i>Penicillium coprobiuum</i> PF1169		
45	<400> 96		
	Val Asp Val Asn Pro Ala Val Leu Lys Asp Ala Pro Leu Pro Trp Asp		
	1	5	10
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

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
<212> PRT
<213> *Penicillium coproblium* PF1169

20 <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 coproblium* PF1169

40 <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

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

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

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

Gly Ser Ile Lys

5

<210> 99
<211> 69
<212> PRT
<213> *Penicillium coprobiuum* PF1169

10

<400> 99

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

15

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

20

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

25

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

Asp Leu Thr Gly Pro

30

65

<210> 100
<211> 79
<212> PRT
<213> *Penicillium coprobiuum* 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

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

50

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

55

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

5 <210> 101
<211> 52
<212> PRT
<213> Penicillium coprobiuum PF1169

10 <400> 101
1 Glu Leu Arg His Gly Lys Asn Ile Asp Lys Pro Glu Tyr Ser Gln Pro
5 10 15

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

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

25 Ala Ala Tyr Val
50

30 <210> 102
<211> 34
<212> PRT
<213> Penicillium coprobiuum PF1169

35 <400> 102
1 Val Gly Phe Val Phe Thr Gly Gln Gly Ala Gln Trp His Gly Met Gly
5 10 15

40 Lys Glu Leu Leu Ser Thr Tyr Pro Ile Phe Arg Gln Thr Met Gln Asp
20 25 30

45 Val Asp

45 <210> 103
<211> 63
<212> PRT
<213> Penicillium coprobiuum PF1169

50 <400> 103
1 Phe Asp Ala Ala Phe Phe Asn Met Ser Pro Arg Glu Ala Gln Gln Thr
5 10 15

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

2009274832 06 Jun 2011

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

<212> PRT

<213> Penicillium coprobiuum 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 coprobiuum 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

Val Val Leu Glu Ser Tyr Thr
65 70

55

<210> 106

2009274832 06 Jun 2011

C:\NR\Protein\NCU\LL\3467399_1.DOC-146-2011

45

<211> 28
<212> PRT
<213> *Penicillium coprobiuum* 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 coprobiuum* 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

65 Val Val Leu Glu Ser Tyr Thr
65 70

40 <210> 108
<211> 50
<212> PRT
45 <213> *Penicillium coprobiuum* PF1169

<400> 108

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

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

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

2009274832 06 Jun 2011

C:\WRP\pub\NCBI\BLAST\667299_1.DOC-1\K\2\H1

46

35 40 45

5 Gly Asn
50

10 <210> 109
<211> 82
<212> PRT
<213> *Penicillium coprobiuum* 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

35 Val Lys

40 <210> 110
<211> 30
<212> PRT
<213> *Penicillium coprobiuum* PF1169
<400> 110

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

50 Ser Leu Arg Asn Gly Glu Ser Thr Glu Ala Leu Ile Ala Gly
20 25 30

55 <210> 111
<211> 38
<212> PRT
<213> *Penicillium coprobiuum* PF1169

2009274832 06 Jun 2011

C:\NR\PenbNDCCV\LL\J667299_1.DOC-1\IG2011

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

Ser Leu Val Ala Leu His
35

15

<210> 112

<211> 72

<212> PRT

<213> Penicillium coprobiuum PF1169

20

<400> 112

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

25

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

30

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 coprobiuum 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

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 coprobiuum* 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
<211> 53
<212> PRT
<213> *Penicillium coprobiuum* 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 coprobiuum* PF1169
55 <400> 116

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 coprobiuum 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

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

65 Val Phe Gly Ala Asp Ile Ser Lys
70

35 <210> 118
<211> 80
<212> PRT
<213> Penicillium coprobiuum PF1169
40 <400> 118

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

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

C:\VRP\bondDC\KLL0667249_1.DOC-146/2011

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 coprobiun 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

<210> 120
<211> 63
<212> PRT
<213> Penicillium coprobiun 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 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 coprobiun PF1169

<400> 121

2009274832 06 Jun 2011

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 coprobiuum 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

<210> 123

<211> 80

35 <212> PRT

<213> Penicillium coprobiuum 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

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

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

2009274832 06 Jun 2011

C:\NRP\Port\NDCC\KLL\3667299_1.DOC (16/201)

52

5 <210> 124
<211> 28
<212> PRT
<213> *Penicillium coprobiuum* 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
15 20 25

20 <210> 125
<211> 64
<212> PRT
<213> *Penicillium coprobiuum* 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
30 20 25 30

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

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

45 <210> 126
<211> 52
<212> PRT
<213> *Penicillium coprobiuum* PF1169
<400> 126

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

55 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

5 Ala Ala Tyr Val
50

10 <210> 127
 <211> 38
 <212> PRT
 10 <213> **Penicillium coprobiuum PF1169**

15 <400> 127

15 Gln Pro Leu Cys Thr Ala Ile Gln Ile Ala Leu Val Glu Leu Leu Glu
1 1 5 10 15

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

20 Ile Ala Ala Ala Tyr Val
35

25 <210> 128
 <211> 86
 <212> PRT
 25 <213> **Penicillium coprobiuum PF1169**

30 <400> 128

35 Arg Leu Pro Gly Asp Val Ser Thr Pro Glu Glu Phe Trp Asp Leu Cys
1 1 5 10 15

35 Ser Arg Gly Arg Gly Ala Trp Ser Pro Val Pro Lys Asp Arg Phe Asn
20 20 25 30

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

45 Ala Ala Gly Ala His Phe Leu Thr Glu Asp Ile Gly Leu Phe Asp Ala
50 50 55 60

50 Pro Phe Phe Asn Ile Thr Leu Gln Glu Ala Gln Thr Met Asp Pro Gln
65 65 70 75 80

55 Gln Arg Ile Phe Leu Glu
85

55 <210> 129

06 Jun 2011

2009274832

<211> 69
<212> PRT
<213> *Penicillium coprobiuum* 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 coprobiuum* PF1169

<400> 130

35

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

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

40

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

<211> 67

<212> PRT

<213> *Penicillium coprobiuum* PF1169

55 <400> 131

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

2009274832 06 Jun 2011

C:\WRD\bndRDCC\KLLU667299_1.DOC 1/06/2011

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

20 Glu Leu Leu
65

25 <210> 132
<211> 21
<212> PRT
<213> *Penicillium coprobitum* PF1169

30 <400> 132

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

35 Ile Ala Asp Val Phe
20

40 <210> 133
<211> 71
<212> PRT
<213> *Penicillium coprobitum* PF1169

45 <400> 133

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

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

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

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

2009274832 06 Jun 2011

C:\NR\Proteins\DCCKLL\1667399_1.DOC-1\1672011

56

Val Asp Val Ala Tyr His Ser
65 70

5 <210> 134
<211> 75
<212> PRT
<213> *Penicillium coprobiuum* 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

35 <210> 135
<211> 53
<212> PRT
<213> *Penicillium coprobiuum* 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

<213> Penicillium coprobiun 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

Ser Leu Val Ala Leu His
35

15

<210> 137

<211> 76

<212> PRT

20 <213> Penicillium coprobiun PF1169

<400> 137

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

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

30

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

35

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 coprobiun 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

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

<210> 139

<211> 57

20 <212> PRT

<213> Penicillium coprobiuum 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

35

Gly Ile Pro Thr His Glu Ile Thr Gly
50 55

40

<210> 140

<211> 68

<212> PRT

<213> Penicillium coprobiuum 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

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

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

Gly Ser Asn Gln Asp Gly Arg Thr Ala Gly Ile Thr Val Pro Asn Gly
5 50 55 60

5

Ala Ala Gln Glu
65

10

<210> 141
<211> 37
<212> PRT
<213> *Penicillium coprobiuum* 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 coprobiuum* PF1169

<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

Asp Thr Ala Cys Ser Ser Ser Leu
65 70

55

<210> 143

2009274832 06 Jun 2011

C:\NRPI\BIN\DC\XLL\U672W_1.DOC-1\62\111

60

<211> 83
<212> PRT
<213> *Penicillium coprobiuum* 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 coprobiuum* PF1169

35

<400> 144

40

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

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

Ala Gly Leu Ile Asn

55 65

2009274832 06 Jun 2011

C:\NRP\bombDCC\LL36672\W_1.DOC-1\162\111

61

5 <210> 145
<211> 64
<212> PRT
<213> *Penicillium coprobiuum* PF1169

10 <400> 145

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

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

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

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

25 <210> 146
<211> 81
<212> PRT
<213> *Penicillium coprobiuum* PF1169

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

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

50 His

55 <210> 147
<211> 38
<212> PRT

2009274832 06 Jun 2011

C:\NRPA\BIN\DCCKLL0667249_1.DOC-1/6/2011

62

<213> Penicillium coprobiun 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

Ser Leu Val Ala Leu His
35

15

<210> 148

<211> 53

<212> PRT

20 <213> Penicillium coprobiun PF1169

<400> 148

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

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

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 coprobiun 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

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
<211> 65
<212> PRT
<213> *Penicillium coprobiuum* PF1169

20 <400> 150

20 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 35 40 45

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

40 Gly
65

45 <210> 151
<211> 78
<212> PRT
<213> *Penicillium coprobiuum* PF1169

50 <220>
<221> misc_feature
<222> (45)..(45)
<223> Xaa can be any naturally occurring amino acid

55 <400> 151

55 Gly Pro Arg Leu Leu Pro Arg Pro Glu Gly Thr Tyr Leu Ile Thr Gly
1 5 10 15

2009274832 06 Jun 2011

C:\WRP\09274832\KLLU667299_1.DOC-1\167111

64

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

5 Gly Ala Arg Arg Val Leu Leu Ile Ser Arg Arg Ala Xaa Pro Pro Arg
35 40 45

10 Arg Thr Trp Asp Gln Val Ala Thr Glu Phe Gln Pro Ala Ile Thr Lys
50 55 60

15 Ile Arg Leu Leu Glu Ser Arg Gly Ala Ser Val Tyr Val Leu
65 70 75

20 <210> 152
<211> 76
<212> PRT
20 <213> *Penicillium coprobiuum* PF1169

<400> 152

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

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

50 <210> 153
<211> 38
<212> PRT
<213> *Penicillium coprobiuum* PF1169

<400> 153

55 Asn Arg Ile Ser Tyr Phe Phe Asp Leu Arg Gly Pro Ser Ile Thr Ile
1 5 10 15

55 Asp Thr Ala Cys Ser Ser Ser Leu Val Ala Leu His Tyr Ala Val Gln
20 25 30

Ser Leu Arg Asn Gly Glu
35

5
<210> 154
<211> 74
<212> PRT
<213> *Penicillium coprobium* PF1169

10
<400> 154

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

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
<211> 57
<212> PRT
<213> *Penicillium coprobium* PF1169

<400> 155

40 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

55
<210> 156
<211> 72

2009274832 06 Jun 2011

C:\NRP\pmb\ADCC\KL\U667299_1.DOC-1\062011

66

<212> PRT

<213> Penicillium coprobiuum PF1169

<400> 156

5

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

25

Ser Cys Lys Ser Phe Asp Asp Asp

65 70

30

<210> 157

<211> 81

<212> PRT

<213> Penicillium coprobiuum PF1169

35

<400> 157

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

40

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

45

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

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

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

Ser

55

<210> 158

<211> 82
<212> PRT
<213> *Penicillium coprobiuum* 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
<211> 75
<212> PRT
<213> *Penicillium coprobiuum* 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 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 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

5 <210> 160

5 <211> 28

5 <212> PRT

5 <213> **Penicillium coprobiuum PF1169**

10 <400> 160

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

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

20 <210> 161

20 <211> 64

20 <212> PRT

20 <213> **Penicillium coprobiuum PF1169**

25 <400> 161

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

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

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

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

45 <210> 162

45 <211> 60

45 <212> PRT

45 <213> **Penicillium coprobiuum PF1169**

50 <400> 162

55 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
<211> 34
<212> PRT
<213> *Penicillium coprobiuum* 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 coprobiuum* 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

45 Gly Ile Leu
50
50 <210> 165
<211> 71
<212> PRT
<213> *Penicillium coprobiuum* PF1169

55 <220>
<221> misc_feature
<222> (5)...(5)
<223> Xaa can be any naturally occurring amino acid

2009274832 06 Jun 2011

C:\WRP\ParbNDCC\ALL\J667299_1.DOC (146201)

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

<213> Penicillium coprobiuum PF1169

<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

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

40

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

45

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

50

<210> 167

<211> 85

<212> PRT

<213> Penicillium coprobiuum PF1169

55 <400> 167

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

2009274832

06 Jun 2011

C:\NRP\pub\NDCC\ALL\3667299_1.DOC (1462101)

71

1 5 10 15

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

20 Gln Asn Phe Pro Thr Tyr Phe Ala Ser Gly Thr Ala Arg Ser Ile Leu
65 70 75 80

25 Ser Asn Arg Val Ser
85

30 <210> 168
<211> 60
<212> PRT
<213> *Penicillium coprobiuum* PF1169

35 <400> 168
Val Val Ala Cys Val Asn Ser Pro Ala Ser Thr Thr Leu Ser Gly Asp
1 5 10 15

40 Val Asp Tyr Ile Asn Gln Leu Glu Ala Arg Leu Gln Gln Asp Gly His
20 25 30

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

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

55 <210> 169
<211> 56
<212> PRT
<213> *Penicillium coprobiuum* PF1169

55 <400> 169
Phe Tyr Gly Met Thr Ser Asp Asp Tyr Arg Glu Val Asn Ser Gly Gln
1 5 10 15

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

10 Asp Thr Ala Cys Ser Ser Ser Leu
50 55

<210> 170

<211> 53

15 <212> PRT

<213> Penicillium coprobiuum PF1169

<400> 170

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

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

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

Ala Leu Val Thr Lys
50

35

<210> 171

<211> 40

<212> PRT

<213> Penicillium coprobiuum PF1169

40

<400> 171

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

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

Phe Ala Ala Leu Asn Ala Ala Gly
35 40

55 <210> 172

<211> 69

<212> PRT

2009274832 06 Jun 2011

C:\HR\Port\BIOCCV\LL\3667295_1.DOC-1A\62111

73

<213> Penicillium coprobiun 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

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

25 <210> 173
<211> 51
<212> PRT
<213> Penicillium coprobiun 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

50 <210> 174
<211> 71
<212> PRT
<213> Penicillium coprobiun 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

C:\WRP\pmb\DC\CKLL\3672\9_1.DOC-1467011

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

<210> 175

<211> 41

20 <212> PRT

<213> Penicillium coprobiuum 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

35 Glu Ile Pro Pro Glu Arg Phe Asn Pro
35 40

40 <210> 176

<211> 64

<212> PRT

<213> Penicillium coprobiuum PF1169

<400> 176

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

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

55 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 Leu Thr Glu Gln Ile Glu Gly
50 55 60

5 <210> 177
<211> 74
<212> PRT
<213> *Penicillium coprobiuum* PF1169
<400> 177

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

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

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

25 Gln Thr Thr Gln Glu Ala Phe Asn Arg Val Leu Ala Pro Lys Ile Ala
50 55 60

30 Gly Ala Leu Ala Leu His Glu Pro Phe Pro
65 70

35 <210> 178
<211> 72
<212> PRT
<213> *Penicillium coprobiuum* PF1169
<400> 178

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

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

50 Ile Ser Met Arg Thr Ala Ile Ile Leu Ala Tyr Tyr Arg Gly Lys Val
35 40 45

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

60 Pro Asp Glu Val Ala Gln Tyr Met
65 70

65 <210> 179

06 Jun 2011

2009274832

<211> 70
<212> PRT
<213> *Penicillium coprobiuum* PF1169

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

30 <210> 180
<211> 51
<212> PRT
<213> *Penicillium coprobiuum* PF1169

35 <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

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

45 Val Thr Ser
50

50 <210> 181
<211> 64
<212> PRT
<213> *Penicillium coprobiuum* PF1169

55 <400> 181

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

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

20 <210> 182
<211> 26
<212> PRT
<213> Penicillium coprobiuum PF1169

25 <400> 182

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

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

40 <210> 183
<211> 65
<212> PRT
<213> Penicillium coprobiuum PF1169

45 <400> 183

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

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

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

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

70 Thr

2009274832 06 Jun 2011

<210> 184
<211> 96
<212> PRT
<213> *Penicillium coprobiuum* PF1169

5

<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 coprobiuum* 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

<210> 186
<211> 59
<212> PRT

2009274832 06 Jun 2011

<213> *Penicillium coprobiuum* PF1169

<400> 186

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

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

20

<210> 187

<211> 31

<212> PRT

<213> *Penicillium coprobiuum* 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 coprobiuum* 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

<211> 38

<212> PRT

<213> *Penicillium coprobiuum* PF1169

55

<400> 189

2009274832 06 Jun 2011

C:\NKP\bombDCC\KLL3667299_1.DOC-1/16/2011

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

<210> 190

<211> 76

15 <212> PRT

<213> Penicillium coprobiuum PF1169

<400> 190

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

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

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

45 <210> 191
<211> 65
<212> PRT
<213> Penicillium coprobiuum PF1169

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

55 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

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

<211> 53

<212> PRT

<213> Penicillium coprobiuum 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

<210> 193

<211> 40

35 <212> PRT

<213> Penicillium coprobiuum PF1169

<400> 193

40 Val Asp Thr Ala Cys 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

<210> 194

<211> 68

<212> PRT

55 <213> Penicillium coprobiuum PF1169

<400> 194

```

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

```

5 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
<210> 195
<211> 62
<212> PRT
25 <213> *Penicillium coprobiuum* PF1169
<400> 195

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

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

45 <210> 196
<211> 76
<212> PRT
<213> *Penicillium coprobitum* PF1169

50 <400> 196

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

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

06 Jun 2011

2009274832

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

5

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

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

15 <210> 197
<211> 79
<212> PRT
<213> Penicillium coprobium PF1169

20 <400> 197

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

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

45 <210> 198
<211> 40
<212> PRT
<213> Penicillium coprobium PF1169

<400> 198

50 Phe Asn Leu Lys Gly Ile Ser Gln Ser Ile Ala Ser Ala Cys Ala Thr
1 5 10 15

55 Ser Ala Asp Ala Ile Gly Tyr Ala Phe His Leu Ile Ala Ala Gly Lys
20 25 30

Gln Asp Leu Met Leu Ala Gly Gly

35 40

5 <210> 199
<211> 70
<212> PRT
<213> *Penicillium coprobiuum* PF1169

10 <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

35 <210> 200
<211> 284
<212> PRT
<213> *Penicillium coprobiuum* PF1169

40 <400> 200
Leu Ser Val Lys Arg Val Gly Ile His Asp Asp Phe Phe Glu Leu Gly
1 5 10 15

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

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

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

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

2009274832 06 Jun 2011

C:\WRP\onbADCCV\LL3667799_1.DOC-1\162\H1

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

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

20 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
275 280

<210> 201

<211> 73

55 <212> PRT

<213> Penicillium coprobiuum PF1169

06 Jun 2011

2009274832

<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

<210> 202

<211> 63

25

<212> PRT

<213> Penicillium coprobiuum 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

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

45

<210> 203

<211> 69

<212> PRT

<213> Penicillium coprobiuum PF1169

50

<400> 203

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

55

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

2009274832 06 Jun 2011

C:\NRP\enbNDCC\KL\J\67299_1.DOC-(1/16/2011)

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		
	50	55	60
	Asp Leu Thr Gly Pro		
	65		
15	<210> 204		
	<211> 96		
	<212> PRT		
	<213> Penicillium coprobiuum PF1169		
20	<400> 204		
25	Phe Asp Ala Ala Phe Phe Asn Met Ser Pro Arg Glu Ala Gln Gln Thr		
	1	5	10
			15
	Asp Pro Met Gln Arg Leu Ala Ile Val Thr Ala Tyr Glu Ala Leu Glu		
	20	25	30
30	Arg Ala Gly Tyr Val Ala Asn Arg Thr Ala Ala Thr Asn Leu His Arg		
	35	40	45
35	Ile Gly Thr Phe Tyr Gly Gln Ala Ser Asp Asp Tyr Arg Glu Val Asn		
	50	55	60
40	Thr Ala Gln Glu Ile Ser Thr Tyr Phe Ile Pro Gly Gly Cys Arg Ala		
	65	70	75
			80
45	Phe Gly Pro Gly Arg Ile Asn Tyr Phe Phe Lys Phe Leu Gly Pro Ala		
	85	90	95
50	<210> 205		
	<211> 58		
	<212> PRT		
	<213> Penicillium coprobiuum PF1169		
	<400> 205		
55	Phe Leu Gln Ile Ser Gly Pro Ser Phe Ser Ile Asp Thr Ala Cys Ser		
	1	5	10
			15

2009274832 06 Jun 2011

C:\NRP\bombiDCCKLL366729\1.DOC-1A&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

<210> 206

<211> 52

15 <212> PRT

<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

<210> 208
<211> 28
<212> PRT
5 <213> **Penicillium coprobiuum 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

15

<210> 209
<211> 71
<212> PRT
20 <213> **Penicillium coprobiuum 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

45 <210> 210
<211> 80
<212> PRT
<213> **Penicillium coprobiuum 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

C:\NR\Protein\NCBI\BLAST\467294_3.DOC-1\162011

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 <210> 211

<211> 38

<212> PRT

<213> Penicillium coprobiuum PF1169

20 <400> 211

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

40 <210> 212

<211> 76

<212> PRT

<213> Penicillium coprobiuum PF1169

45 <400> 212

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

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

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

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

70 75

5 <210> 213
<211> 69
<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

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

30 Ala Asp Ile Ser Lys
65

35 <210> 214
<211> 53
<212> PRT
<213> Penicillium coprobium PF1169

40 <400> 214
Thr Ser Asp Asp Tyr Arg Glu Val Asn Ser Gly Gln Asp Ile Asp Thr
1 5 10 15

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

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

55 Ser Ser Ser Leu Ala
50

55 <210> 215
<211> 63
<212> PRT
<213> Penicillium coprobium PF1169

06 Jun 2011

2009274832

<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

<210> 216

20

<211> 68

<212> PRT

<213> Penicillium coprobiuum PF1169

<400> 216

25

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

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

40

Ala Tyr His Ser

65

45

<210> 217

<211> 39

<212> PRT

<213> Penicillium coprobiuum 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

20 25 30

5 Gln Gln Arg Ile Phe Leu Glu
35

10 <210> 218
<211> 76
<212> PRT
<213> *Penicillium coprobiuum* PF1169
<400> 218

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

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 coprobiuum* PF1169

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

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

2009274832 06 Jun 2011

C:\WRP\mb\NDCC\KLL\3667299_1.DOC-1\06\2011

94

<210> 220
<211> 79
<212> PRT
<213> *Penicillium coprobiuum* PF1169
5
<400> 220

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

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

15 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

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

<210> 221
<211> 81
30 <212> PRT
<213> *Penicillium coprobiuum* PF1169

<400> 221

35 Val Tyr Thr Gly Arg Ile Ser Leu Lys Asp Leu Gly Met Arg Cys Leu
1 5 10 15

40 Pro Leu Cys Leu Phe Leu Phe Leu Trp Thr Ile Tyr Phe Asn Thr Ala
20 25 30

45 Tyr Ser Tyr Gln Asp Ile Lys Asp Asp Cys Lys Leu Asn Val Asn Ser
35 40 45

50 Ser Tyr Val Leu Ala Gly Ser His Val Arg Gly Met Leu Leu Leu Gln
50 55 60

55 Ala Ile Ala Val Val Leu Val Ile Pro Trp Ile Leu Tyr Thr Ser Ala
65 70 75 80

55 Ser

<210> 222
<211> 82
<212> PRT
5 <213> Penicillium coprobum PF1169

<400> 222

Arg His Phe Gly Leu Trp Asp Glu Pro Arg Glu Leu Glu Asp Val Glu
10 1 5 10 15

Phe Leu Leu Lys Ala Asp Val Arg Asn Asn Ser Ala Trp Asn His Arg
15 20 25 30

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

Met Val Asn Ala Gly Asp Leu Ser Thr Ala Pro Ala Glu Lys Gly Arg
25 50 55 60

Leu Ser Val Val Asp Glu Asp Met Val Asp Gly Glu Leu Lys Phe Ala
30 65 70 75 80

Gln Glu
35

<210> 223
<211> 35
<212> PRT
35 <213> Penicillium coprobum PF1169

<400> 223

Ile Met Arg Gly Ala Gly Cys Ala Ile Asn Asp Leu Trp Asp Arg Asn
40 1 5 10 15

Leu Asp Pro His Val Glu Arg Thr Lys Phe Arg Pro Ile Ala Arg Gly
45 20 25 30

Ala Leu Ser
50 35

<210> 224
<211> 86
<212> PRT
55 <213> Penicillium coprobum PF1169

<400> 224

2009274832 06 Jun 2011

C:\NRPonNDCC\KLL366799_1.DOC (1462011)

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
<213> Penicillium coprobium PF1169
30
<400> 225

35 Asn His Arg Tyr Met Leu Arg Phe Gly Pro Arg Asp Thr Ser Leu Pro
1 5 10 15

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

55 Trp Trp Tyr Ala Arg Gly Val Leu Arg Ala Ala Gly Arg Gly Leu Gly
65 70 75 80

Glu Trp

55

<210> 226

<211> 45
<212> PRT
<213> *Penicillium coprobium* PF1169

5 <400> 226

Arg Pro Thr Ser Arg Lys Leu Gly Val Tyr Pro Gln Tyr Ile Leu Gly
1 5 10 15

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

20 <210> 227
<211> 20
<212> DNA
<213> Artificial Sequence

25 <220>
<223> a primer sequence for PCR

<400> 227
tacaggcgcc ctaaattgtc 20

30 <210> 228
<211> 20
<212> DNA
<213> Artificial Sequence

35 <220>
<223> a primer sequence for PCR

<400> 228
40 gaacacagcg caagagatca 20

<210> 229
<211> 20
45 <212> DNA
<213> Artificial Sequence

<220>
<223> a primer sequence for PCR

50 <400> 229
cgcaagactt gaggaacaag 20

55 <210> 230
<211> 20
<212> DNA

5 <210> 235
<211> 20
<212> DNA
<213> Artificial Sequence

10 <220>
<223> a primer sequence for PCR

10 <400> 235
tctcttgcgc tgtgttcact 20

15 <210> 236
<211> 19
<212> DNA
<213> Artificial Sequence

20 <220>
<223> a primer sequence for PCR

20 <400> 236
atgcggcctt tttcaacat 19

25 <210> 237
<211> 20
<212> DNA
<213> Artificial Sequence

30 <220>
<223> a primer sequence for PCR

35 <400> 237
cgacgtaagg agctgtgagc 20

40 <210> 238
<211> 18
<212> DNA
<213> Artificial Sequence

45 <220>
<223> a primer sequence for PCR

45 <400> 238
acctcgatcc tgctgcaa 18

50 <210> 239
<211> 21
<212> DNA
<213> Artificial Sequence

55 <220>
<223> a primer sequence for PCR

55 <400> 239

2009274832 06 Jun 2011

C:\WRP\mbNDCCV\KL\4667299_1.DOC-IMM\2011

100

ggttgtcagg atttgtcaga a 21

5 <210> 240
<211> 20
<212> DNA
<213> Artificial Sequence

10 <220>
<223> a primer sequence for PCR

<400> 240
ttacttcattt cccgggtggta 20

15 <210> 241
<211> 20
<212> DNA
<213> Artificial Sequence

20 <220>
<223> a primer sequence for PCR

<400> 241
25 agagcatatgc ccgggttggta 20

<210> 242
<211> 20
30 <212> DNA
<213> Artificial Sequence

<220>
<223> a primer sequence for PCR

35 <400> 242
cccacccatgttga tttgctcagt 20

40 <210> 243
<211> 20
<212> DNA
<213> Artificial Sequence

45 <220>
<223> a primer sequence for PCR

<400> 243
50 agagcatatgc ccgggttggta 20

<210> 244
<211> 20
<212> DNA
55 <213> Artificial Sequence

<220>

101

<223> a primer sequence for PCR

5 <400> 244
5 gccacacctga tttgctcagt 20

10 <210> 245
<211> 22
<212> DNA
<213> Artificial Sequence

15 <220>
<223> a primer sequence for PCR 22

20 <400> 245
15 aagaacacag agattggtgtt gg

25 <210> 246
<211> 20
<212> DNA
<213> Artificial Sequence

30 <220>
<223> a primer sequence for PCR

35 <400> 246
30 ccaggaagac acttggaaagg 20

40 <210> 247
<211> 20
<212> DNA
<213> Artificial Sequence

45 <220>
<223> a primer sequence for PCR

50 <400> 247
45 agagcatatgc ccggttgtta 20

55 <210> 248
<211> 20
<212> DNA
<213> Artificial Sequence

55 <220>
<223> a primer sequence for PCR

55 <400> 248
55 ccacaccttcga tttgctcagt 20

55 <210> 249
<211> 20
<212> DNA

5 <210> 254
<211> 20
<212> DNA
<213> Artificial Sequence
10 <220>
<223> a primer sequence for PCR

15 <400> 254
tgctcccgcc ttgtctcttt 20

20 <210> 255
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> a primer sequence for PCR

25 <400> 255
agacgtggag ttcctcctga 20

30 <210> 256
<211> 20
<212> DNA
<213> Artificial Sequence

35 <220>
<223> a primer sequence for PCR

<400> 256
caaacttcag ttgcgcata 20

40 <210> 257
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> a primer sequence for PCR

45 <400> 257
caactttccc acccaaagaa 20

50 <210> 258
<211> 20
<212> DNA
<213> Artificial Sequence

55 <220>
<223> a primer sequence for PCR

<400> 258

2009274832 06 Jun 2011

C:\NRP\ntbNDCCVLLJ6672v9_1.DOC (1062111)

104

aagcatgtat cggtggttcc 20

5 <210> 259
<211> 20
<212> DNA
<213> Artificial Sequence

10 <220>
<223> a primer sequence for PCR

<400> 259
cgatacatgc ttctgtttgg 20

15 <210> 260
<211> 18
<212> DNA
<213> Artificial Sequence

20 <220>
<223> a primer sequence for PCR

<400> 260
25 cagcagccct cagcacac 18

30 <210> 261
<211> 20
<212> DNA
<213> Artificial Sequence

35 <220>
<223> a primer sequence for PCR
<400> 261
ggcgatctatc cgcaatacat 20

40 <210> 262
<211> 20
<212> DNA
<213> Artificial Sequence

45 <220>
<223> a primer sequence for PCR

<400> 262
gagacacccgc atacccagat 20
50

<210> 263
<211> 406
<212> DNA
55 <213> Penicillium coprobiuum PF1169

<400> 263

105

cccagccaa gacttgagta gactatattt attctctttg atatccatct cagcatcaag 60
tttttgacgt tgttattacta tcctcgtttg gaattctcct cccaggtctt gcttcattgc 120
5 ttatagcatt ctaccaaaaaa cgtcactgtc atggacgggt ggtcagacat atcatcagcg 180
cctgccggat acaaggatgt tgtttggata gcagatcggg ctctgcttagc ccaaggattg 240
10 ggtatggtcaa tcaactacctt ggcattgata taccaatcgc gcaaagaccg cacatacggc 300
atggccattt tgccactatg ttgcaacttt gcgtggaaat tcgtctacac tgtcatctat 360
ccttctcaaa atcccttcga gagagctgtc ctcacaacat ggtatgg 406
15
<210> 264
<211> 20
<212> DNA
<213> Artificial Sequence
20
<220>
<223> a primer sequence for PCR

<400> 264
25 cccagccaa gacttgagta 20

<210> 265
<211> 20
30 <212> DNA
<213> Artificial Sequence

<220>
<223> a primer sequence for PCR
35
<400> 265
ccatccatgt tgtgaggaca 20

40 <210> 266
<211> 39008
<212> DNA
<213> Penicillium coprobiuum PF1169

45 <400> 266
gccagccaat gtctcgacga gactctcggt gtcaagctgc ttgaggaggc cattgtgaag 60
atcgaagagc gtatcagggtc gcacggcggt agctgcacccg tgaagatggc acccaaggcc 120
50 gtcaccggagc aggacgatgc gatcctgcag gagcttatgg agaagcgcga acgtgagaac 180
acccaggtca gcggagatga ggactctgaa agtgtatgagg gtgttcccgaa gtaagcgcacg 240
55 ggctacaaat tcgagtcgag gggcatacag cggtcaccag cgctaaaatt caaagctgg 300
atcaccggcta gaggggagtt ggtgaaagat ggatagaaaa aacttgacaca tatcggaaaa 360

	aaggctcgat	ggccagttgt	gctgatgggc	aggattacag	tcagaactcg	cccaggtaag	420
	tcgcctggac	ttcggggtct	ggatatgaca	tattcacacc	tgtgtatgcg	gtattccat	480
5	tgcgtcgaa	atcctcggtc	ccggcatcaa	atacactggg	tccgcacagg	gtgcaagttc	540
	tgtgcacat	aatgtttgat	gcaaccgata	cgttcaatgc	cagtcatgct	tttagatgca	600
10	attatccctg	tagaggccat	gtagcaatgt	atgttagcaat	gtatgttagca	atgtatata	660
	caatgtatgt	agcaatgtat	gtaagatatc	ataacaatcg	agctcatgaa	atggcgggga	720
	gagctgaagc	ttatctaccg	ccggcgtatca	ttgggtccct	caaagccatc	gagaacttcc	780
15	cttcggcac	ttctcttttt	ccaccaactt	tcattctacg	cgatatggg	cattggcaaa	840
	agatcttac	cgccgatcgt	ggcaggaccc	ggcatcggt	cgaggtggaa	cgtcgctggc	900
20	gtacgtcatt	ttccaaacat	gcgaaacact	actgacaagc	cgcagtgc	ccggccat	960
	cggccgatga	gtctgacgcc	tcggatgctt	caaaggaaat	tgcaaagg	ttcttcggt	1020
	tgaaatatca	aattgagcag	gttgtctct	gtgaagg	ggagaacg	ttgaccgacc	1080
25	caaacagccg	tatcatcacg	gatgatgtgg	ttgcgactgc	taagcaggcc	ggtgaggat	1140
	aatacaaagc	atgcattgtt	tattgtctcc	tggtttgtct	gcgatgg	tttc	1200
30	catccgtcga	gtttggat	tccgatctcc	atgagattcg	agctgtgg	tgcgagg	1260
	tcgccaagcg	catgtaatgc	cccttttca	ttccatgttc	tcggccattt	cctgacccaa	1320
	acagtatcga	atccgaggcag	aaccaagaat	acgtgc	aaaat	ttta	1380
35	actcaatctt	cagtgaaggt	gtggagactg	atcccgc	tgtcatt	gaa	1440
	atctccatgc	tttaaggatc	atcagctgt	ctgcgtacca	gaagtgtatc	cagtatct	1500
	ggagaggttg	gatctgccc	gaagaaggca	acccaactaa	cttgcgaa	tacagt	1560
40	agtcaaacc	caattattgg	tttcatttcc	atcctgatcg	gatgcgg	actcgtatc	1620
	agaatgtctg	ccaaattttg	ttttccttga	tttaccttgc	gacttatacc	gcagttat	1680
45	ataccgtgaa	tcccaccggt	gacctggatg	tagctgaagc	catactgtat	gttatgactc	1740
	tcgcgttcat	ctgcgacgag	gcggtaaaat	tctggaaaggt	tggatggaat	tatctcgaat	1800
50	tctggaaatgc	gttcaactca	acgctctact	ctatcctggc	agtgtctt	gtttgcgt	1860
	ttattgcctt	ggcacactca	tcatctacgc	acgatgaaac	aaggcaggca	tacaatgaac	1920
	tcagctacaa	cttcctcgcc	tttgcgggccc	ctatgttctg	gatgcggatg	atgctatatc	1980
55	ttgactcg	tttgcgttcc	ggtgcctatgt	tctggatctt	tctggat	atgaaagaaa	2040
	gtttgatatt	ctttgcgttt	ctattcgtgg	ttatggctgg	tttctccag	ggcttcgtcg	2100

gcatggccca agtggatgt gatatccccca tccaccgaaa tattctccag ggaatgatca 2160
5 atagtatcat gcaaagccct gagtttgaca cttttcagga atttgattt ccctttggta 2220
tcatccctcta ttatgtgttc aacttcattt ttagtactgg taagtctgtt ttacatttgt 2280
ttgggggtgtc gctaaacatt ttttagttctg ttgaatattc tcattgcctt gtacaacagc 2340
10 gcatatgaag atatctctgg caatgccacg gacgagttca tggccatctt cgccgcagaaa 2400
accatgcagt tcgtccgcgc cccagatgaa aatgtcttca tcccacgtac gtgtttactc 2460
15 aattctgata tagcatacgt atgactaact ttggtctggg taatagcctt caatctcatc 2520
gagattctct gtttgatagc tccattcgaa tggtggtttt cgccggagac ttacgccaag 2580
gtgaatgaca ttgtatggc cgtgatataat tctccgctgc ttgtcggtgc agcctgggtt 2640
20 gagaccggc aggcgcataa gattcgatgg aatcgccgtc atggcgaaga agacgatgac 2700
tgcgctcagg aatgggagca tggccaaag gaggtaattt ttgatcttga cgataacctgg 2760
25 aaacagcacg taattgagtc cacgccccat atcaagggtt atagttgtac atatgaactc 2820
cgagagctga gggagcaggtaaaatgtt acggggatgg tgaaggaattt gactcaggag 2880
atggaaaaga aggcggatgg agcaagctag gaagtcctgt tgaattgtac agcaagaata 2940
30 ctacactgag catgggacat cgcaaagggtt atttgctact gcagttcac caatattaca 3000
ttgcgaaaac tggatattct cttaatgtct aatagcagca atcagcccg tggcacggag 3060
35 gaaagtcacc gtcctgttaag gcaaatactt gtgcttcaaa tgaattttga ctattttca 3120
tgcgataact ggcaaagggc agggggagaa aaaatgtca ttattcaacc caagcaaact 3180
gtccagaaag tgacatgcc actttgcaag taaagaagat atgtgacaat ctaacagtct 3240
40 cagtagaca ttgcgttcc attaaaatcc atgcgttgct cggccgtagcc caattcgaag 3300
cactggccaa cccacatcga gacctaaaaa tgggtgtatc atcacacagc aacaggctca 3360
45 gcaagaatgg aggcaatgtt ctccctttga tgatccagct gtgagagctt cgctcgatgg 3420
tgcttgccaa tacatatccg aggaatgtca tccacaaaata caacacccctt atctagggt 3480
ttatagctgg ccagttggct ttgaatcaga cctgccactt gatcgccgt cgtctccggg 3540
50 gacgtatcat tgcggacgac ataagctcga ggaacctcgc tgctgccatc tgggagcatg 3600
actccgatca cggctgcgtc cttgatactc gggtccttgc gtggatccc ttcaatctct 3660
55 gcgggagcga cggagtatct aaatcaagca cgatatgtt aatgtatcatc tgctgcatacg 3720
gatgtcatat ggagaggaaa gaaagcgaag gatgtgaagg atgaaggccta gaggactggg 3780

taacttgcgg tcgaactttg atgagatctt tggccgtcc gatgacatgg tagttccgt 3840
cttccacatg gaacatgtct ccagtccgga accatcccttgc tctatctttg gcatcagtcc 3900
5 gtcctttgtat tgctagaagg agtcccggtc cacggacata caactctcca ggggagtcgt 3960
gtgtcccggtc gacatcttcg cccgtgtcgg gattgacaaa ggcgcagctca tatctggca 4020
aaagagtccc tacactgcca aattgtggtt gtatccgtat ggcattctgg aaaaccactc 4080
10 caacccatcaga catgccccac agatttcccg ctatagcgtc cgggtatagc aggctctgg 4140
attgtgtcat agagtacccg tctatggag cacccgaaat accgatatacg cgccagagaag 4200
15 acaagctctc ggctacattc aaggaggacc tattgagaat gtggatcatg gcaggaacca 4260
tgtacgtttc cgtgtatgtgg tgctggcgga tgccgtcgag caaagcgggtg atttgcgaagc 4320
20 gcgggataat gtacagagggc tggccgtacc gaatggggaa gatgttgcggc cagaagtcgc 4380
caaaaagaatg gtacagtggc agtgccatca aacgaacgac ggggtatggc acttcatagt 4440
agacgctcag atgggtggaa atgatcggtt ggtgggttcg aattgcggct ttggggagac 4500
25 cgctggtgcc actgggttagg aacatagccg ccggcgtgat cttgtctcc tcgctatctt 4560
cgaaacgaag ccaatccaac tcgcccatact ggagcagact ctccaggcgg ataggttgg 4620
30 ccactgtctg ggtgtcgaga tcctccgtct gctccgcctg gccatgtgca aattggacta 4680
cactttcgat agacttctca tccatcagaa ggacttggtt tgaggacatt ctttgattat 4740
tgcaaacttc caggactctg gtcagcgcac tcggagcagt aataatcaac cgaggctcgg 4800
35 cgacacgaag cagatgagcc acttcatggg ggcgcactagc gacatcaaac cccatataca 4860
ctccgcccgc accaacgatg gcaaaagaaaa gagcagagtg tagaacctag gctgtatgtta 4920
40 gcagcatatc attgttatgg ggttagtgtga ttacactgtt ctccagttgc acgagtacac 4980
aatcgccctcg ttccacaccc cgggcttga ggcccgcataat gagtgtatgc accagccgtc 5040
ggaatttggat ggcattgaaa gcacgcgaag ggttgcgggc atcaatatacg atgggcttag 5100
45 attggtcataa ggcaggacca ctaaaagcaa agctgactag gtctgtctcg tgctccatata 5160
cgatgtttat attgtacagt tctcgtgtgc tattgacatg cagaacttgc tgcaggattt 5220
50 gtgcctcactt taagtagtag tacatggaaat gctcagacactt cccatatacactt tttgtatgcac 5280
actgcacggg acaagtatca tgcagaagac tattgagaag aatgccacgc caccaattcg 5340
tattatacta atcttagccta agccaataca tgtaaagagt actatttagg acccacactg 5400
55 tcattgcaga gctttgaagc agctgcattgc gctaattcactt ccacagatac gccactaaga 5460
atcaaaaatta ccccgatgtc gacgctcagc tctttcgtaa accattgact cagcccaatg 5520

5 gcgataagcg agtcaatccc gagctcagga atcagcgtgt cagcagagag cggtgcaccc 5580
tcggccaagt tcaagctggc ccgaattttc tccatttagtg gtctcacgac tgcttcggct 5640
ttttcttcca agcttgcgc tgcagtgaca agatcttgg tcgaccgggt ctcaatcaat 5700
gcaagtatcc gatcttgaga cgccgtggcc gtataggagt agaatggcca taacttcggt 5760
10 atcgggcact cgccatagcc acacttcaaa ctctggtgc ggagtcccc gattaactca 5820
gcgttggaat tggaaatctga gcgcggcag aggattgcct cggcgagtat ctcgtcgaca 5880
15 tccccctggg atacagctac cgggccacac caaagaggct gactgggaga aggactagaa 5940
atgccgtgga tctcgcccaag atgtactaga ctgtgcggc tgccctggc tcgacgggtgg 6000
cggaccagca gggccatttt ctggacatc gtcgcgtca ttgcctggc cgcatggcct 6060
20 aacactcctg caatagatcc gatgagcacc caaaagtcca gagttgggtt cttgttagac 6120
tcatctagct gctgcagccc cttcaagacc ggatgcagat ggttccggag ggaatctatc 6180
gtgagctggg acaaggaaca gtcaggtaga ggccggaggct gaattagaac tcctccact 6240
25 accgggggga acgcatacgaa aatagtttgta tgcaaactgg tggcagaaat gccatcgatc 6300
aggttctgg cattgattag ctgtgcatt tgaattcatg ggtacgtaaag taacttacat 6360
30 tttcgagacc gctatgcgcg ttccccgtcg agaaacttct tccaaccacc acgcacatctga 6420
gtcaagtcta gagccagcaa ggaggatcca ttttgcggcg tgggttgcta gccaaagaca 6480
35 gatagcatgg gctagttcac tgcctagacc tacaagtatg taggttttt tctctgacag 6540
ctgcacctgg gaaccagcag tcggtatctg ggccgcgcacc ggagttgttag agtcccaatc 6600
taccacggca tcctgggagt caaggattgg gtactcgaa atcctgctga ttggtaacga 6660
40 gtccacagaa tttgggtggga gcccctgcgc gcccgtgtac gccactaagc aggccgttag 6720
gaaaggccttg gctatgagtg acgaatcatc cgcgttgcgc ggccctgtcg atgcagaggt 6780
45 aaggtagaaa tcctgcaaattt ggattcgtgt cgcgttgcac ggcaagagcg atagcatgcg 6840
atcatagaca ccctgtccac gacgatgttag aatcgctatg gccgacacat ctgagggaag 6900
tacctgagac aattgccgcg cagtgcatac ctgcgtcaaa agcaacatcg gtctttctt 6960
50 gtctgggttg cttttgcttag tgctgaagat aaccttacca tcccgccggg tcagcatttg 7020
gtggaaaaca gactgaagga ccccatcagc ctgcgtgcact acaagcgtgc ctgattgtgg 7080
55 gacttgcgtcg accaggtatc ctgccaacag agcagccgcgt gtggcgccgga gataagactg 7140
ctcggtggct tccaaacaccg tgcgtggcac tgaccacgac caggagtctg gaacgataac 7200

110

	atgagatgca atgtgggacg acagagcgat cattctcttg ttgctcttaa cgtccagccc	7260
	5 tatgaccaggc cgccagaata tggccctgc aacccgcacg gctgctatac tggaatgtcg	7320
5	aactcgcaga tgaagggttggccataact tgccgttatac ggcggatcag ccatcgaaag	7380
	aaggcggAAC ccttcacaag tcttgcggc cgtacgacggaggatttgcggatccatccctt	7440
10	10 gtcgagggtcc acgcagtctg tcactttttgc cgcctggct agatggcgca gcccagtagc	7500
	atggtcgtga tattgccag gaacacggaa catagagccg tcgtattgaa tttcgggctc	7560
	gatgttggtg agtccgcacg aatgaggatt ttcttgagct gcactagctt gaacaaagtgc	7620
15	15 tccaaagact gtcgccaata tttccgttgt aactccaacc ggatctgtga tatgttagcag	7680
	ttgttagcaga gaagacgatc gttcagaagc caggaagggtg gaaaggagac ctttcaccag	7740
20	20 tccggcatca ggggtggtcga ccttctcaca ggtcaccact aacattcgct tactgactgt	7800
	catcagtctg cacaactcgc tcaaggttagc gtttgcaga tctcgatcat cgacgaggta	7860
	cagaacagtc aatttagata aatcgatc ttcaatgaga tcttaggtctg ggcgcattgac	7920
25	25 aacccgtacg aagtcatcct gtaccaattc aaaaagctct gaggttaagac aatctgcctc	7980
	ctcagcgtca ccggccaaagca acagcaggatc tccgcggctc tgggttccg caggactgtc	8040
30	30 tggagtgcga cagagaagga gtgaaaagtc cccaaaggctt tcactctt gggacgcata	8100
	gaacgaatcc aagccataga atccgcgtt agacagtagc tcaacccagt ccctcctgg	8160
	agtaattggc tcaccagagc agtagcctt cccggctct gtgcattca ttggaggacc	8220
35	35 gaatagaaga ttcaaataatg tagtgctggg attcggttgt actagtagga ccaagaatcc	8280
	accaggcttgc acgaaggcgac gaacatgagc caccgcgacc tcttgcagaa ataccgcggc	8340
40	40 tgtgatcagc accatatcgt agaattgctc gcggcagccc tgctcgacag gatcctcatt	8400
	gatgtccaaac gttttgtgcg acacctcgcc aggttgctca aggtttccct caatcgcttg	8460
	taggccagaa acggagagtc cagcataagt aaatgaccga taagtccgac ccattttttt	8520
45	45 cagtcagatgc tgaacatggc ctccaaatttgc gccgatctga aggttattca tttgtggaaa	8580
	gcggaaacac gcctggctga cgacagatac gagtcgtct tctagatcca agacttgcaa	8640
50	50 gtcctccttc agatattgac ttccatc gatggccggc caggcctcta tttgaagacc	8700
	agaatcagc agaacgcgag gtagccgtg gcctaccgca gcaatggccg tgagaccagg	8760
	gtcattcaag agtgacgggc tcactccagc tgtgaagtcc tcaatcttct ggtccaagca	8820
55	55 ctcggattcc ccgacaggggt ctggcttttg gctagcattt gcaatgcatt ggttcatcca	8880
	tgcaaggcaga cgagcaccat cgaaatccaa tccgcttcgc tccaagtccg tcagtcatt	8940

gcgagcctgc ttgagataca gtagcgcaag ctcctctcg agagagtgt a gctgttagcat 9000
5 ggtagctggc aacttccgtg atcccttctt cagagtggc tcaagcggtc cccacgcgg 9060
ctggaaaga acctgcaagt tgtttggtgc agttccagat ggctggcata tgagagaaac 9120
accttcgagc tggacagcct tttccccatt catggtgaag atgtcaatat caccgcgaat 9180
10 tcgatctcca ttaacgcagg tcagatagct tgctaccgtc aattccttgc cttgccaatc 9240
tgaagcacat aacaccggat ttatccaggt actgtcaaca tttctcgata agaatggtcc 9300
15 cgtcagcagc gtctcttcaa gcccaccaat tgcagcaatc attgtttgaa caccaagg 9360
caaaatagcc gggtaagag ccatgggctc atccgaatca tttgaggaa cgggcacact 9420
cccagtggct agatcacgccc ttttgcggag tcccgtcaag gtagagaatg ggccagtaca 9480
20 gtggtagtca ggcggcgca ggctgtcata gaattcagtg ctgtccacgg gctccaaggc 9540
ctgaggttagc tgcgtctgtg ggggttaggag agcacggtca gaatcccctg gatgcatgat 9600
25 catctggct gttgcacact gaacgagctc tccggataca acagcttcgc agcagaacca 9660
agcagtaatg gctccatcat gcgagtgaat actacccacg gtgacaagca cttcagtgcc 9720
gatggatca ttctgaatcg ggagctgagt gtggatggtc aagtccctga cattcaacaa 9780
30 gcgtaggcct tgcgtctgtg ccattatcat acctgcctcc agtgcacatcg atatgtatcc 9840
tgtctcagg aagacagatc ccgaatcggc acgacggtgc gccaaccagg gcagctcctc 9900
35 tggtcgtaga tagttcgcc aacggaactt ttctgctccg gtctctggac tgagagaacc 9960
gagaagtgcg ttaggagatg tagcacgatg gttatggttc gaagacattc gcgactgtgt 10020
ccagtatgtc tgagtatggt cgaagggta gaatggtagc gattctacca acacaggcca 10080
40 atgatttgg a tcaaagagtg agacatagtc tgcgtctgtg acgacatttg ggccgagg 10140
tgcccaggaa gatcctaggg ccgttgcaca tgcgtctgtg ccgggcttcc ctcgctcagc 10200
45 aagagcaagg taaggaattt ccgagttggc cgagtgcatt ttggagaggg tctgttaggac 10260
aggcccttc agtgcggat gggcccgat ctcaatgtatg agatctggtg gcccagcg 10320
tcgtgccg 10380
50 gtactcccct gtcaatttcc tgcgtctgtg ccagtcatg acctcccctg ggttagacact 10440
cgagtaccag cgcgaggcag aagctgacag ggcgacagga tacgtttca ttgcgtc 10500
55 atatggatct gcacaaggat tcatatgcgg agagtgtat ggcgtgtcca ctcgaagcat 10560
acgcggagtg aggcccaggc tcttcagcag ccactccagc tcccgcaggc actctgcgtc 10620

	gccggataac gtgacgctgg atggcgagtt ggcggcagca acgcttatac gtccggagta	10680
	ggcttctaaa gcacagatat tctgcgcctg ctgccatgtc aaattcacgg ccatcatccg	10740
5	acctgtcgga tcgcgtgact tatcaatggt catcccccta aggtacgcga tgccgattgc	10800
	atccgaggcc gtcagcacac ccgcagcata ggctgctaca atctcgccgg aagagtgacc	10860
10	gaccacaatg gtaagctcaa tccctaccgc acggagcatg ttgacttgca tgatttgcaa	10920
	cgctgtccgt agggggagag aaaggaggcc ctcggttacg cgcgaggacg atgccggctg	10980
	tgacaactcg tcgagaagag aaaactgtgg acgaaggct agtggaaagct catccaaagc	11040
15	ttcctccaga ttcataatcc attttcaat tgagggactt gcctcaatca gatcaagtcc	11100
	catttgcgtt cattggactc cttggccctgt gaagatgccc atgacgcgtc tggcccgagt	11160
20	gttggatctg gagacgacag aggctggtt acccggtgacc cttcgactta ttctgtatt	11220
	gatctggtct ttcaactctt gtattgagtg tgccattagc gtcaaccggt ggcgtatgagt	11280
	ggaacgtcga tcccacaaag agagcgccag accaacgaga ctgactgtt cgtgttccctg	11340
25	gaggaatgtt gcgtatgatt ccatcacaca agtgggggtc cgctcagacg cagcagaaaa	11400
	gacaaagggc agactggagg gatatgtt ggacggactg agctccgagc gagttgttagct	11460
30	ttcttaggacg acatgcacat tggcaccccc gaatccaaag gagttcaccc aggtcgacg	11520
	aggacagcca tctggactg caggccacgg gatgcattct gtggggacag aaagcttggc	11580
	ggcaaacggt ttaattttt gattgagatg ctgcatcagg agattaggag caatcatccc	11640
35	gtgttgcagc gagagggatg cttgtatcaa tcccgctagt ccagcagtgg cctctgtatg	11700
	tccaaattatt gtcttaattt acccaacata cagccgatcg gtcgaatctg gaacggattc	11760
40	gggccccaaag aagctttagt tgattgcggc tgcttcctgc ggatctccgg cctgggttcc	11820
	cgtgccatga gcctcgaagt actggcacccg atcttccggg ttgtttttag gagagagccc	11880
	cgcgcgtgca tagttgcga ggtcaatga ttgttgcgc tttggattag gcatcgtat	11940
45	ccccatagtt cgcccatccg agtttgcctt tgaggcacgg atcacgcatt cgataggatc	12000
	tccatcatta atcgcgttccgcagacgttt tagtacaacc gaagccacac cctcgccacg	12060
	tccgttagccg tggccttgc tggccacat tctactccgg ccggtagggg acagcatccg	12120
50	tgttttagaa tccgcaatat aggcatggg agacaggatc aggttgcttc ctactgccac	12180
	tgccatggaa cagtcatcgt tctgcagagc ctcgactccc agatgaacag ccaagagact	12240
55	cgaagaacat cgggtgtcaa cggccataga aggaccttgc cagtcaaagt agtaagagat	12300
	acgattggcc atgattgacg gtgagttcc cgttaaccaca tacgcgggaa acgcctgagg	12360

atccatggcc tggatttgat tgtaatcggt gcgaagtgtt ccgcagaaca ccccggtctt 12420
5 tgagcgctgc agcgcatcca tccgttaaccc ggccgcacatcg agcgattcgt acacggcttc 12480
taggagcaat cgctgttgtg gatccattgc taccgcttca gttggcgaga tattgaagaa 12540
ggccgcacatca aaggctttaa tgcctcgctt caagaagtat gactctttaa cgtttgtt 12600
10 gccatggtgg tctccatctg gatgataaaa ggcacatctata ttgaatctgt cggccggaac 12660
tttgcgcgcg atatcccggag ggctttgaag aagctccac agtttgcgaag gagaggaagc 12720
15 gccaccggga aagcggcatc ctgtaccaat aatagcaaca ggctctgttg ctttcattgt 12780
gagattataa gagaggtgtt aacacgttgc tcaaaataat ttgcagttgg gtggctgttag 12840
ctctactgag agtacgttca tagatataag caatgcgttgc ttgccttact tacttccacg 12900
20 atcttgcag catatctatc gaacgaatag caaaaacttggc cctatagagc aatttccggc 12960
catcgataga tcattggata gctgtcctat ttgggaagta tgcgttacaa ttatgcage 13020
25 cacaactat acaaagtggt ccatgcgcag atttggcgat gagcagcggt gtggaaatagt 13080
gactttgatg aacatgtcag gtccgtcattc tacatgttgc ggtgtccaag gatgtcctt 13140
gcgcaagaa gtggagtagg gacattcagg taccttcctt tctttccct tcttttaatg 13200
30 ctcactctgt gcataataat agtggcgaat atcgaagcat cggaaatccaa cgacatttgc 13260
acaacatggc taacatggac aacatgttgc acacaccctt aggttcaac tgggcctggg 13320
35 cagtcatcat ctcttcctt ggtctgttgc cttttcctt tgcgttgcac caccttttc 13380
cttcaagatt gacggtgatt aatggtgaa gagcctggaa tatcttttgtt accaaggcc 13440
aaaagcgatt tcgctcgac gcagcacgtc ttataaagaa cggcttcgag gaggtgagta 13500
40 tggaaaaact gcatcatatggaa ggataaagtgc taaacgttc ctcttactc cagtcttc 13560
atgcctttcg cattatcactg gataacggtc ctttgctggc cttgtcaccc caatacgctc 13620
45 gtgaggttcg cagcgatgtt agactcagcc ttgaccattt cattgcctcg gtttgccttgc 13680
cttcatgtcc aacgttttc tagttggcgat cgctaaatgc ttactgttta ggaatttcac 13740
ccaaacatcc caggtttcgat gcccgttcaaa ttgatcttgg atccaaagaa cccgttgaac 13800
50 acgatcctca agtccaatct cacacaagca ctgggtactg acatgttgc ctccgttgc 13860
atgcagccca ttacatgttgc aacattgtttt acctggatag cttatctgac agaggacttgc 13920
55 tctgcggagg taacagagggc actatctgca acctgttaccg atgaccctgg taagctataa 13980
aacatggttt tccaaagggtt ctggtatcaa tactaactttt ctttttctc ttaatcaaag 14040

agtggcacga ggtcagcgtt agtcaaacgg ctctcaaaat tatcgacaa atggcggtcca 14100
aagccttcat tggacaagaa agatgccggg atgccaagtg gcataacatt atcatcacgt 14160
5 acacgcacaa cgtctatgga gcagcacagg cactccactt ttggccagt ttcctacgac 14220
ccatagtggc acagttttg ccagcatgcc gaactttgca ggctcagatt gctgaagcgc 14280
10 gagagatctt ggagccattt gtagcccaga gacgagccga gagagccacc cgagccgctc 14340
aggagaagcc tcatccgtct ggtgggata tcattgactg gctggAACAG ttttatgggg 14400
accaaccgta tgatcccgtg gcccacacg tactgctctc atttgctgct atccatggaa 14460
15 cttccaatct cctggcgcaa gcgcctcatag atctctgtgg ccaaccggag ctgtacagg 14520
atctccggga agaagctgtg tccgtgtgg gtaaagaggg atggaccagg gcccgttgt 14580
20 accaactcaa actaatggac agcgcctga aagaaagcca ggggttggcg ccaaacagat 14640
tgtgtgagtg ggccttcct cttcccccc aatttgcacca ttcaactggc cattagagac 14700
taattcaggt gtgttttac agtacgtatg ggacgcattt cgcaaggcga tatggacctg 14760
25 tctgtggtc tccgtatcca cggggcacg accctcatgg tgtctgccc caacatgtgg 14820
gatcctgaaa tctaccctga tccccgaaaa tacgtatggct accgattcca taagttgcga 14880
30 caaacatcag ggcaagaggg ccagcaccaa ctcgtatctt cgacgcccga tcacatggga 14940
ttcggatacg gaaagcatgc ttgccccggc cggttttcg ccgcagccc gatcaaagtt 15000
gcattgtgca atatcctctt caagtatgtat attgaataca ggggtggcaa gtccccaggt 15060
35 gtgtgggttc agggcataca tctgtttccc gatccgacgt ctaggatcca cgtccgtcgt 15120
cgaaaaagagg agattaactt gtgatactat tgtctaacta tgccgatgtg gttgaatgca 15180
40 aggactctct ctctctctt gtctgatttga tatttgagtt ttctatggc atcgagcaag 15240
atttttgcacca tgtggagccc atgcattgtc atgaggccta ttggccgat ctcttcgaga 15300
tcgtgatcga gagcaaattt gagaacctca gaccttgcattt atttgaaagt agcagatgaa 15360
45 caatagaatt gttttactt ttggaaatggt tccacaataa tcctagtcta gatttaagat 15420
accaatattt aagtgtttagt tttgcattgtc tcttcagctg ctccacccgc gtggagtgtat 15480
50 tattagctta ttagcgcctt ctcattataa cgcctccag ttccagcctc tcaaaagtaa 15540
tatgctggaa tgatagaggt aattggctaa tggcctcaag gcaaccctgc agatagtgaa 15600
gcaaaaagcaa taaatattca atattcacac ataaatttgcac atacggagta ctccgtactc 15660
55 cgtttaagat cgggcatagt attggatgtat gtttagaatat atcttggcaa ggtgacatata 15720
acaatgtact ccgtatgtt gtagtgcata atggctttgtt ggagctgaag atgcgggtat 15780

ttctttctt gatgcacat caagtccgga aaattgatga aaatctacga gtacctcgag 15840
5 ggtatgtttt ccctgcacag atcatgacat acatataaac tattgttcca cttgcattag 15900
cgggagtctt gcaagagcaa gtctatgtat tccctacatg gtcgaggagg taagttcggg 15960
ctgaaaaata cgatgcagca tacactaccc ttacaactag ctgtttatc agaaaaagca 16020
10 aatagaaaatt agggcacaat ttactctta ctgccaaccc cccgtcgtaa cccttgctgc 16080
tagcattgtat tggctgtcag tcgtacaacg aagaaacgac actgtctgtt attatattct 16140
15 attccatcac aaacgttagcc cggagtgcgg ttcccagagt cttgtcttg tacaccgtgc 16200
ttgtcttagc attttcatta tgatcgagct caaagatgtt tcgtatggggg ctgtattgt 16260
gacatgcgtc cttgtgtttt cagggctata tctcattcga ttgacgttat caagcgacca 16320
20 attggacaag tttccttagca tcaatcctcg gaagccctgg gaaatcgtaa atgtcttcgc 16380
ccaaagaaga tttcaacagg atggccctag gtatctggaa gctgggtatg caaagggtgt 16440
25 ttccataagc aactgctcca aaaggcgaat aaggctgaaa gttactacag tccccatct 16500
tttagcgttgtt caccgacccg gggccaaaat tagtggtttgc ggggcatttc atcgaggaat 16560
tcaaggatga aaagctgttgc gaccattatc ggtcaatgtat cgaggtttgt acgacgttag 16620
30 tgattatgaa agagcaagcg cttacttgtt caaggacttc atggcagagg tacctggttt 16680
tgagtcgtatg ttcctggggaa atctacacaa tacggtaactt cgcgatgtga tttctgtcat 16740
35 cactcgcgaa ctaggtaaat attctttctt tttgactgtc cggttatccg ctgagttcta 16800
atttataga acaactgcta gcacccctctt cggatgaagt atcagcggct ctggtagata 16860
cttggacgga ctcaccagggt gggtaaaagc acacttccca atagaaatca ggagggaaata 16920
40 aaaactaata tcaatataga ctggcatgag gtagcactgc ttccaagcat gctgggcttg 16980
atcgcaaaagg tttcatctctt cgtcttcgtt ggtgaaccgt tttgtccgcac cccagtcgg 17040
45 ttggagacag tgatcaactt caccctcattt cgacacaacg caatcttagc cctccaccag 17100
tgcctctgtt tacttcggcc cgtccttcac tgggttcttc caccatgcac gaaactccga 17160
cgagagatca gaaactgcacg gacactgatc gactctgttc tggaaaaatc aagaaaagaat 17220
50 ccgcaccccg agaaattttc cagcggtgcc tgggttgatg cttttgcacaa aggcaacaag 17280
tataatgcacg ccatggtgca gttaaagactg gcaaatgcgtt ccatccactc cagcgccgat 17340
55 ctcctggta agatcttat caatctatgc gagcagccag aattgattcg ggacctccgg 17400
gacgagatca tctctgttctt tggggagaat ggatggcgat cctcgacact gaaccaatta 17460

	aagctccttg atagtgttct gaaggagagc cagcggttgc atccagtcac aaccggatg	17520
	catcgctggc ttttcaaaact gcgtgcccag tgcataatgct gaccatttac ttttaggagca	17580
5	ttttcgcgct ttactcgca agatatcaag ttgaccaatg gcactgagat tccttcagga	17640
	acacccatta tggtcactaa tgatgtcgcc gggatgcc a gatctatga t gatcccgat	17700
10	gtcttcgatg ggtatcggtt cttcagaatg cgtgaaggag ccgataaggc cccggcacca	17760
	ttcacaacga cgggccaaaaa tcaccttggg tttgggtacg ggaagtatgc ttgtcctggt	17820
	cgattctttg ctgctaccga gattaagata ggcctctgcc atatgttggt gaagtatgaa	17880
15	tggaggctag taaaggacag gccgcattggg atagttacaa gcggggttgc agcattccgt	17940
	gacccacgag caagcataga agtccgcaga cgcgcggtgg cgggagaaga gctcgaggta	18000
20	ttgactggaa agaagtgtac tagggaaaat tacgaactca tagtatgagc aaccataccc	18060
	aaaacaaaga gacttaccaa ccccatcatc aaggtagact ggggatttg actatgtcga	18120
	tgtaaatcggtt ccaacagcct tatttaggata tataaattat acgcttctca ggctttaaag	18180
25	catcacccag cacgataatt tctctggatt attgcaaaac caagaaattc tctgatccac	18240
	agctgtatac tccgtactcc gttcatcatc ttacagtcat gcagagggtg aaagggtca	18300
	gtgtgtgacg gtatttcggt atctcgccctc gtaatttgac agatccagcg ttaaaccagg	18360
30	cccaagactt gagtagacta tatttattct ctttgatatac catctcagca tcaagttttt	18420
	gacgttgtat tactatccctc gtttggaaatt ctcctccag gtcttgcctc attgcttata	18480
35	gcattctacc aaaaacgtca ctgtcatggc cgggtggtca gacatatacat cagcgctgc	18540
	cggatacaag gatgttgttt ggatagcaga tcgggcctcg ctggccaaag gattgggatg	18600
40	gtcaatcaac tacctggcca tggatatacca atcgcgcacaa gaccgcacat acggcatggc	18660
	cattttgcca ctatgttgcac actttgcgtg ggaattcgatc tacactgtca tctatccctc	18720
	tcaaaatccc ttcgagagag ctgtcctcac aacatggatg gtcctgaacc tctacctcat	18780
45	gtacactacc atcaaattcg ctcccaacga atggcagcac gccccgcctcg tccagcgaat	18840
	tcttccagtg atattccctg tggcaatcgcc ggcatttacg gcggggcatc tcgccttggc	18900
	tgcgacagtg ggagtggcca aggcagtcac ctggagcgcc tttctgtgtt ttgagctatt	18960
50	gactgcccgtt ggcgtgtgcc agctcatgag tcggggatct agcagagggg cgtcgatatac	19020
	aatctggat gttcttttg cttgtggat cttgttggg tttattggct aatgtgaatt	19080
55	gtggttggca gggctcaag atttctggc tcgtatatacg gtatgtatctt tatgcatgtt	19140
	cqaaqaccc actggccgca qgagtttgcac tggatcagct accctttcggtt ggcgtggcat	19200

ggcatcatgt gcttcctcgct ggatatttct tatgtggct tactgtggta cattcgctgg 19260
5 caggagcgcc agggccaatt gaagaaagct atgtgatcga caggaccatg catgatggag 19320
gtccgcacta acctcaactg tactttgtac aggtctgagt gctatatgac gatagtcaca 19380
aaacagagtt ggaggttatt tgcgcacatt gactaaaaat gggagagctg atggatata 19440
10 gcaaggggga tcaggtctcg atctgatcgt gccgatcgac aagaacaatg ctgtctgg 19500
gcgggtccaa ttgtctagcc tagaagtcta aatttcaatt ttcttcggac ttttacata 19560
15 gtaactactg cctaggactc gggatatgaa gtataatggc gagaaatggc tggctgcagg 19620
ggacatacag gtgataattt gcctcgatc tggcagctag ttacgtcaat atcttggtag 19680
taaacaccag ttgttagatct ttgcgtatata atgaaactca aaagcatttgc tgcgtactcc 19740
20 gtaattacct tcccaacccc tccagtgcca ttgaaaccat gaaggtcatc attgtcggag 19800
ggtcacatcgc gggtctcgcc ctgcgcatt gcttggacaa ggccaaacatt gactatgtca 19860
ttctagaaaa gaagaaagaa attgcccccc aggaagggtgc ttccattggat atcatgccta 19920
25 atgggtcg gatcctggaa cagcttgggt tatacgacca gatcgaggag ctgatcgac 19980
ctttggtag ggcgcatgta acttaccccg acggcttcaa ctatacaagt cgataccctg 20040
30 cactcataca gcagcggtgc gtcaatataa gcttctact ttctgatttgc aaactaatgc 20100
gagaggtctt aggtttggct atccacttgc attcttggat cgacagaagt tactgcaa 20160
tctggcaact cagccggtcc aatccagccg agtggaaacta gaccacaagg ttgagagcat 20220
35 tgaggtctcc ccatgtggcg tcacggtgat aacaagcaac ggacacacccat atcaggcgaa 20280
tcttgcgtc ggggctgatg gagtgcatag tcgggtacga gggagatgt ggcgactggc 20340
40 agatgcctcg caggggaacg tatgtggaaa tggagacaaa ggtacacattt ttcctactgt 20400
tttgcctat cctcgctttt tttttcttg gccaagtgtt ttgactttga gctggaaagc 20460
taatataatttgc atttatagca ttacgatca actatgcctg catcttggat attcgtcac 20520
45 acgtcgatca attggacccct ggcgagcaaa taacctgtta caatgtgggg tggagtatcc 20580
ttagtgtgat cggacagaat ggcaggatct actggttccct ttttatcaag ctggaaaaag 20640
50 aattcgttta tcatggatca cacaaaaccc agtccactt tagccgtgaa gacgcccgg 20700
ctcattgcga gaggctggcg caggagcctc tctggaaaga tggacattt ggtcagggtct 20760
gggctcgatg tgaggtcttt caaatgacac ctttggaaaga aggggtgctt ggcaaatggc 20820
55 actggagaaa cattatctgc atcgagaca gcatgcataa ggtcagcagc tcattatcac 20880

tcctggctta ctgacttttgc taattaatttgc acattctcat gcagttcgca ccgcattttgc 20940
gacagggtgc taattgcgt atcgaggatg cagctcagct cagcaatagt ttgcacactt 21000
5 ggctgagcgg atctggaaag gaggcatcaac taaaaaccga tgatttgaca gagattctgg 21060
ctcaatttgc acaaactcgc ctccagaggc taggtccgac gccatggcc gctcgatctg 21120
10 ctatgcgtct gcatgcgcgg gaagggtca aaaactggat actgggacgc tacttcttgc 21180
cctacgctgg tgacaagccg gcccactggg cctcccgagg aatcgcaggt gggaaatactt 21240
tggacttcgt agagcctccc acgcgggctg gtcctggctg gattcagttc agccagtcgg 21300
15 gtaaaaggac ttctttccc atggcagtgg caggtctgtg cctagtgagc attgtggccc 21360
gaatcatgtt tttgaaatattt gttgcataaga gaggcccacc atatctggag tacttcatac 21420
20 agagtgtttt atgggacaat ataaacttta gggcaattta gcgcatttgc atagatcatc 21480
tgcatactag taaggcaacc ctgaagggtga tcgacacgat ctgaaaaat caatatcgtg 21540
cttcgttacg gagtattgtt ttctacatgt catagtgcgc gctgccccag tggggctatg 21600
25 cagaaaagtga ttctcgatgtt ttgtacttta cagtgtatgtg gtccagcatg tcagccatttgc 21660
ctctagtgcg tgcgtgtact gaccacatcg cggccatttc cattttatcta gggtcctgtc 21720
30 gcctcaaaag cttgtggtca caaatcgttt gatccttcga gatcatactg aatttttgc 21780
caatctgtca tcatggctgg ctctcgtct acggcgcagt tggctcgct tctcattgtat 21840
atctcccgat ttgacaaata caactgtttt tttgtatata tccctggagg tacggagtag 21900
35 tgcagaccac ttcaacatcca taccaccgcg ctcacaattt catatagtct ggtctatctt 21960
ccttgcagca gcctcacgac acgctgtatgg cgaccccgtc cctctggact ttgtattggg 22020
40 ccgcgcagga ctggccttca tgtacacgtt tatgtgtatgc ggccgcaggaa tggtatggaa 22080
cgactggatc gaccgcgata tcgatgccc ggtggccgtt accaagaatc ggcccctcgc 22140
ctccggtcgg ctttccacca gagctgcctt catttggatg cttgtccagt acgcagcctc 22200
45 ggtctggctg atggaccgcg tgggtgagcgg gcaggatgtg tacgtctttt ttcctcctcg 22260
taccccaaac aattattctg ttgattgaaa actgacccta atcattctcc agatggacat 22320
50 acatgcttcc tctcacaacc gggattatct tttatccctt cggcaagcga ccgacaagtc 22380
gcaagctggg cgtctatccg caatacatcc tcggtgcaag cagcgcctt actatcctcc 22440
cagcctgggc ctccgtctac acaggccgtt tatctttgaa ggatctgggt atgcgggtgc 22500
55 tcccgcttttgc tctcttcctg tttctgttggaa ccatctactt caacaccgc ttcacgtatc 22560
aggatattaa ggatgactgt aagctgaatg tgaattcgatc gtacgtcctc gcggggagcc 22620

atgtgcgtgg aatgcttctg ttacaggcta ttgctgtggc gctgggtatc ccctggattc 22680
5 tctacaccag cgccctccact tggctctggg tctcatggct gggggatgg acggcatctc 22740
tcggcgagca gctttatctc tttgatgtga aggatccgag tagcggtgga aaggttcatc 22800
ggcggaaattt cgcactgggg atttggaatg tgctggcctg ctttggtagag ctgctatatg 22860
10 ctteaggctc tctgtgaatg atgttaatac gatgtggtcc ggttggact tggggagtag 22920
agtctgagag gcttaaaatg ggttaatggt gcgatgttgg cacagtgtga actattcata 22980
15 aatctttgct acgaagttgg gcttcacett tcaattgaga agttgttact ggaattttc 23040
gacactcaaa attcgaagag acttgttatta ttagagggat atagcctatg tcttccaatt 23100
ggtgtagaat cccaaactacg agaccgcttc agaacgttgg agcacaagga tagaaagttc 23160
20 acctattcga aattctctac tgtcgtacat atgctatgtt catgttactc ctttgcttgc 23220
gcacctatacg cccagcaaaa caagggatcc tttgctaaca ggagctgtatc atcacggttc 23280
agagtcagat gcaaataccca cggctccgtt ctcgcacat catcctgacc ctttggaaagg 23340
25 ataaagcaca tccccctaa gacaggcaaa tggatgttgg accctcgagg ttgcgttcca 23400
aggctctccc caaagtccag tccgaagatt tcaaaattcc taaagctgtt cagagggata 23460
30 ggaactcccc gaaatccat atccgcccag tcaggctgcg agtggagatg ggatagagcg 23520
tcttgaatat attctgcgtc aaccgcacaa agactctggc gtatacgagc tgcaatttgc 23580
35 gtgagatcct ccagacactc ctggcgcaat tccacccaaag gatctgtgcc atcaactagg 23640
gcctcgtttc tcccagcttgc aattggcgta tatgtcaata gcaccatgtt tcccagatag 23700
tcatcaaagg ctggagttt gaaattccca cgcatatcca ccgcgttgc cagttcggtt 23760
40 gatttaccag ccaattgtcc cgcttgcga agtacatgtt cccaaaggcc gctcacgtt 23820
tcgttactgg acaggaaccc tggactcggtt ctaccatcgat cctggaaaga cgtttgcctt 23880
45 ttgatcaacg tattgcaacg ctccttcaaa tactcgatct taggaccggg gattttcgtt 23940
cgccaggtga caagctcggtt ggctctcgcc cggacgaacg ccgaccaatt ctttgcgtt 24000
agtgtgtccc agtctccgag gccacagtag tgcttgcata aatccatcct ggaaagaccg 24060
50 gagctgtttt ctgggacaag acgctcaatc tccgatcgta actgcccgtc tggcgacaca 24120
cttgcagaag acatcgccgtt cgggtctctg cagcaatcggtt ctagaaggcc caagactcgc 24180
55 gcagcgcctt caccatccat tgcggaaatgtt tgaaacgtca tggcgagaat gatccatcg 24240
cgcatgacat ttgtttgtttt tcgttaggatc ggccttcgtt gcaacgaaat atccatgtcg 24300

ataggcaatg gcgccagccg acttatgatt tcctgctcct cagtgccgt taggaggcat 24360
tttgatttggaa ttcccttcaa tgactcgccc tggtagtgcc gtatccggag tataggaaac 24420
5 tggacaagcg actctgagggc ttctggttcg atttgccagg tgtacttcgt ttggctggac 24480
tctgtccgcc gagtcacgtc ccctgcgagg aagggggtgta cttcaatag cagctcgatg 24540
10 ccattctcgaa gaacaccaat gctcttctca gttgcgtgg tctggaaaaa cagcagaaag 24600
gtgacgttca ttccgagggg attgtggtcg agagaagata aagggttaagc agagcggtcg 24660
ccagttcttc gggcatcgca cattccatct tcacatagac cgtggagtct cacaggtccc 24720
15 tctttgacct gatctcttg actgactggg agacatactt cctgggtgct catgatttct 24780
gggtgttatac ctattgagtt gagttgtgtc ttgatcttt ttttatttt ttttgattt 24840
20 ctgaccttgt ttgccttata ttggactttg cttttcttg tatattgtat tgcattaccg 24900
tacaacaaag catgggatc tctgtgttct gcatgattgt ggagcgtatt ttccctcgatt 24960
tggtatacaa tcaggtcgat ccctggcgga ttccggatct gatgcgttgc tacaggtcat 25020
25 atatctgctt tcctcggtat ttttgagctg aatatcacta tatactgttt ggagacgatc 25080
aatcgcaaga gagggttagt gattaaatca gttagtctca tccatagtgg gcattagac 25140
30 caataaaaga tggtttccac cttgagatgt gatcgccaca agaagatttt gtaaatagta 25200
tgtatttcc aggcctgtat ttctatctgc atatttgta gcttgcata cgaggtacat 25260
cttactgctt ttagatactg acagcagcaa aactcccggt tgaaggacga gctttgacac 25320
35 aaggtcaggc acttctctag tacacaaatc ctaatcatcc gacgacatac tactccgtat 25380
gctgtacata gagatccatg tccaaattttt gagtctgccc ctctttgatc cacagtccag 25440
ctcagccagg cgcaatctgc atgcattggc atggaagcta ggagctgaca ttggctggaa 25500
40 ctacgccatc tggggcacaa tgcaagctag gcaactgacc atgtactggg tcagtttga 25560
ttgagttatgc tatacggaag aaagcgacta gtactccgtt gtttgcgttata ctacctgcaa 25620
45 gtggaaagag ataccttagat aggtgacatt agtgtccgaa ccaatgacca atggccctta 25680
tgcacccata tcccttacat ctttcagaaa gagaaaagcc acaagtataat catgtactcc 25740
50 gtactccgtt caacggaaatt acttgatctc tatattacat tcttcctgaa gaccgtttct 25800
cgctattgtc agttacacac acaatggatt ccctattgac gagcccgta tggctcaaaa 25860
ttgcacatga gctagcactt tacctctttt ttattgtgcc aaccgccttt ctcatacataa 25920
55 caactcaaaa atcatccatt attcgatggg cctggacacc atgtctgtt tatactctgt 25980
accaattctc tcttcgggtt ccctctctgt cgacaagtca attttgaag ggcgttgcag 26040

tcattgcttc cctgattcgt atgagtcttt aattttctgg ttgcttgact ctgaccgcgg 27780
tcactagatt gcccaccatg tgcgttacta gaacctttcc ccgattcttt gctgcagcta 27840
5 acactataca gggcaaagct cgtggacgac catcagatcc atactgcctc gttgcataac 27900
ccgattcctt ggcaattgca tacatacgtc tggccttcc tgcgttacta gcccgtgttc 27960
10 tttgcctttt acctctctcc cgagcgctat gatacctaca ttcagggaca ggagtggacc 28020
tttgtgtttg cggggtctat catcacagtc cagtcgtct tctggctgat gaccaagtgg 28080
aacatcgata ttaacaccct attcacaact actcgatcca aatccatcga cactgcccgg 28140
15 cttatcaaag tggttccgat caccaatgcc ggctctgccc agatctgtaa cctgattaga 28200
gagcacattg gcccgaagaa gaccctttcg ttccctttcc agaagcgccc cttcctcttt 28260
20 taccggaga ctgcgtcctt cgccacccctt tcttacgccc tcgacgcccga gccgaagccg 28320
gcctcaaga ctttccagca gagcgagggc ttcacgtcga aggccgagat tgagcgcgtc 28380
caaaaccact atggtgacaa taccttcgtat attcccgatcc ccgggttcat tgagctcttc 28440
25 caggagcatg ccgtcgccgc gttcttcgtc ttccagatct tctgtgttgg attgtggatg 28500
ttggatgaat actggtaacta ctgcgtcttc accctcttca tgctcgtgat gtttggatgt 28560
30 accgttgtgt ggcagcgcca gaggacattg agcgagttcc gtggggatgag catcaagcct 28620
tacgatgtct gggtataccg tgaacggaaa tggcaggaga tcaccagtga taagcttctt 28680
cccggtgatc tcatgtcggt gaaccgcacc aaggaggaca gcgggttgc ttgtgatatt 28740
35 cttctggttg aaggcagtgt cattgtcaac gaggctatgc tttctggcga gagcaccct 28800
cttctgaaag actctatcca gtcggcgttcc ggcgtatgact tgattgagcc agatggattg 28860
40 gataagctct cgtttgtgca tggaggtacc aaagtcctcc aggttactca ccctaattcg 28920
actggcgacg cgggcttgaa gaacttggcc agcaacgtta ccatgcctcc agacaatgg 28980
gccttgggtg tgggtgtgaa gaccggtttc gaaaccagcc agggtagcct cgtccgtact 29040
45 atgatctact cgactgaacg tgtctctgcc aacaatgtt aagctctgct gttcattctc 29100
ttccttttga ttttcgtccat tgcgttccgt tggtacgtgt ggcaagaagg tgtgattcgg 29160
50 gatcgcaaac gtcggcgttcc gtcgttccgttcc ttttcgttccgttcc ttttcgttccgttcc 29220
cccgaaattgc ctatggaaact cagcttggcc gtcaacacta gtcttgcgtc tctgagcaag 29280
tatgccattt tctgcactga gccattccgt atccccttg ctgggtgtgt tgatatcgct 29340
55 tgcttcgata agactggtaac cctgaccgga gaggatcttgc tgcgttgcgttgc 29400
ctcactttgg gtgaggctgg ttcaaaggtaa gaagctgtatg gtgctcacac cgagttggcc 29460

aattttctg ctgctggacc cgacaccact ctcgttctcg ccagtgctca tgccttggtg 29520
5 aaattggatg agggtgaagt cgtcggtgac cccatggaga aggctacttt ggaatggctt 29580
ggctggactc tgggcaagaa cgacactttg tcttccaagg gcaacgctcc cgttgttct 29640
ggtcgccagcg ttgagtcgt tcaaatacg agaagattcc agttctcctc ggccctgaag 29700
10 cgtcagagca ctatcgac cattacgacc aatgaccgca atgcttccaa gaagaccaag 29760
tctacttttgc tgggtgtcaa ggggtcccccc gagaccatca acactatgct ggtcaacaca 29820
15 cctcccaact acgaggagac ctacaaggcac ttcacccgta acggtgctcg tgtgcttgc 29880
cttgcattaca agtacccccc ttccggagacc gagcttccc agagccgtgt gaacaattat 29940
gtccgcgaag agatcgaatc cgaactgatt tttgcgggtt tccttgcct gcagtgcgg 30000
20 ctgaaggacg atgccatcaa gtctgtccaa atgttaaatg aaagcagtca ccgtgttgc 30060
atgatcaccg gtgataaccc attgactgct gtccacgtcg cacgcaaggt tgaaattgtt 30120
25 gaccgtgagg ttctcattct tcatgcggccca gaacatgaca actctggAAC caagattgtc 30180
tggcgtagca ttgacgataa gctcaacctt gaagtcgacc ccactaagcc tcttgatcct 30240
gaaatcttga agactaagga tatttgcgtt actggatatg ccttggcaaa gttcaaggc 30300
30 cagaaggctc tccctgatct gctccgtcac acctgggtt acgctcggtt ctctccaaag 30360
cagaaggaag agattctcct tgggtttaaa gatgctggat acaccactct gatgtgcgg 30420
35 gatggAACca acgatgttgg tgctctgaag caggcccacg tcgggtgcgc gcttctgaac 30480
ggctcgcaag aggtctcac caagatcgct gaacactacc ggaacactaa gatgaaggag 30540
ctgtacgaga agcaggtcag catgtgcaaa agatttAACcc tccagtaccc 30600
40 gttctgatcg ctcacctgta tccccccggc cctaccaacc cacactacga gaaagcgatg 30660
gagagagagt cgccagcgaa ggggtgtcgat atcaccgctc ccggcagcac tcccgaagct 30720
45 attccgacta tcacatcccc tggcgacacg gcccgtgcac aatcgaaactt gaaccccccag 30780
cagcagaaaa agcagcaggc ccaggcagct gcagctggcc ttgcagacaa gtcacatcg 30840
tctatgtgg aacaggagct ggatgacagc gagccccca cttatcaagct ggggtgtgca 30900
50 tccgtcgctg ctcccttcac tagcaagttg gccaacgtta ttgttatccc gaatattatc 30960
cgtcaaggcgtt gttgcacccct ggtcgacact attcagatgt ataaaatcct cgcttgcac 31020
55 tgcttgcata gtcctacacg tcttagtgc atctacctgg atggatcaaa gtttgggtat 31080
ggacaggtca ctatcagcgg tatgctgtatg agtgcgtcttgc tcccttcaat ttcccgcc 31140

aaggtagtc gtatcccgtgtcgaccaa atgatttgcataatgttac tgtgtgaagt 31200
ctgtcgaggg tctgtccaag gaacgcccgc aacccaatat tttcaacgtc tacatcattg 31260
5 gatctgttct tggacagttt gccatccaca ttgcgactct gatctacctt tccaaactatg 31320
tctataagca cgagccgtac gtgatgaaaa cttcccttt catttgcctt acttcatagc 31380
10 taacataatc aacaggagag attctgatat tgatctcgag ggcgagtttgcgacttccct 31440
tctgaacagt gccatctacc tcctccagct gattcagcaa atctccacct tctcgattaa 31500
ctaccaaggc cgtcccttcc gtgagtcaat ccgcgagaac aagggcatgt actggggcct 31560
15 cattgcgcgc tccgggtgtcg cattctctcg cgccactgaa ttcattcccg agctgaatga 31620
gaagttgcgc ctcgtccct tcaccaacga atttaaggtg acattgactg tgctgatgat 31680
20 ctgcactac ggtggctgtt ggtttagtga gaacgtccctc aagcacctgt tcagtgactt 31740
ccgtcccaag gacattgcca ttcgtcgccc tgaccagctc aagcgggagg cggaacggaa 31800
gttgcaagag caagtcgacg ctgaggccca gaaggagctg caaaggaagg tctagaggtt 31860
25 ggtggtttga agatttgat ctgtaaacat agagaggagg ttgttgaatt ttagaaatgt 31920
tcaagtggtg tgtgacattt aatacattt ttttggctt ttattgaagc attcttggaa 31980
30 actatatgtt gaaacaaatt cgtatagttt aatggctcct actctgtact gtccaaatcgt 32040
cgtgaggcca ggtattgcct tggtagagaa cagtgttagac tcaaatgtgg cgatcgccg 32100
atcagcttgt tacgaggtt gggctcgaaa tgatcgcccc accataactt cttgtagctc 32160
35 cttgtttgag aggatgcagt ctacccgtta tggtagaccta attatccagg atggtcgaga 32220
atacttctca atacacaggg ttagacccca gatatatgtat atgtcacctc agagagggc 32280
aaagactggg taattccaaa aaatgtgatt ttgcagaggg tcaaagctat atcggatact 32340
40 gcttcttttc tctgcctcat agtgaaggaa acactatatt cttcatggta ggcaaagagg 32400
taaaagtgtt cgtgccccaa ttcggtagaa ggataggccc tggggaaaa ttccacattt 32460
45 tgaccgatatactatagaa acatatgtttagccgttgcgttccatttgcgttccatttgcgtt 32520
tcgagctgac gtggacttca aatgcaggat gctttgttcc tttgtactgc catgcaatata 32580
aatgttgcct tgaactcgag attataatgc gaaaacctcg tagagccgat cgccggccgaa 32640
50 gccaacattt ttctataata cataggtaaa cgatctgtga attcagaaag ctcccacattt 32700
gtattataag catgaatcat tcaacgcgag acttcaagct tcatgaaatc cttcaggaac 32760
55 ccaacagttt gaaagaccacc aattcccttag atcccactga tttcgattac gacattccgg 32820
attgttagtag ggcataatggc gatgcgggt ttgattgcaa agaataatatt cccatacattt 32880

5	gcagtaccca ccttgacaa tccaggattc aggtgcgtcg acgcccgcgt gattacaact	32940
	agtgctggag ggcgcgttgt catagaacca tttgcaggag ttaccgtcaa ttgattcggt	33000
	ggtagactt cggctggc gcaggcattt gaggtccttc gtccagccca cattatgtcc	33060
	tttcatttgg atgtagttgc cttgcagtt cttcttggtg tacatttcca ctgtccatgg	33120
10	atggacatata ggcgcatcgc tgggtgttgt gcttggtttc gtgttggatg ttttggttga	33180
	cttgaccgat gtagtttgc tagtggttgt ggcccttggtg gttgtggtgc ttgacgcage	33240
15	actggccgat atcgtttat tggatgggaa caaggcaagt ctctgaatag tcataaattt	33300
	tgcaaaagca ttaaccgcata tagtaggcata ggcagtcattt tttgaggtgg tcgctccaaat	33360
	ggtgcggttt gtagaagtac agaattccga agtgcattcca caataaccat acgaagagca	33420
20	ctatgaattt attagatgtt gaagagcagg atttagagcc tcttgactca cacatttatt	33480
	agctgcacat ggattcaaag aacccagatc agaccagttt ctagggcgct ttgtcccagg	33540
25	cacccggggg ccacaaacag cattggaaag cgtagctggc atcatgggtt ctccagaact	33600
	gaggcatata acagcaccctt gcactaaatg gtcacatccca agccattccc aggtttgagc	33660
	attgttaggtc tcgatatacag ccaaggcgag agagtgttct tgagcaatcg tggtgcaggt	33720
30	ttcgcccccc tggacaacat atttgtgaca aagaccatca ctacctgctt gaggtgcacc	33780
	tgatattgct gtcggactgg cccaaaggac gcctatgaga gtcacgaagt tgcttggtcg	33840
35	gaagcaccac atcatattga ctgatggaga gtagttgctt tgcttgcattt tattttgcaa	33900
	ggcagggtttc atcttatct gttcaaagag gaaaacatgt gccaaactgc caaggataga	33960
	tgcacgcattt aatatgacat tgccggggag gggcaaatgt ttgtgaaaga actaggatac	34020
40	tgtgccaggg ccattagcat agtattgaag caaattatag aatggcactg cataaaaatg	34080
	tggaatcctc gaatttttctt tttgtcttctt aacgccttagt gcatgtctt ccaggttgc	34140
45	cttgaaggct ttgtctggc tcccagaaat ggaaggactc aagggtatgt atacagcttc	34200
	taaaacgtaa atgattcacc cgagaaagga attcataatc cgaggaaggt cagacacata	34260
	aggctgtctc gaaaccctt gaatgccccaa ggaagaaagg aaattcctac ggctgggtca	34320
50	gactagcaag aaaacgtcac ttgacttctg agatccactc agatagcaga agaacgtttt	34380
	tggtagttt cgttcttgcgaaatgcata gaccagatga ttcgaggaat cttcttgcata	34440
	gcacccttaa tccaaatctt ctgttagacca agcactcgcc tattgataact gtttcgagag	34500
55	tctgtaaatgcata tctgatcacatcgtggaa tggaaacatttgcggctttc	34560

gaatgacatt gggttgacta acgaaggccc cttcacgcag tgacgaggcc ccaaagttca 34620
aggccaaacgc gcaaagcggg accaacatcg aactccccat tctcggggag ctgagggccc 34680
5 gccttgattt tgacatcttc ccatttgtca aagtcattat tgaacgcctg cgtcatttcc 34740
gaaggattca gaatgcgggc caaccggaag tgacaatcta ggagatcagc acttggcaga 34800
cttcgtacat tgtctgcgtc cttgttgctg aaaaccacct ctatcgaaa aaagaccttg 34860
10 taatttgaag aaattccttc gaacgtgtga actttgtatt tattgtccac atccttgaa 34920
atttcggtta gcaaggtaat aaagtaaaat cattccacaa atgggagcgc actcactgtg 34980
15 gtttcaaagg caagacggaa cgcgcgaag tgttcatgta cccagctgt aagggtgaga 35040
ccgttggata agctgttgc ctcttggtgc ctgaaattca tttccggat cctggaaaaa 35100
20 cagcgccaga ggacctccca agtcgtgac gcatttgta tgagatcggtt cttgagcccg 35160
tgagtcatttgc atcttaatga ggatccgaga acttactcgc cggggatccc agactgcata 35220
tgacaaggaa attatgtgtg cgcttctgt attcccgagt tttcaatat cctctgattt 35280
25 gcctagactg tcccaccgat cgagatccat gtcaccagtg acaacacagc agtaaccgtc 35340
gcgttttagt aaaccttcct tgaactcccc tgcggggac ataggctggt ccaaagtgg 35400
30 atttttcttc gcatggaaat cctgccgttc gtttgcggat atagttactg atgggggctt 35460
agaccgagaa cgcactattt cgagtgaacg tataagttagt ggcttgcgtac tttttgtta 35520
acttactacg gtttcttgc ccagtgtaca gtttgcggaaat gtattgataa attccatcat 35580
35 cgtctccag gctgatgata tccttcgcaa tggagctttt gccgcgtatct gggagggaaat 35640
tcaaaaaagc ttgtaatgtat ttttgcggaaat tgcggggatctt ttttccacgg ctcttgggtt 35700
40 tgtaactttc tattcttaag cgtcggtttt caatatcctg cttttgttgc gtttccgtatg 35760
cttcagtaga ttgtgaaaac agcgattgtg cgcggggata ggcacgtgtt ggaccagaaa 35820
cttcgtcttc ttgggtttgc ttttgcgtt ttttgcgtt ggtgggtgggt cgatctgattt 35880
45 caggcaattc agaacggcgc ctttaagat tagttttggaa tgggtggcgac gaggaagttt 35940
gagggatgtt ttgggttcggg gcatcggtttt gtgtgtttt ggtgggtgggtt gatagctccg 36000
50 tctttttctt ccccatgtat tctgtatgtt gtgtgtttt ttttccatgtt 36060
actgagatgg tagtttgcgtt ggtgggtttt ggtgggtgggtt gatagctccg 36120
atgagaacaa tttttttttt ggggggggggg ggtgtgtttt gatagctccg 36180
55 agatattcctt ccggcagaat cgtccaccg aaaaacagtc cggccggacgg gtcattttttttt 36240
ttttggagaa aatgttatttgc tagttacaga aaggcatttttgc cccacagaac aagaattttttttt 36300

ccatatttca ttgtttcca tcaagcaatt actcgccaa tcgtctctcg gagggtgcag 36360
5 agaataggct ctctctggaa ggccgctgga aaaagtggga aaaggataca ttctgtggcc 36420
acaggcgtgg gacagggttt cccctgaca ctgggggaga aatgtggaaa tgtggggaa 36480
ctctgcggag acggaagaac aaaaggcggt caactgctgc ctccacgtga tgtcacgtgg 36540
10 agcttagccg tccagcttgg aagataaccc tagaggaata tgagcatatt ctacggagaa 36600
ctactccgta caacatacgg agtactcata caactctgta gcaacccctg atgtgatctg 36660
15 tatttgaagt gtggacctga taccgactgc tcctcaaacc ccttaaaccct gtatcgagta 36720
ctccgtaata tgtacaccgt tcactgactc acattgatta atcacattag atctctcggt 36780
ttcatgtacg tggatcatta tgagttcgag cattgaatat aagctaaaac cataccccc 36840
20 gaccctaagg ggccttctgg aaagaaaaat ctgtctttt gcaaatcaaa atatataatag 36900
agttgtttac ccgaactgtc gggttatgca tcttcaggcc tgtggagctg tgtcatcatt 36960
25 ttgttactcc cccttatcta ccgcaggatc gccaaaatgc cttagcgagac tgctacaggt 37020
gactttggtc cagcgccgccc tgggatagac ttgacagaga accaaaactgg cgacttgcta 37080
ggagcagtga ttctgttagc ggtggtcgag acgactgcgg tggatattgcg gacgattgcg 37140
30 ccgacgagga tcaaagagat ccgacaaaaca gctattgatg actatctcat tggcgccg 37200
cttttattct cttggggAAC ggcaatatca tgcttcatca gtgagttgac catgaggcca 37260
35 aagccgatgg gcccagtact cacaacagac tctttaggca ttccatatgg caacggttat 37320
catttgcaat ctgtgacaaa agcagagttt aacactgttt ggaaagtaag gaatccaata 37380
ttaaatgaga tgcctggat agacgttgac cagacattca gatcctttc gcctatgtca 37440
40 tgatttacgc tacagccgtt acctgcacca aagcctcgat cgtcttattt tacggccgca 37500
tcttcactt tcgctggtca ctggccatct gcctgtttct ggtcggttggaa tattgggttg 37560
45 ccattattgt cacgggtggg atggcctgtc gaccactgcc acatttctgg ttggcttaca 37620
cagatccatc agcccttgggt gtctgcattt gatattccac gttcttttc gcaaatggca 37680
ttgctgcccattt ggcgattgtat gtgatcatac tggatgttgc gatgccagca atataccagt 37740
50 ctcagatgca gttgtcgcaa aaggtagcggtt tcgtgggtat cctactcttgc ggaagttgt 37800
atgtacctct gccccggggcc tcctacgaga aggactgttag ctaatttttc tcagtgttttgc 37860
55 cgtggcaagt atctgcccggaa tcatcgact tcagaatatc accgacggga cagatacgac 37920
gtgggctatc gccccagttct ttattttggtc gtccgtggaa ccattttgttgc ggatttttg 37980

128

cgcatgcctc	ccaacatgg	ggccttctt	tcggcaatgg	cggccatcg	ctcggacgct	38040	
ctcatcaact	gatggcagta	ccgatccaag	ctctgagcta	ccatctgaga	caacgacctg	38100	
5	gctccgaaga	tcccgaacca	aaaaacctgc	caaggactca	atattcagta	tcaatgattt	38160
	ttgctgtgtc	gatgaggtcc	aactaatgaa	cgatataat	gccactcggt	cgctggggga	38220
10	cgaggctgct	agtgaccatc	aggacgtgga	gggaggctgt	atcacagtc	aaaaagatgt	38280
	ggaagtgaca	tggccaaatgt	acaagtcaagg	aaaaaaaaat	gatctggct	tcaagtatca	38340
	taaaggggct	tgatcagtt	tgcaaataatt	tcgacttgac	acggactata	tttgcgtttt	38400
15	gtgtatattt	aataaaaata	gacgccactg	gcaatttgtt	attgataaag	gttaagtctt	38460
	ttccgtaatc	cataccccgt	actctataca	aagtactctg	tgctccgtac	ggagtacacg	38520
20	gaaacaaacg	gggatatagt	cgtggcacct	ttcccgtgtt	ggcggacttg	cccgtaacgt	38580
	aaacactccg	cagatccctt	ccaacacagt	acataatcct	gcagcgaaga	gcgatctgat	38640
	agacgctatg	tgccgtcg	acttgttatg	ccaattaacg	gtggcagaat	tgtggagcaa	38700
25	tctagcagag	gaaagtttcg	atgtgcattgc	cgagccctaa	aaagtcccag	tgcggagaat	38760
	gtagtaatcg	actggacatt	ccatgtactt	tgcacgctat	aacatatttc	tatgccatat	38820
30	acccctctgg	taatcatgtt	gatcctcttg	cttactgcgt	tggctcctt	gtatcgact	38880
	ttcccggtcg	cagcattata	agaggataga	gagaccgcat	gagagaatac	acaagagaaa	38940
	tcactaattc	actacgtat	cccccaattc	actcaacatg	tctcacatc	acacttccag	39000
35	attgcaaa						39008

<210>	267			
<211>	556			
40 <212>	PRT			
<213>	Penicillium coprobiuum PF1169			
<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	
Ala Phe Asn Ala Ile Gln Phe Arg Arg Leu Val Arg Ser Leu Ile Ala	35	40	45	
55 Gly Leu Lys Ala Arg Gly Val Glu Arg Gly Asp Cys Val Leu Val Gln				

2009274832 06 Jun 2011

C:\NRP\b6\BDCCKLLU667799_1.DOC (1/6/2011)

129

50 55 60

5 Leu Glu Asn Ser Val Leu His Ser Ala Leu Phe Phe Ala Ile Val Gly
65 70 75 80

10 Ala Gly Gly Val Tyr Met Gly Phe Asp Val Ala Ser Arg Pro His Glu
85 90 95

15 Val Ala His Leu Leu Arg Val Ala Glu Pro Arg Leu Ile Ile Thr Ala
100 105 110

20 Pro Ser Ala Leu Thr Arg Val Leu Glu Val Cys Asn Asn Gln Gly Met
115 120 125

25 Ser Ser Asn Gln Val Leu Leu Met Asp Glu Lys Ser Ile Glu Ser Val
130 135 140

30 Val Gln Phe Ala His Gly Gln Ala Glu Gln Thr Glu Asp Leu Asp Thr
145 150 155 160

35 Gln Thr Val Asp Gln Pro Ile Arg Leu Glu Ser Leu Leu Gln Tyr Gly
165 170 175

40 Glu Leu Asp Trp Leu Arg Phe Glu Asp Ser Glu Glu Ser Lys Ile Thr
180 185 190

45 Pro Ala Ala Met Phe Leu Thr Ser Gly Thr Ser Gly Leu Pro Lys Ala
195 200 205

50 Ala Ile Arg Thr His His Thr Ile Ile Ser His His Leu Ser Val Tyr
210 215 220

55 Tyr Glu Val Pro Tyr Pro Val Val Arg Leu Met Ala Leu Pro Leu Tyr
225 230 235 240

60 His Ser Phe Gly Asp Phe Trp Gly Asn Ile Phe Pro Ile Arg Tyr Gly
245 250 255

65 Gln Pro Leu Tyr Ile Ile Pro Arg Phe Glu Ile Thr Ala Leu Leu Asp
260 265 270

70 Gly Ile Arg Gln His His Ile Thr Glu Thr Tyr Met Val Pro Ala Met
275 280 285

06 Jun 2011

2009274832

130

Ile His Ile Leu Asn Arg Ser Ser Leu Asn Val Ala Glu Ser Leu Ser
290 295 300

5

Ser Leu Arg Tyr Ile Gly Ile Ser Gly Ala Pro Ile Asp Gly Tyr Ser
305 310 315 320

10

Met Gln Gln Phe Gln Ser Leu Leu Ser Pro Asp Ala Ile Ala Gly Asn
325 330 335

15

Leu Trp Gly Met Ser Glu Val Gly Val Val Phe Gln Asn Arg Tyr Gly
340 345 350

20 25

Ile Gln Pro Gln Phe Gly Ser Val Gly Thr Leu Leu Pro Arg Tyr Glu
355 360 365

Leu Arg Phe Val Asn Pro Asp Thr Gly Glu Asp Val Ala Gly Thr Pro
370 375 380

25

Asp Ser Pro Gly Glu Leu Tyr Val Arg Gly Pro Gly Leu Leu Ala
385 390 395 400

30

Tyr Lys Gly Arg Thr Asp Ala Lys Asp Glu Gln Gly Trp Phe Arg Thr
405 410 415

35

Gly Asp Met Phe His Val Glu Asp Gly Asn Tyr His Val Ile Gly Arg
420 425 430

40 45

Thr Lys Asp Leu Ile Lys Val Arg Gly Gln Val Thr Gln Tyr Ser Val
435 440 445

Ala Pro Ala Glu Ile Glu Gly Ile Leu Arg Lys Asp Pro Ser Ile Lys
450 455 460

45

Asp Ala Ala Val Ile Gly Val Met Leu Pro Asp Gly Ser Ser Glu Val
465 470 475 480

50

Pro Arg Ala Tyr Val Val Arg Asn Asp Thr Ser Pro Glu Thr Thr Ala
485 490 495

55

Asp Gln Val Ala Gly Leu Ile Gln Ser Gln Leu Ala Ser Tyr Lys Ala
500 505 510

06 Jun 2011

2009274832

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

15 <210> 268
<211> 2447
<212> PRT
<213> Penicillium coprobiuum PF1169

20 <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

40 Glu Ser Tyr Phe Leu Asp Glu Asp Ile Lys Ala Phe Asp Ala Ala Phe
65 70 75 80

45 Phe Asn Ile Ser Pro Thr Glu Ala Val Ala Met Asp Pro Gln Gln Arg
85 90 95

50 Leu Leu Leu Glu Thr Val Tyr Glu Ser Leu Asp Ala Ala Gly Leu Arg
100 105 110

55 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

C:\WRP\mBio\DC\7ULLU667299_1.DOC (3/16/2011)

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
225 230 235 240

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
305 310 315 320

55 Thr Gln Ala Gly Asp Pro Gln Glu Ala Ala Ala Ile Asn Ser Ser Phe
325 330 335

340 345 350

355 360 365

370 375 380

2009274832 06 Jun 2011

CWRPordbNDCCX1.LJ667299_1.DOC (106201)

133

Asn Leu Leu Met Gln His Leu Asn Pro Lys Ile Lys Pro Phe Ala Ala
385 390 395 400
5 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 Arg Val Asn
595 600 605

5 Glu Gly Leu Leu Ser Leu Pro Leu Arg Thr Ala Leu Gln Ile Met Gln
610 615 620

10 Val Asn Met Leu Arg Ala Val Gly Ile Glu Leu Thr Ile Val Val Gly
625 630 635 640

15 His Ser Ser Gly Glu Ile Val Ala Ala Tyr Ala Ala Gly Val Leu Thr
645 650 655

20 Ala Ser Asp Ala Ile Arg Ile Ala Tyr Leu Arg Gly Met Thr Ile Asp
660 665 670

25 Lys Ser Arg Asp Pro Thr Gly Arg Met Met Ala Val Asn Leu Thr Trp
675 680 685

30 Gln Gln Ala Gln Asn Ile Cys Ala Leu Glu Ala Tyr Ser Gly Arg Ile
690 695 700

35 Ser Val Ala Ala Ala Asn Ser Pro Ser Ser Val Thr Leu Ser Gly Asp
705 710 715 720

40 Ala Glu Cys Leu Arg Glu Leu Glu Trp Leu Leu Lys Ser Leu Gly Leu
725 730 735

45 Thr Pro Arg Met Leu Arg Val Asp Thr Ala Tyr His Ser Pro His Met
740 745 750

50 Lys Pro Cys Ala Asp Pro Tyr Arg Asp Ala Met Lys Ala Tyr Pro Val
755 760 765

55 Ala Leu Ser Ala Ser Ala Ser Arg Trp Tyr Ser Ser Val Tyr Pro Gly
770 775 780

60 Glu Val Met Thr Gly Tyr Asp Gln Gln Glu Leu Thr Gly Glu Tyr Trp
785 790 795 800

65 Val Glu Asn Met Leu Arg Pro Val Gln Phe Ser Gln Ala Leu Glu Ala
805 810 815

70 Ala Ala Arg Asp Ala Gly Pro Pro Asp Leu Ile Ile Glu Ile Gly Pro
820 825 830

2009274832 06 Jun 2011

C:\NRP\ord\NDCC\KLL\3667244_1.DOC (44/201)

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

Pro Asn His Trp Pro Val Leu Val Glu Ser Leu Pro Phe Tyr Pro Phe
900 905 910

20 Asp His Thr Gln Thr Tyr Trp Thr Gln Ser Arg Met Ser Ser Asn His
915 920 925

25 Asn His Arg Ala Thr Ser Pro Asn Ala Leu Leu Gly Ser Leu Ser Pro
930 935 940

30 Glu Thr Gly Ala Glu Lys Phe Arg Trp Arg Asn Tyr Leu Arg Pro Glu
945 950 955 960

35 Glu Leu Pro Trp Leu Ala Asp Arg Ala Asp Ser Gly Ser Val Phe
965 970 975

Pro Glu Thr Gly Tyr Ile Ser Met Ala Leu Glu Ala Gly Met Ile Met
980 985 990

40 Ala Gln Thr Gln Gly Leu Arg Leu Leu Asn Val Lys Asp Leu Thr Ile
995 1000 1005

45 His Thr Gln Leu Pro Ile Gln Asn Asp Pro Ile Gly Thr Glu Val
1010 1015 1020

50 Leu Val Thr Val Gly Ser Ile His Ser His Asp Gly Ala Ile Thr
1025 1030 1035

55 Ala Trp Phe Cys Cys Glu Ala Val Val Ser Gly Glu Leu Val Gln
1040 1045 1050

Cys Ala Thr Ala Lys Met Ile Met His Pro Gly Asp Ser Asp Arg

1055 1060 1065

5 Ala Leu Leu Pro Pro Gln Gly Gln Leu Pro Gln Ala Leu Glu Pro
1070 1075 1080

10 Val Asp Ser Thr Glu Phe Tyr Asp Ser Leu Arg Arg Ala Asp Tyr
1085 1090 1095

15 His Cys Thr Gly Pro Phe Ser Thr Leu Thr Gly Leu Arg Lys Arg
1100 1105 1110

20 Arg Asp Leu Ala Thr Gly Ser Val Pro Val Pro Ser Asn Asp Ser
1115 1120 1125

25 Asp Glu Pro Met Ala Leu His Pro Ala Ile Leu Asp Leu Gly Val
1130 1135 1140

30 Gln Thr Met Ile Ala Ala Ile Gly Gly Leu Glu Glu Thr Leu Leu
1145 1150 1155

35 Thr Gly Pro Phe Leu Ser Arg Asn Val Asp Ser Thr Trp Ile Asn
1160 1165 1170

40 Pro Val Leu Cys Ala Ser Asp Trp Gln Gly Lys Glu Leu Thr Val
1175 1180 1185

45 Ala Ser Tyr Leu Thr Cys Val Asn Gly Asp Arg Ile Arg Gly Asp
1190 1195 1200

50 Ile Asp Ile Phe Thr Met Asn Gly Glu Lys Ala Val Gln Leu Glu
1205 1210 1215

55 Gly Val Ser Leu Ile Cys Gln Pro Ser Gly Thr Ala Pro Asn Asn
1220 1225 1230

Leu Gln Val Leu Ser Gln Thr Ala Trp Gly Pro Leu Glu Pro Thr
1235 1240 1245

60 Leu Lys Lys Gly Ser Arg Lys Leu Pro Ala Thr Met Leu Gln Leu
1250 1255 1260

65 His Ser Leu Arg Glu Glu Leu Ala Leu Leu Tyr Leu Lys Gln Ala
1265 1270 1275

Arg Asn Gly Leu Thr Asp Leu Glu Arg Ser Gly Leu Asp Phe Asp
1280 1285 1290

5

Gly Ala Arg Leu Leu Ala Trp Met Asn Gln Cys Ile Ala Asn Ala
1295 1300 1305

10

Ser Gln Glu Pro Asp Pro Val Gly Glu Ser Glu Cys Leu Asp Gln
1310 1315 1320

15

Lys Ile Glu Asp Phe Thr Ala Gly Val Ser Pro Ser Leu Leu Asn
1325 1330 1335

20

Asp Pro Gly Leu Thr Ala Ile Ala Ala Val Gly Gln Arg Leu Pro
1340 1345 1350

25

Arg Val Leu Arg Asp Ser Gly Leu Gln Ile Glu Ala Trp Pro Ala
1355 1360 1365

Ile Asp Glu Glu Ser Gln Tyr Leu Lys Glu Asp Leu Gln Val Leu
1370 1375 1380

30

Asp Leu Glu Asp Glu Leu Val Ser Val Val Ser Gln Ala Cys Phe
1385 1390 1395

35

Arg Phe Pro Gln Met Asn Ile Leu Gln Ile Gly Gln Phe Gly Gly
1400 1405 1410

40

His Val His Ser Gly Leu Lys Lys Met Gly Arg Thr Tyr Arg Ser
1415 1420 1425

45

Phe Thr Tyr Ala Gly Leu Ser Val Ser Gly Leu Gln Ala Ile Glu
1430 1435 1440

50

Glu Asp Leu Glu Gln Pro Gly Glu Val Ser His Lys Thr Leu Asp
1445 1450 1455

55

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

25 Glu Ser Glu Ser Leu Gly Asp Phe Ser Leu Leu Leu Cys Arg Thr
1565 1570 1575

30 Pro Asp Ser Pro Ala Glu Pro Gln Ser Arg Gly Asp Leu Leu Leu
1580 1585 1590

35 Leu Gly Gly Asp Ala Glu Glu Ala Asp Cys Leu Thr Ser Glu Leu
1595 1600 1605

40 Phe Glu Leu Val Gln Asp Asp Phe Val Lys Val Ala His Ala Pro
1610 1615 1620

Asp Leu Asp Leu Ile Glu Asp Arg Asp Leu Ser Lys Leu Thr Val
1625 1630 1635

45 Leu Tyr Leu Val Asp Asp Arg Asp Leu Thr Asn Ala Thr Leu Ser
1640 1645 1650

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

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

His Ala Thr Gly Leu Arg His Leu Ala Arg Arg Gln Lys Val Thr
1760 1765 1770

20 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

Val Arg His Ser Ser Ile Ala Ala Val Arg Val Ala Gly Ala Ile
1820 1825 1830

35 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

Tyr Leu Val Glu Gln Val Pro Gln Ser Gly Thr Leu Val Val His
1895 1900 1905

55 Glu Ala Asp Gly Val Leu Gln Ser Val Phe His Gln Met Leu Thr

2009274832 06 Jun 2011

C:\NR\PenNDCC\KL\U667299_1.DOC-1\162\111

140

	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

C:\NRP\9274832\CCV\LL366299_1.DOC (146211)

141

Arg Ile Ala Val Ser Lys Ile Asn Leu Ile Asp Gly Ile Ser Ala
2135 2140 2145
5
Thr Ser Leu His Gln Thr Ile Pro Tyr Ala Phe Pro Pro Val Val
2150 2155 2160
10 Gly Gly Val Leu Ile Gln Pro Pro Pro Leu Pro Asp Cys Ser Leu
2165 2170 2175
15 Ser Gln Leu Thr Ile Asp Ser Leu Arg Asn His Leu His Pro Val
2180 2185 2190
20 Leu Lys Gly Leu Gln Gln Leu Asp Glu Leu Tyr Lys Thr Pro Thr
2195 2200 2205
25 Leu Asp Phe Trp Val Leu Ile Gly Ser Ile Ala Gly Val Leu Gly
2210 2215 2220
His Ala Asp Gln Ala Met Thr Ala Ala Met Ser Glu Lys Met Ala
2225 2230 2235
30 Leu Leu Val Arg His Arg Arg Ala Gln Gly Arg Pro Ala Ser Leu
2240 2245 2250
35 Val His Leu Gly Glu Ile His Gly Ile Ser Ser Pro Ser Pro Ser
2255 2260 2265
40 Gln Pro Leu Trp Cys Gly Pro Val Ala Val Ser Gln Arg Asp Val
2270 2275 2280
Asp Glu Ile Leu Ala Glu Ala Ile Leu Cys Gly Arg Ser Asp Ser
2285 2290 2295
45 Asn Ser Asn Ala Glu Leu Ile Gly Gly Leu Arg His Gln Ser Leu
2300 2305 2310
50 Lys Cys Gly Tyr Gly Glu Cys Pro Ile Pro Lys Leu Trp Pro Phe
2315 2320 2325
55 Tyr Ser Tyr Thr Ala Thr Ala Ser Gln Asp Gln Ile Leu Ala Leu
2330 2335 2340

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

25 Val Ile Leu Ile Leu Ser Gly Val Ser Val Gly Glu Leu Ala His
2420 2425 2430

30 Ala Ala Ala Ser Lys Leu Cys Asn Val Ser Val Gly Lys Pro
2435 2440 2445

35 <210> 269
<211> 509
<212> PRT
<213> Penicillium coprobiuum PF1169

40 Met Asp Asn Met Asp Asn Met Asn Asn Thr Pro Leu Gly Phe Asn Trp
1 5 10 15

45 Ala Trp Ala Val Ile Ile Ser Phe Leu Gly Leu Leu Thr Phe Ser Phe
20 25 30

50 Val Ser Pro His Leu Phe Pro Ser Arg Leu Thr Val Ile Asn Gly Gly
35 40 45

55 Arg Ala Trp Asp Ile Phe Arg Thr Lys Ala Lys Lys Arg Phe Arg Ser
50 55 60

65 Asp Ala Ala Arg Leu Ile Lys Asn Gly Phe Glu Glu Ser Pro Asp Ala
65 70 75 80

75 Phe Arg Ile Ile Thr Asp Asn Gly Pro Leu Leu Val Pro Gln

2009274832 06 Jun 2011

C:\NR\Protein\DCCKLLJ667299_1.DOC-1\62\111

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

10

15 Leu Ile Leu Asp Pro Lys Asn Pro Leu Asn Thr Ile Leu Lys Ser Asn
130 135 140

15

Leu Thr Gln Ala Leu Glu Asp Leu Ser Ala Glu Val Thr Glu Ala Leu
145 150 155 160

20

Ser Ala Thr Cys Thr Asp Asp Pro Glu Trp His Glu Val Ser Val Ser
165 170 175

25

Gln Thr Ala Leu Lys Ile Ile Ala Gln Met Ala Ser Lys Ala Phe Ile
180 185 190

30 Gly Gln Glu Arg Cys Arg Asp Ala Lys Trp His Asn Ile Ile Ile Thr
195 200 205

30

Tyr Thr His Asn Val Tyr Gly Ala Ala Gln Ala Leu His Phe Trp Pro
210 215 220

35

Ser Phe Leu Arg Pro Ile Val Ala Gln Phe Leu Pro Ala Cys Arg Thr
225 230 235 240

40

Leu Gln Ala Gln Ile Ala Glu Ala Arg Glu Ile Leu Glu Pro Leu Val
245 250 255

45

Ala Gln Arg Arg Ala Glu Arg Ala Thr Arg Ala Ala Gln Glu Lys Pro
260 265 270

50 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 Ser Phe Ala
290 295 300

55

Ala Ile His Gly Thr Ser Asn Leu Leu Ala Gln Ala Leu Ile Asp Leu
305 310 315 320

06 Jun 2011

2009274832

144

Cys Gly Gln Pro Glu Leu Val Gln Asp Leu Arg Glu Glu Ala Val Ser
325 330 335

5

Val Leu Gly Lys Glu Gly Trp Thr Arg Ala Ala Leu Tyr Gln Leu Lys
340 345 350

10

Leu Met Asp Ser Ala Leu Lys Glu Ser Gln Arg Leu Ala Pro Asn Arg
355 360 365

15

Leu Leu Ser Met Gly Arg Ile Ala Gln Gly Asp Met Asp Leu Ser Asp
370 375 380

20

Gly Leu Arg Ile His Arg Gly Thr Thr Leu Met Val Ser Ala His Asn
385 390 395 400

Met Trp Asp Pro Glu Ile Tyr Pro Asp Pro Arg Lys Tyr Asp Gly Tyr
405 410 415

25

Arg Phe His Lys Leu Arg Gln Thr Ser Gly Gln Glu Gly Gln His Gln
420 425 430

30

Leu Val Ser Ser Thr Pro Asp His Met Gly Phe Gly Tyr Gly Lys His
435 440 445

35

Ala Cys Pro Gly Arg Phe Phe Ala Ala Ala Gln Ile Lys Val Ala Leu
450 455 460

40

Cys Asn Ile Leu Leu Lys Tyr Asp Ile Glu Tyr Arg Gly Gly Lys Ser
465 470 475 480

Pro Gly Val Trp Gly Gln Gly Ile His Leu Phe Pro Asp Pro Thr Ser
485 490 495

45

Arg Ile His Val Arg Arg Arg Lys Glu Glu Ile Asn Leu
500 505

50

<210> 270
<211> 505
<212> PRT
<213> Penicillium coprobiuum PF1169
55 <400> 270

2009274832 06 Jun 2011

C:\NRP\Downloads\3667299_1.DOC-1\62011

145

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

5 Val Leu Val Leu Ala Gly Leu Tyr Leu Ile Arg Leu Thr Leu Ser Ser
20 25 30

10 Asp Gln Leu Asp Lys Phe Pro Ser Ile Asn Pro Arg Lys Pro Trp Glu
35 40 45

15 Ile Val Asn Val Phe Ala Gln Arg Arg Phe Gln Gln Asp Gly Pro Arg
50 55 60

20 Tyr Leu Glu Ala Gly Tyr Ala Lys Ser Pro Ile Phe Ser Val Val Thr
65 70 75 80

25 Asp Leu Gly Pro Lys Leu Val Val Ser Gly Ala Phe Ile Glu Glu Phe
85 90 95

30 Lys Asp Glu Lys Leu Leu Asp His Tyr Arg Ser Met Ile Glu Asp Phe
100 105 110

35 Met Ala Glu Val Pro Gly Phe Glu Ser Met Phe Leu Gly Asn Leu His
115 120 125

40 Asn Thr Val Leu Arg Asp Val Ile Ser Val Ile Thr Arg Glu Leu Glu
130 135 140

45 Gln Leu Leu Ala Pro Leu Ser Asp Glu Val Ser Ala Ala Leu Val Asp
145 150 155 160

50 Thr Trp Thr Asp Ser Pro Asp Trp His Glu Val Ala Leu Leu Pro Ser
165 170 175

55 Met Leu Gly Leu Ile Ala Lys Val Ser Ser Leu Val Phe Val Gly Glu
180 185 190

50 Pro Leu Cys Arg His Pro Val Trp Leu Glu Thr Val Ile Asn Phe Thr
195 200 205

55 Leu Ile Arg His Asn Ala Ile Leu Ala Leu His Gln Cys Pro Ala Val
210 215 220

Leu Arg Pro Val Leu His Trp Val Leu Pro Pro Cys Gln Lys Leu Arg

2009274832 06 Jun 2011

C:\WRP\en6\RDCC\KLL\3667199_1.DOC-1\162011

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

Asp Ala Phe Ala Lys Gly Asn Lys Tyr Asn Ala Ala Met Val Gln Leu
275 280 285

15 Arg Leu Ala Asn Ala Ser Ile His Ser Ser Ala Asp Leu Leu Val Lys
290 295 300

20 Ile Leu Ile Asn Leu Cys Glu Gln Pro Glu Leu Ile Arg Asp Leu Arg
305 310 315 320

25 Asp Glu Ile Ile Ser Val Leu Gly Glu Asn Gly Trp Arg Ser Ser Thr
325 330 335

30 Leu Asn Gln Leu Lys Leu Leu Asp Ser Val Leu Lys Glu Ser Gln Arg
340 345 350

Leu His Pro Val Thr Thr Gly Ala Phe Ser Arg Phe Thr Arg Gln Asp
355 360 365

35 Ile Lys Leu Thr Asn Gly Thr Glu Ile Pro Ser Gly Thr Pro Ile Met
370 375 380

40 Val Thr Asn Asp Val Ala Gly Asp Ala Ser Ile Tyr Asp Asp Pro Asp
385 390 395 400

45 Val Phe Asp Gly Tyr Arg Tyr Phe Arg Met Arg Glu Gly Ala Asp Lys
405 410 415

50 Ala Arg Ala Pro Phe Thr Thr Gly Gln Asn His Leu Gly Phe Gly
420 425 430

Tyr Gly Lys Tyr Ala Cys Pro Gly Arg Phe Phe Ala Ala Thr Glu Ile
435 440 445

55 Lys Ile Ala Leu Cys His Met Leu Leu Lys Tyr Glu Trp Arg Leu Val
450 455 460

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 coprobiuum 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

C:\NRP\pmb\000\KLL36672\W_1.DOC 3/16/2011

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 Met

35 <210> 272
<211> 464
<212> PRT
<213> Penicillium coprobiuum PF1169

40 <400> 272
35 Met Lys Val Ile Ile Val Gly Gly Ser Ile Ala Gly Leu Ala Leu Ala
1 5 10 15

45 His Cys Leu Asp Lys Ala Asn Ile Asp Tyr Val Ile Leu Glu Lys Lys
20 25 30

50 Lys Glu Ile Ala Pro Gln Glu Gly Ala Ser Ile Gly Ile Met Pro Asn
35 40 45

55 Gly Gly Arg Ile Leu Glu Gln Leu Gly Leu Tyr Asp Gln Ile Glu Glu
50 55 60

60 Leu Ile Glu Pro Leu Val Arg Ala His Val Thr Tyr Pro Asp Gly Phe
65 70 75 80

65 Asn Tyr Thr Ser Arg Tyr Pro Ala Leu Ile Gln Gln Arg Phe Gly Tyr
85 90 95

149

Pro Leu Ala Phe Leu Asp Arg Gln Lys Leu Leu Gln Ile Leu Ala Thr
100 105 110

5 Gln Pro Val Gln Ser Ser Arg Val Lys Leu Asp His Lys Val Glu Ser
115 120 125

10 Ile Glu Val Ser Pro Cys Gly Val Thr Val Ile Thr Ser Asn Gly His
130 135 140

15 Thr Tyr Gln Gly Asp Leu Val Val Gly Ala Asp Gly Val His Ser Arg
145 150 155 160

20 Val Arg Ala Glu Met Trp Arg Leu Ala Asp Ala Ser Gln Gly Asn Val
165 170 175

25 Cys Gly Asn Gly Asp Lys Ala Phe Thr Ile Asn Tyr Ala Cys Ile Phe
180 185 190

30 Gly Ile Ser Ser His Val Asp Gln Leu Asp Pro Gly Glu Gln Ile Thr
195 200 205

35 Cys Tyr Asn Asp Gly Trp Ser Ile Leu Ser Val Ile Gly Gln Asn Gly
210 215 220

40 Arg Ile Tyr Trp Phe Leu Phe Ile Lys Leu Glu Lys Glu Phe Val Tyr
225 230 235 240

45 Asp Gly Ser His Lys Thr Gln Leu His Phe Ser Arg Glu Asp Ala Arg
245 250 255

50 Ala His Cys Glu Arg Leu Ala Gln Glu Pro Leu Trp Lys Asp Val Thr
260 265 270

55 Phe Gly Gln Val Trp Ala Arg Cys Glu Val Phe Gln Met Thr Pro Leu
275 280 285

60 Glu Glu Gly Val Leu Gly Lys Trp His Trp Arg Asn Ile Ile Cys Ile
290 295 300

65 Gly Asp Ser Met His Lys Phe Ala Pro His Ile Gly Gln Gly Ala Asn
305 310 315 320

2009274832 06 Jun 2011

C:\NRP\bio\NCBI\BLAST\667299_1.DOC.JNK\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

40 <212> PRT

<213> Penicillium coprobiuum 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

C:\NR\Prot\NDCCV\KL136672\9_1.DOC (146/211)

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

06 Jun 2011

2009274832

C:\INK\Perl\NDCC\KL13867249_1.DOC-1/16/2011

152

Gly Lys Val His Arg Arg Asn Phe Ala Leu Gly Ile Trp Asn Val Leu
290 295 300

5

Ala Cys Phe Val Glu Leu Leu Tyr Ala Ser Gly Ser Leu
305 310 315

10

<210> 274
<211> 522
<212> PRT
<213> *Penicillium coprobiuum* PF1169

15

<400> 274

Met Ser Thr Gln Glu Val Cys Leu Pro Val Ser Gln Arg Asp Gln Val
1 5 10 15

20

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

Leu Leu Leu Lys Val His Pro Phe Leu Ala Gly Asp Val Thr Arg Arg
85 90 95

40

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

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

Leu Arg Ser Glu Ile Glu Arg Leu Val Pro Glu Ser Ser Ser Gly Leu
225 230 235 240

20 Ser Arg Met Asp Phe Ser Lys His Tyr Cys Gly Leu Gly Asp Trp Ala
245 250 255

25 Ala Leu Leu Ala Lys Asn Trp Ser Gly Phe Val Arg Ala Arg Ala Thr
260 265 270

30 Glu Leu Val Thr Trp Arg Leu Lys Ile Pro Gly Pro Lys Ile Glu Tyr
275 280 285

35 Leu Lys Glu Ala Cys Asn Thr Leu Ile Lys Gly Gln Thr Ser Phe Gln
290 295 300

40 Ala Asp Gly Arg Pro Ser Pro Gly Phe Leu Ser Ser Asn Asp Ile Val
305 310 315 320

45 Ser Ala Leu Leu Ala Met Ile Leu Arg Gln Ala Gly Gln Leu Ala Gly
325 330 335

50 Lys Ser Thr Glu Leu Ser Ile Ala Val Asp Met Arg Gly Asn Phe Lys
340 345 350

55 Thr Pro Ala Phe Asp Asp Tyr Leu Gly Asn Met Val Leu Leu Thr Tyr
355 360 365

Thr Pro Ile Gln Ala Gly Arg Asn Glu Ala Leu Val Asp Gly Thr Asp
370 375 380

Pro Ser Val Glu Leu Arg Gln Glu Cys Leu Glu Asp Leu Thr Gln Ile

2009274832 06 Jun 2011

C:\NRP\on\NDCC\LLJ667299_1.DOC (1/16/2011)

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

Gly Phe Arg Gly Val Pro Ile Pro Leu Ser Ser Phe Arg Asn Phe Glu
435 440 445

15 Ile Phe Gly Leu Asp Phe Gly Glu Ser Leu Gly Ala Gln Pro Arg Gly
450 455 460

20 Phe Gln Leu His Leu Pro Val Leu Gly Gly Met Cys Phe Ile Leu Pro
465 470 475 480

25 Lys Gly Gln Asp Asp Val Ala Ser Thr Glu Pro Trp Asp Leu His Leu
485 490 495

30 Thr Leu Asn Arg Asp Asp Gln Leu Leu Leu Ala Lys Asp Pro Leu Phe
500 505 510

35 Cys Trp Ala Ile Gly Ala Gln Ala Lys Glu
515 520

40 <210> 275
<211> 434
<212> PRT
<213> Penicillium coprobiuum PF1169

45 <400> 275

50 Met Asp Ser Leu Leu Thr Ser Pro Leu Trp Leu Lys Ile Ala His Glu
1 5 10 15

55 Leu Ala Leu Tyr Leu Ser Phe Ile Val Pro Thr Ala Phe Leu Ile Ile
20 25 30

60 Thr Thr Gln Lys Ser Ser Ile Ile Arg Trp Ala Trp Thr Pro Cys Leu
35 40 45

65 Leu Tyr Ile Leu Tyr Gln Phe Ser Leu Arg Val Pro Ser Leu Ser Thr
50 55 60

2009274832 06 Jun 2011

C:\NRP\ord\NRPC\LT\366724W_1.DOC (1467/211)

155

65 Ser Gln Phe Leu Lys Gly Val Ala Ala Gly Gln Ala Thr Val Ala Ala
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

20

Lys Gly Lys Asp Gln Ser Met Ser Arg Lys Arg Phe Val Leu Arg Glu
145 150 155 160

25

Ile Ala Ile Ile Val Trp Gln Tyr Leu Leu Leu Asp Phe Ile Tyr Glu
165 170 175

30 Ser Thr Lys Gly Thr Ser Ala Glu Asp Leu Met Arg Leu Phe Gly Pro
180 185 190

35 Gly Met Glu Ile Lys Tyr Leu Asp Ala Thr Phe Glu Gln Trp Met Gly
195 200 205

40 Arg Leu Ser Val Gly Ile Phe Ser Trp Leu Val Pro Ser Arg Val Cys
210 215 220

40

Leu Asn Ile Thr Ser Arg Leu Tyr Phe Leu Ile Leu Val Val Leu Gly
225 230 235 240

45

Ile Ser Ser Pro Glu Ser Cys Arg Pro Gly Phe Gly Arg Val Arg Asp
245 250 255

50 Val Cys Thr Ile Arg Gly Val Trp Gly Lys Phe Trp His Gln Ser Phe
260 265 270

55 Arg Trp Pro Leu Thr Ser Val Gly Asn Tyr Ile Ala Arg Asp Val Leu
275 280 285

2009274832 06 Jun 2011

C:\WRP\pub\NDCC\KL\3667299_1.DOC (1462011)

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

Pro Leu Ala Ile Met Ile Glu Asp Gly Val Gln Glu Ile Trp Arg Arg
340 345 350

15

Ala Thr Gly Gln Thr Lys Asp Ser Asp Arg Ala Val Pro Phe Trp Gln
355 360 365

20

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 coprobiuum PF1169
45 <400> 276

Met Leu Tyr Arg Ala Lys Leu Val Asp Asp His Gln Ile His Thr Ala
1 5 10 15

50

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

C:\NRP\pub\DCCKLLU667299_1.DOC-1\462011

157

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

5

Gly Ser Ile Ile Thr Val Gln Ser Leu Phe Trp Leu Met Thr Lys Trp
65 70 75 80

10 Asn Ile Asp Ile Asn Thr Leu Phe Thr Thr Thr Arg Ser Lys Ser Ile
85 90 95

15 Asp Thr Ala Arg Leu Ile Lys Val Val Pro Ile Thr Asn Ala Gly Ser
100 105 110

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

Phe Val Phe Gln Ile Phe Cys Val Gly Leu Trp Met Leu Asp Glu Tyr
210 215 220

50 Trp Tyr Tyr Ser Leu Phe Thr Leu Phe Met Leu Val Met Phe Glu Ser
225 230 235 240

55 Thr Val Val Trp Gln Arg Gln Arg Thr Leu Ser Glu Phe Arg Gly Met
245 250 255

55 Ser Ile Lys Pro Tyr Asp Val Trp Val Tyr Arg Glu Arg Lys Trp Gln
260 265 270

2009274832 06 Jun 2011

C:\WRR\Port\BN\CC\KL\366799_1.DOC-1\W\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

Pro Asp Gly Leu Asp Lys Leu Ser Phe Val His Gly Gly Thr Lys Val
340 345 350

20

Leu Gln Val Thr His Pro Asn Leu Thr Gly Asp Ala Gly Leu Lys Asn
355 360 365

25

Leu Ala Ser Asn Val Thr Met Pro Pro Asp Asn Gly Ala Leu Gly Val
370 375 380

30

Val Val Lys Thr Gly Phe Glu Thr Ser Gln Gly Ser Leu Val Arg Thr
385 390 395 400

35

Met Ile Tyr Ser Thr Glu Arg Val Ser Ala Asn Asn Val Glu Ala Leu
405 410 415

Leu Phe Ile Leu Phe Leu Leu Ile Phe Ala Ile Ala Ala Ser Trp Tyr
420 425 430

40

Val Trp Gln Glu Gly Val Ile Arg Asp Arg Lys Arg Ser Lys Leu Leu
435 440 445

45

Leu Asp Cys Val Leu Ile Ile Thr Ser Val Val Pro Pro Glu Leu Pro
450 455 460

50

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

55

2009274832 06 Jun 2011

C:\WRP\bnd\OCCKLLU67299_1.DOC-14K2011

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

C:\NRP\bNDCC\KL\J467299_1.DOC (16K9811)

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
785 790 795 800

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
865 870 875 880

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

60 Asn Gln Pro Ala Pro Pro Val Pro Val Leu Ile Ala His Leu Tyr Pro
915 920 925

65 Pro Gly Pro Thr Asn Pro His Tyr Glu Lys Ala Met Glu Arg Glu Ser
930 935 940

70 Gln Arg Lys Gly Ala Ala Ile Thr Ala Pro Gly Ser Thr Pro Glu Ala
945 950 955 960

2009274832

06 Jun 2011

C:\WRP\pmb\DC\KLL\U667299_1.DOC-1\6/2011

161

Ile Pro Thr Ile Thr Ser Pro Gly Ala Gln Ala Leu Gln Gln Ser Asn
965 970 975

5

Leu Asn Pro Gln Gln Gln Lys Lys Gln Gln Ala Gln Ala Ala Ala Ala
980 985 990

10

Gly Leu Ala Asp Lys Leu Thr Ser Ser Met Met Glu Gln Glu Leu Asp
995 1000 1005

15

Asp Ser Glu Pro Pro Thr Ile Lys Leu Gly Asp Ala Ser Val Ala
1010 1015 1020

20 20

Ala Pro Phe Thr Ser Lys Leu Ala Asn Val Ile Ala Ile Pro Asn
1025 1030 1035

Ile Ile Arg Gln Gly Arg Cys Thr Leu Val Ala Thr Ile Gln Met
1040 1045 1050

25

Tyr Lys Ile Leu Ala Leu Asn Cys Leu Ile Ser Ala Tyr Ser Leu
1055 1060 1065

30

Ser Val Ile Tyr Leu Asp Gly Ile Lys Phe Gly Asp Gly Gln Val
1070 1075 1080

35

Thr Ile Ser Gly Met Leu Met Ser Val Cys Phe Leu Ser Ile Ser
1085 1090 1095

40

Arg Ala Lys Ser Val Glu Gly Leu Ser Lys Glu Arg Pro Gln Pro
1100 1105 1110

Asn Ile Phe Asn Val Tyr Ile Ile Gly Ser Val Leu Gly Gln Phe
1115 1120 1125

45

Ala Ile His Ile Ala Thr Leu Ile Tyr Leu Ser Asn Tyr Val Tyr
1130 1135 1140

50

Lys His Glu Pro Arg Asp Ser Asp Ile Asp Leu Glu Gly Glu Phe
1145 1150 1155

55

Glu Pro Ser Leu Leu Asn Ser Ala Ile Tyr Leu Leu Gln Leu Ile
1160 1165 1170

5 Gln Gln Ile Ser Thr Phe Ser Ile Asn Tyr Gln Gly Arg Pro Phe
1175 1180 1185

10 Arg Glu Ser Ile Arg Glu Asn Lys Gly Met Tyr Trp Gly Leu Ile
1190 1195 1200

15 Ala Ala Ser Gly Val Ala Phe Ser Cys Ala Thr Glu Phe Ile Pro
1205 1210 1215

20 Glu Leu Asn Glu Lys Leu Arg Leu Val Pro Phe Thr Asn Glu Phe
1220 1225 1230

25 Lys Val Thr Leu Thr Val Leu Met Ile Phe Asp Tyr Gly Gly Cys
1235 1240 1245

30 Trp Leu Ile Glu Asn Val Leu Lys His Leu Phe Ser Asp Phe Arg
1250 1255 1260

35 Pro Lys Asp Ile Ala Ile Arg Arg Pro Asp Gln Leu Lys Arg Glu
1265 1270 1275

40 Ala Glu Arg Lys Leu Gln Glu Gln Val Asp Ala Glu Ala Gln Lys
1280 1285 1290

45 Glu Leu Gln Arg Lys Val
1295

50 <210> 277
<211> 27
<212> DNA
<213> Artificial Sequence

55 <220>
<223> a primer sequence for PCR

60 <400> 277
gcgcggtaacc attgagacaa catggat

65 <210> 278
<211> 26
<212> DNA
<213> Artificial Sequence

70 <220>
<223> a primer sequence for PCR

2009274832 06 Jun 2011

C:\NRPort\NDCC\KLL\W67249_1.DOC-1\162411

163

<400> 278
atttaaatag ttagacaata gtagca 26

5 <210> 279
<211> 35
<212> DNA
<213> Artificial Sequence

10 <220>
<223> a primer sequence for PCR

<400> 279
gcgcggggta ccatgttgc ttagcattt catta 35

15 <210> 280
<211> 40
<212> DNA
20 <213> Artificial Sequence

<220>
<223> a primer sequence for PCR

25 <400> 280
accacgctaa agatggggga ctttgcatac ccagcttcca 40

30 <210> 281
<211> 40
<212> DNA
<213> Artificial Sequence

<220>
35 <223> a primer sequence for PCR

<400> 281
tggaagctgg gtatgcaaag tccccatct ttagcgtgg 40

40 <210> 282
<211> 40
<212> DNA
<213> Artificial Sequence

45 <220>
<223> a primer sequence for PCR

<400> 282
50 ggtacctctg ccatgaagtc ctgcgttatt gaccgataat 40

<210> 283
<211> 40
55 <212> DNA
<213> Artificial Sequence

2009274832 06 Jun 2011

C:\NRP\orbNDCC\KLLU667299_1.DOC (1/06/2011)

164

<220>
<223> a primer sequence for PCR

5 <400> 283
attatcggtc aatgatcgag gacttcatgg cagaggtacc 40

10 <210> 284
<211> 40
<212> DNA
<213> Artificial Sequence

<220>
<223> a primer sequence for PCR
15 <400> 284
gagaggtgct agcagttgtt ctagttcgcg agtgatgaca 40

20 <210> 285
<211> 40
<212> DNA
<213> Artificial Sequence

25 <220>
<223> a primer sequence for PCR

<400> 285
tgtcatcact cgcgaaactag aacaactgct agcacctctc 40
30

<210> 286
<211> 40
<212> DNA
35 <213> Artificial Sequence

<220>
<223> a primer sequence for PCR

40 <400> 286
cagtgctacc tcatgccagt ctggtgagtc cgtccaagta 40

45 <210> 287
<211> 40
<212> DNA
<213> Artificial Sequence

50 <220>
<223> a primer sequence for PCR

<400> 287
tacttggacg gactcaccag actggcatga ggttagcactg 40

55 <210> 288
<211> 40

<212> DNA
<213> Artificial Sequence

5 <220>
<223> a primer sequence for PCR

<400> 288
agtaaaagcgc gaaaatgctc cggttgtgac tggatgcaac 40

10 <210> 289
<211> 40
<212> DNA
<213> Artificial Sequence

15 <220>
<223> a primer sequence for PCR

<400> 289
20 gttgcacatcca gtcacaacccg gagcattttc gcgctttact 40

25 <210> 290
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> a primer sequence for PCR

30 <400> 290
attttcccta gatcacttct 20

35 <210> 291
<211> 40
<212> DNA
<213> Artificial Sequence

40 <220>
<223> a primer sequence for PCR

<400> 291
45 ccgaattcga gctcggtacc ttgtcttagc attttcatta 40

<210> 292
<211> 38
<212> DNA
50 <213> Artificial Sequence

<220>
<223> a primer sequence for PCR

55 <400> 292
ctactacaga tccccgggtc gtaattttcc ctagatca 38