DESCRIPTION

WEST NILE VIRUS VACCINE, AND METHOD FOR PRODUCTION THEREOF

Technical Field

[0001]

The present invention relates to an artificial signal peptide enabling high secretion expression of virus-like particles of West Nile virus, a recombinant expression vector comprising a nucleic acid encoding a West Nile virus-derived protein including the signal peptide, a transformant harboring the vector, a production method of virus-like particles of West Nile virus by using the transformant, West Nile virus vaccine containing virus-like particles obtained by the method and the like.

Background Art

[0002]

West Nile fever is a systemic acute fever disease caused by infection with West Nile virus (WNV). Occasionally, the virus invades and grows in the central nervous system to cause lethal brain meningitis. WNV is widely distributed in Africa, Middle East, part of Europe, Russia, India, Indonesia and the like. The virus is maintained and propagated by an infection ring between Culex species as a vector and birds (wild and domestic) as an amplification animal. During the process, human, horse and domestic animals become accidental hosts.

In summer 1999, WNV invaded and was indigenized in New York, USA, and has continuously been expanding since then. It was confirmed that more than 2300 persons were infected by the end of September last year (2007) throughout the United States, thus causing a serious problem for the public health. A WNV vaccine for human does not exist in the world at present.

[0003]

Under the circumstances, propagation of the virus to Asian countries including Japan has been feared, and practicalization of human vaccine has been desired. while culture Vero cell-derived virus inactivation vaccines are

being urgently developed at present, there is an increasing need for the development of a subunit vaccine that can be produced safely at a low cost without using a biosafety level 3 virus in the production step.

[0004]

WNV was first isolated in the West Nile region of Uganda, Africa, in 1937 and is classified to belong to the Flaviviridae family falvivirus genus (non-patent document 1). The structure of virus particles consists of a spherical structure wherein a capsid protein (C protein) is bonded to one (+) chain RNA virus gene, and a lipid bilayer membrane surrounding the spherical structure. The lipid membrane includes two kinds of proteins: envelope protein (E protein) and membrane protein (M protein). M protein is produced as a precursor prM protein and cleaved with a protease called furin to become a mature protein.

[0005]

The present inventors previously reported a production method of JEV VLP, comprising introducing an expression vector containing a cDNA fragment encoding prM protein and E protein of Japanese encephalitis virus (JEV) into an animal cell, culturing the obtained transformed cell and harvesting virus-like particles (VLP) secreted in the medium (patent document 1). However, no report has ever been made relating to the production of WNV subunit vaccine.

patent document 1: JP-A-2004-65118

non-patent document 1: Agrawal, A. G., L. R. Petersen, J. Infect. Dis., 188: 1-4 (2003)

Disclosure of the Invention

Problems to be Solved by the Invention

[0006]

An object of the present invention is to provide a high secretion production method of WNV VLP useful as a safe and economical WNV vaccine component, thereby enabling the development and stable supply of WNV vaccine. Another object

of the present invention is to provide a WNV DNA vaccine capable of producing the aforementioned VLP in the animal body.

Means of Solving the Problems

[0007]

The present inventors have conducted intensive studies in an attempt to achieve the aforementioned objects and succeeded in efficiently secreting VLP in a medium by introducing an expression vector containing a DNA fragment encoding prM protein and E protein of WNV into an animal cell, and cultivating the obtained transformed cell in the medium. Moreover, the present inventors have found that, unexpectedly, the formation or secretion efficiency of VLP can be remarkably improved as compared to the native full length signal sequence of WNV by altering a signal peptide located upstream of the prM protein.

The present inventor have conducted further studies based on these findings and completed the present invention.

[0008]

Accordingly, the present invention provides

[1] a signal peptide comprising an amino acid sequence the same or substantially the same as the amino acid sequence of the following (a) or (b)

(a) the amino acid sequence shown by SEQ ID NO: 1

(b) a partial amino acid sequence of the amino acid sequence of the above-mentioned (a), comprising at least the amino acid sequence shown by amino acid NOs 11 - 25;

[2] an isolated nucleic acid substantially consisting of a base sequence encoding the signal peptide of the above-mentioned [1];

[3] a West Nile virus-like particle expression vector comprising a nucleic acid comprising a base sequence encoding the signal peptide of the above-mentioned [1], and prM protein and E protein derived from West Nile virus;

[4] a transformant obtained by transformation with the vector of the above-mentioned [3];

[5] a method of producing a West Nile virus-like particle, comprising culturing the transformant of the above-mentioned [4], and recovering a virus-like particle secreted in a medium;

[6] a West Nile virus-like particle obtained by the method of the above-mentioned [5];

[7] a West Nile virus vaccine comprising the particle of the above-mentioned [6] as an active ingredient;

[8] a West Nile virus vaccine comprising the vector of the above-mentioned [3] as an active ingredient,

and the like.

Effect of the Invention

[0009]

By connecting the signal peptide of the present invention to a polyprotein comprised of prM protein and E protein of WNV, the efficiency of formation and extracellular secretion of VLP in a host cell can be improved. Therefore, a secretion expression vector containing the signal peptide, and a nucleic acid encoding prM protein and E protein is useful for the preparation of a WNV VLP highly secreting or producing cell line, and moreover, the vector itself can be used as a WNV DNA vaccine.

Brief Description of the Drawings

[0010]

Fig. 1 shows polypeptides encoded by recombinant expression vectors (pWPME1 - pWPME14).

Fig. 2 shows the results of Western blot of WNV E protein expressed and secreted by HEK293 T cells harboring recombinant expression vectors (pWPME1 - pWPME14), wherein the left chart shows the results of Western blot of the proteins obtained by solubilizing the cell, and the right chart shows the results of Western blot of the proteins secreted in a culture medium. In the upper left chart and the upper right chart, anti-Japanese encephalitis virus (JEV) antibody was used as a primary antibody, and in the lower left chart and the lower

right chart, anti-M protein polyclonal antibody was used as a primary antibody. In the Figures, E, prM and M show the positions of E protein, prM protein and M protein, respectively.

Fig. 3 shows the results of comparison by the sandwich ELISA method of quantified E proteins secreted in a culture medium by HEK293T cells harboring recombinant expression vectors (pWPME1 - pWPME14), wherein 402 monoclonal antibody was used as a primary antibody.

Fig. 4 shows the results of Western blot of WNV E proteins expressed by HEK293T cells harboring recombinant expression vectors (pWPME1, 7, 9, 12, pWPME388A, pWPME406A). An anti-Japanese encephalitis virus (JEV) antibody was used as a primary antibody, wherein E and prM show the positions of E protein and prM protein, respectively.

Fig. 5 shows the results of comparison by the sandwich ELISA method of quantified E proteins secreted in a culture medium by HEK293T cells harboring recombinant expression vectors (pWPME1, 7, 9, 12, pWPME388A, pWPME406A), wherein 402 monoclonal antibody was used as a primary antibody.

Fig. 6 shows the results of comparison by the sandwich ELISA method of quantified E proteins in each sample fraction after sucrose density gradient centrifugation of the protein secreted in a culture medium by HEK293T cell harboring a recombinant expression vector (pWPME12).

Fig. 7 shows electron microscopic images of WNV-like particles secreted in a culture medium by HEK293T cells harboring a recombinant expression vector (pWPME12). The left charts show virus-like particles contained in sample fractions with density 1.16, and the right charts show virus-like particles contained in sample fractions with density 1.11.

Fig. 8 shows the results of quantification by the sandwich ELISA method of E proteins in each sample fraction after sucrose density gradient centrifugation of the protein secreted in a culture medium by HEK293T cells harboring a

recombinant expression vector (pWPME12). A shows the results obtained by using WNY-11 as a trapping antibody and WNY-11 as a detection antibody. B shows the results obtained by using WNY-11 antibody as a trapping antibody and 402 antibody as a detection antibody. C shows the results obtained by using 402 antibody as a trapping antibody and 402 antibody as a detection antibody. D shows the results obtained by using 402 antibody as a trapping antibody and WNY-11 antibody as a detection antibody. E shows the density of each sample fraction.

Fig. 9 shows the results of a plaque decrease assay of WNV by using the serum of mouse immunized with WNV-like particles, wherein the serum used was diluted and the plaque number is the total of 3 wells.

Fig. 10 shows the results of comparison by Western blot of the expression level of WNV-like particles after sucrose density gradient centrifugation of the protein secreted in a culture medium by CHO-K1 (parent strain) and each clone acclimated in a serum-free medium, wherein anti-Japanese encephalitis virus (JEV) antibody was used as a primary antibody.

Best Mode for Carrying out the Invention

[0011]

The present invention provides a signal peptide for high secretion production of WNV VLP. The signal peptide comprises

(a) the amino acid sequence shown by SEQ ID NO: 1;

(b) a partial amino acid sequence of the amino acid sequence of the above-mentioned (a), comprising at least the amino acid sequence shown by amino acid NOs 11 - 25; or

(c)an amino acid sequence the same or substantially the same as the amino acid sequence of
the above-mentioned (a) or (b).

[0012]

Here, the "substantially the same amino acid sequence" is the amino acid sequence of the above-mentioned (a) or (b),

wherein one to several (2, 3, 4 or 5) amino acids are substituted, deleted, added or inserted, which has a significantly high WNV VLP secretion ability as compared to WNV native signal sequence.

[0013]

When "substantially the same amino acid sequence" contains substitution of the amino acid, it is desirably similar to the original amino acid in the physicochemical properties. For example, substitution of amino acids classified into the same group such as aromatic amino acid (Phe, Trp, Tyr), aliphatic amino acid (Ala, Leu, Ile, Val), polar amino acid (Gln, Asn), basic amino acid (Lys, Arg, His), acidic amino acid (Glu, Asp), amino acid having a hydroxyl group (Ser, Thr), amino acid having a small side chain (Gly, Ala, Ser, Thr, Met) and the like can be mentioned. It is predicted that the substitution of such similar amino acids does not change the phenotype of a protein (i.e., preservative amino acid substitution). Specific examples of the preservative amino acid substitution are well known in the art, and described in various documents (see, for example, Bowie et al., Science, 247: 1306-1310 (1990)). When an amino acid (sequence) is substituted, deleted or inserted, the position of the substitution, deletion or insertion is not particularly limited as long as the WNV VLP secretion ability of the original signal peptide is substantially maintained. The "substantially maintained" means having a significantly high secretion ability at least as compared to the native signal sequence of WNV.

[0014]

The amino acid sequence shown by SEQ ID NO: 1 corresponds to an amino acid sequence of the 99th - 123rd (i.e., amino acids from immediately before N terminal of prM protein to 25 amino acid residues in the upstream) of the polyprotein precursor (GenBank Accession No. AAF20092.2) of West Nile virus NY99-flamingo382-99 strain (Lanciotti, R. S. et al.,

Science, 286: 2333-2337, 1999) (GenBank Accession No. AF196835). Many variants of WNV have been reported so far, and new mutant virus strains will also be found one after another. Thus, an amino acid sequence corresponding to the above-mentioned particular sequence of the NY99-flamingo382-99 strain in such other mutant WNV strains is also encompassed in the "substantially the same amino acid sequence" of the signal peptide of the present invention. Examples of the mutant WNV strain other than the NY99-flamingo382-99 strain include those described in "Table I" of Ebel, G.D. et al., Am. J. Trop. Med. Hyg., 71(4): 493-500, 2004 and the like. The above-mentioned "substantially the same amino acid sequence" can be acquired easily by obtaining a sequence of each Accession No. shown in the Table and determining the corresponding region.

[0015]

The length of the amino acid sequence of the signal peptide of the present invention is desirably 15 amino acids in the shortest and 25 amino acids in the longest. It preferably has a length other than 20 amino acids. More preferably, the length of the amino acid sequence of the signal peptide is 15 - 19 or 21 - 25 amino acids, more preferably 15 - 18 amino acids.

Therefore, when the "substantially the same amino acid sequence" contains deletion, addition or insertion of amino acids, the number thereof is desirably within a range affording the above-mentioned preferable full-length after alteration. When an amino acid is added or inserted, the kind of the amino acid to be added or inserted is not particularly limited as long as the full-length after the addition or insertion falls within the above-mentioned preferable range.

[0016]

The signal peptide of the present invention can also be produced by a known peptide synthesis method, for example, a solid phase synthesis process and a liquid phase synthesis process, based on the information of the amino acid sequence

thereof. In view of the object of use of the peptide, which is for efficient secretion of WNV VLP from a host cell, the peptide is provided as an N-terminal region of a polyprotein precursor produced by culturing a transformant harboring an expression vector containing a nucleic acid encoding the precursor, wherein the nucleic acid consists of a nucleic acid comprising a base sequence encoding the peptide, which is operably linked to a nucleic acid encoding prM protein and E protein constituting the object VLP.

Therefore, the present invention also provides an isolated nucleic acid substantially consisting of a base sequence encoding the above-mentioned signal peptide of the present invention. Here, "substantially consisting of" means that an amino acid (excluding initiating methionine residue) other than the amino acid sequence of the signal peptide of the present invention is not contained in the N-terminal region (other than prM protein and E protein) of a polyprotein precursor, which is produced by a suitable host cell harboring a suitable expression vector comprising the above nucleic acid operably linked to a nucleic acid encoding the prM protein and E protein of WNV. Therefore, examples of the sequence that can be contained in the nucleic acid besides the base sequence encoding the signal peptide of the present invention include initiating ATG codon, a restriction enzyme recognition sequence that facilitates ligation of the nucleic acid to the downstream of the promoter sequence of an expression vector or the upstream of a nucleic acid encoding prM protein of WNV, and the like.

The nucleic acid may be a DNA or an RNA, or a DNA/RNA chimera, with preference given to DNA. The nucleic acid may be double stranded or single stranded. When the nucleic acid is double stranded, it may be double stranded DNA, double stranded RNA or DNA:RNA hybrid.

[0017]

The base sequence encoding the above-mentioned signal

peptide of the present invention is not particularly limited as long as it generates, after translation, any of the amino acid sequences of the above-mentioned signal peptide of the present invention. When a desired signal peptide is completely the same as a part of a polyprotein precursor sequence of any of the available WNV strains, a base sequence encoding the amino acid sequence can be obtained by PCR method and the like and using, as a template, cDNA prepared from the genomic RNA of the virus strain. Such method is advantageous since not only a signal peptide but also a base sequence encoding prM protein and E protein of WNV can be obtained at once. Depending on the selection of a primer sequence, a nucleic acid encoding signal peptides having different lengths of amino acid sequences can be prepared with ease.

[0018]

On the other hand, in consideration of the expression efficiency in a host cell, it is generally preferable to select a codon highly frequently used for the host cell to be used. The data of use frequency of codon in various biological species can be obtained from the genetic code use frequency database disclosed, for example, in the home page of Kazusa DNA Research Institute (http://www.kazusa.or.jp/codon/index.html). For conversion of a base sequence in accordance with codon use frequency of the host cell, the full-length sequence of a nucleic acid encoding a signal peptide is chemically synthesized by a DNA/RNA automatic synthesizer, or partly overlapping oligoDNA short chains synthesized are connected by PCR method.

[0019]

The present invention also provides a WNV VLP expression vector containing the above-mentioned signal peptide of the present invention, and a nucleic acid comprising a base sequence encoding prM protein and E protein derived from WNV.

Here, the "base sequence encoding prM protein of WNV" means a base sequence encoding a protein comprising an amino

acid sequence the same or substantially the same as the amino acid sequence shown by SEQ ID NO: 3. The "base sequence encoding E protein of WNV" means a base sequence encoding a protein comprising an amino acid sequence the same or substantially the same as the amino acid sequence shown by SEQ ID NO: 5.

[0020]

As the "amino acid sequence substantially the same as the amino acid sequence shown by SEQ ID NO: 3 (or SEQ ID NO: 5)", an amino acid sequence having a homology of not less than about 80%, preferably not less than about 90%, more preferably not less than about 95%, particularly preferably not less than about 97%, with the amino acid sequence shown by SEQ ID NO: 3 (or SEQ ID NO: 5) and the like can be mentioned.

Here, "homology" means a ratio (%) of amino acid residues identical or similar to all overlapping amino acid residues in the optimal alignment where two amino acid sequences are aligned using a mathematical algorithm known in the technical field (preferably, the algorithm can consider introduction of gaps on one or both of the sequences for the best alignment). The "similar amino acid" refers to an amino acid similar in its physiochemical properties, and examples include amino acids classified in the same group as the original amino acids in, for example, a classification similar to the aforementioned classification of the signal peptide of the present invention.

[0021]

Homology of the amino acid sequences in the present specification can be measured under the following conditions (an expectation value = 10; gaps are allowed; matrix = BLOSUM62; filtering = OFF) using a homology scoring algorithm NCBI BLAST (National Center for Biotechnology Information Basic Local Alignment Search Tool). Other algorithm for determining homology of the amino acid sequence is exemplified by an algorithm disclosed in Karlin et al., Proc. Natl. Acad.

Sci. USA, 90: 5873-5877 (1993) [this algorithm is incorporated in NBLAST and XBLAST program (version 2.0) (Altschul et al., Nucleic Acids Res., 25:3389-3402(1997))]; an algorithm disclosed in Needleman et al., J. Mol. Biol., 48: 444-453 (1970) [This algorithm is incorporated in a GAP program in a GCG software package]; an algorithm disclosed in Myers and Miller, CABIOS, 4: 11-17 (1988) [This algorithm is incorporated in ALIGN program (version 2.0) which is a part of a CGC sequence alignment software package]; an algorithm disclosed in Pearson et al., Proc. Natl. Acad. Sci. USA, 85: 2444-2448 (1998) [This algorithm is incorporated in an FASTA program in a GCG software package], etc., and these may be also preferably used.

[0022]

More preferably, the "amino acid sequence substantially the same as the amino acid sequence shown by SEQ ID NO: 3 (or SEQ ID NO: 5)" is an amino acid sequence having an identity of not less than about 80%, preferably not less than about 90%, more preferably not less than about 95%, particularly preferably not less than about 97%, with the amino acid sequence shown by SEQ ID NO: 3 (or SEQ ID NO: 5).

[0023]

The "protein comprising an amino acid sequence substantially the same as the amino acid sequence shown by SEQ ID NO: 3 (or SEQ ID NO: 5)" means a protein comprising an amino acid sequence substantially the same as the aforementioned amino acid sequence shown by SEQ ID NO: 3 (or SEQ ID NO: 5), and having substantially identical activity as a protein comprising the amino acid sequence shown by SEQ ID NO: 3 (or SEQ ID NO: 5).

"Substantially identical activity" in the case of prM protein is, for example, an activity to change conformation of E protein when it has matured as M protein, and the like, and that of E protein is, for example, adsorption to infected host cell, red blood cell coagulation activity and the like.

"Substantially identical" means that these activities are qualitatively the same. Therefore, each of the above-mentioned activities are preferably equivalent (e.g., about 0.5- to about 2-fold), but quantitative factors such as the extent of activity and protein molecular weight may be different.

[0024]

The prM protein (E protein) to be used in the present invention includes, for example, a protein comprising (1) an amino acid sequence shown by SEQ ID NO: 3 (or SEQ ID NO: 5) wherein one or more (preferably about 1 - 30, more preferably about 1 - 10, still more preferably 1 - several (2, 3, 4 or 5)) amino acids are deleted, (2) an amino acid sequence shown by SEQ ID NO: 3 (or SEQ ID NO: 5) wherein one or more (preferably about 1 - 30, more preferably about 1 - 10, still more preferably 1 - several (2, 3, 4 or 5)) amino acids are added, (3) an amino acid sequence shown by SEQ ID NO: 3 (or SEQ ID NO: 5) wherein one or more (preferably about 1 - 30, more preferably about 1 - 10, still more preferably 1 - several (2, 3, 4 or 5)) amino acids are inserted, (4) an amino acid sequence shown by SEQ ID NO: 3 (or SEQ ID NO: 5) wherein one or more (preferably about 1 - 30, more preferably about 1 - 10, still more preferably 1 - several (2, 3, 4 or 5)) amino acids are substituted by other amino acids, (5) a combination of such amino acid sequences, or the like.

When an amino acid sequence is inserted, deleted or substituted as mentioned above, the position of the insertion, deletion or substitution is not particularly limited as long as the activity of protein is maintained.

[0025]

DNA encoding prM protein (or E protein) of WNV can be directly amplified by Reverse Transcriptase-Polymerase Chain Reaction (hereinafter to be abbreviated as "RT-PCR method") and using RNA prepared from a virus stock of WNV as a template. Since prM protein and E protein are continuously encoded in the WNV genome, DNAs encoding prM protein and E protein can be

prepared at once by designing a primer capable of amplifying a base sequence covering the coding regions of the both. Moreover, when a sequence in the WNV genome is utilized as a base sequence encoding the signal peptide of the present invention, since the signal peptide coding region is present immediately before the coding region of prM protein, the three can also be prepared at once.

[0026]

Examples of the DNA encoding prM protein (or E protein) include a DNA containing a base sequence shown by SEQ ID NO: 2 (or SEQ ID NO: 4), a DNA containing a base sequence that hybridizes with a complementary chain sequence of the base sequence shown by SEQ ID NO: 2 (or SEQ ID NO: 4) under stringent conditions, and encoding a protein having a substantially identical activity to the aforementioned prM protein (or E protein), and the like.

Examples of the DNA capable of hybridizing to the base sequence shown by SEQ ID NO: 2(or SEQ ID NO: 4) under stringent conditions include DNA containing a base sequence having not less than about 70%, preferably not less than about 80%, more preferably not less than about 90%, and particularly preferably not less than about 95%, homology to the base sequence shown by SEQ ID NO: 2 (or SEQ ID NO: 4) and the like.

The homology of the base sequence in the present specification can be calculated using homology calculation algorithm NCBI BLAST (National Center for Biotechnology Information Basic Local Alignment Search Tool) under the following conditions (expectancy=10; gaps are allowable; filtering=ON; match score=1; mismatch score=-3). Examples of other algorithms to determine a homology of base sequence preferably include the above-mentioned homology calculation algorithms of amino acid sequence in a similar manner.

[0027]

Hybridization can be conducted according to a method known per se or a method based thereon, for example, a method

described in Molecular Cloning, 2nd edition (J. Sambrook et al., Cold Spring Harbor Lab. Press, 1989) and the like. When a commercially available library is used, hybridization can be conducted according to the method described in the instruction manual attached thereto. Hybridization can preferably be conducted under highly stringent conditions.

Examples of the stringent conditions include a hybridization reaction in 6×SSC (sodium chloride/sodium citrate) at 45°C, and then one or more times of washing in 0.2×SSC/0.1% SDS at 65°C, and the like. Those of ordinary skill in the art can easily adjust to a desired stringency by appropriately changing the salt concentration of hybridization solution, the temperature of hybridization reaction, probe concentration, probe length, number of mismatch, hybridization reaction time, salt concentration of washing, washing temperature and the like.

[0028]

A DNA encoding prM protein and E protein can be obtained from WNV genome as mentioned above. It is also possible to construct a DNA encoding the full length of prM protein and E protein by chemically synthesizing a DNA chain, or connecting, by PCR method, partly overlapping oligoDNA short chains synthesized. The advantage of constructing a full-length DNA by combining chemical synthesis or PCR method is that, as mentioned above, the codon to be used can be designed over the full length of the gene according to the host into which the gene is to be introduced.

[0029]

The DNA cloned as mentioned above can be used as is, or after digestion with a restriction endonuclease or addition of a linker as desired, depending on the purpose of its use. When DNA encoding signal peptide, DNA encoding prM protein and DNA encoding E protein are individually cloned, they are ligated in the above-mentioned order by using suitable restriction enzyme, linker, ligase and the like.

[0030]

Then, the obtained DNA encoding "signal peptide-prM protein-E protein" polyprotein precursor, after introduction as necessary of ATG as a translation initiation codon into the 5′ terminal side, and TAA, TGA or TAG as a translation stop codon into the 3′ terminal side, is ligated to the downstream of a promoter in a suitable expression vector by using a restriction enzyme and a DNA ligase. The translation initiation codons and translation stop codons can be added using an appropriate synthetic DNA adapter. Alternatively, utilizing PCR in advance, they may be introduced into the 5′ side of a base sequence encoding the signal peptide and the 3′ side of a base sequence encoding E protein, respectively.

[0031]

Useful expression vectors include plasmids derived from *Escherichia coli* (e.g., pBR322, pBR325, pUC12, pUC13); plasmids derived from *Bacillus subtilis* (e.g., pUB110, pTP5, pC194); plasmids derived from yeast (e.g., pSH19, pSH15); insect cell expression plasmids (e.g., pFast-Bac); animal cell expression plasmids (e.g., pCAGGS, pA1-11, pXT1, pRc/CMV, pRc/RSV, pcDNAI/Neo); bacteriophages such as λ phage; insect viral vectors such as baculovirus (e.g., BmNPV, AcNPV); animal viral vectors such as retrovirus, vaccinia virus, adenovirus; and the like.

The promoter may be any promoter, as long as it is appropriate for the host used to express the gene.

For example, when the host is an animal cell, useful promoters include CAG promoter, β-actin promoter, SRα promoter, SV40 promoter, LTR promoter, CMV (cytomegalovirus) promoter, RSV (Rous sarcoma virus) promoter, MoMuLV (Moloney murine leukemia virus) LTR, HSV-TK (simple herpes virus thymidine kinase) promoter and the like are used. Among these, CAG promoter is preferable.

When the host is a bacterium of the genus *Bacillus*, the SPO1 promoter, the SPO2 promoter, the penP promoter and the

like are preferred.

When the host is yeast, the PHO5 promoter, the PGK promoter, the GAP promoter, the ADH promoter and the like are preferred.

When the host is an insect cell, the polyhedrin promoter, the P10 promoter and the like are preferred.

[0032]

Useful expression vectors include, in addition to the above, those optionally harboring an enhancer, a splicing signal, a polyA addition signal, a selection marker, an SV40 replication origin (hereinafter sometimes to be abbreviated as SV40 ori) and the like. As examples of the selection marker, dihydrofolate reductase gene (hereinafter sometimes to be abbreviated as dhfr, methotrexate (MTX) resistant), ampicillin resistant gene (hereinafter sometimes to be abbreviated as amp^r), neomycin resistance gene (hereinafter sometimes to be abbreviated as neo^r, G418 resistant) and the like can be mentioned. In particular, when a dhfr-deficient Chinese hamster ($CHO\text{-}dhfr^-$) cell is used in combination with the dhfr gene as the selection marker, a target gene can also be selected using a thymidine-free medium.

[0033]

WNV VLP can be produced by culturing a transformant obtained by transforming a host with an expression vector containing a nucleic acid comprising a base sequence encoding the signal peptide of the present invention, prM protein and E protein.

Useful hosts include, for example, those suitable for secretion expression such as bacterium of the genus *Bacillus*, yeast, an insect cell, an insect, an animal cell, an animal and the like.

Useful bacteria of the genus *Bacillus* include, for example, *Bacillus subtilis* MI114, 207-21 and the like.

Useful yeasts include, for example, *Saccharomyces cerevisiae* AH22, $AH22R^-$, NA87-11A, DKD-5D and 20B-12,

Schizosaccharomyces pombe NCYC1913 and NCYC2036, *Pichia pastoris* KM71, and the like.

[0034]

Useful insect cells include, for example, *Spodoptera frugiperda* cell (Sf cell), MG1 cell derived from the mid-intestine of *Trichoplusia ni*, High Five™ cell derived from an egg of *Trichoplusia ni*, cell derived from Mamestra brassicae, cell derived from *Estigmena acrea*, and the like can be mentioned when the virus is AcNPV. When the virus is BmNPV, useful insect cells include *Bombyx mori* N cell (BmN cell) and the like. Useful Sf cells include, for example, Sf9 cell (ATCC CRL1711), Sf21 cell and the like.

Useful insects include, for example, a larva of *Bombyx mori* and the like.

[0035]

Useful animal cell includes, for example, cells such as COS-7, Vero, CHO, CHO (dhfr$^-$), CHO-K1, L, AtT-20, GH3, FL, HEK293, NIH3T3, Balb3T3, FM3A, L929, SP2/0, P3U1, B16, P388 and the like. Preferably, HEK293, CHO-K1 and the like are used but the animal cell is not limited thereto.

As the animal, mammals with established transgenic system (e.g., mouse, rat, rabbit, sheep, swine, bovine), chicken and the like can be mentioned.

[0036]

Transformation can be carried out according to the kind of host in accordance with a publicly known method.

A bacterium of the genus *Bacillus* can be transformed, for example, in accordance with the method described in Molecular & General Genetics, vol. 168, 111 (1979) and the like.

Yeast can be transformed, for example, in accordance with the method described in Methods in Enzymology, Vol. 194, 182-187 (1991), Proc. Natl. Acad. Sci. USA, Vol. 75, 1929 (1978) and the like.

An insect cell and an insect can be transformed, for example, according to a method described in Bio/Technology, 6,

47-55 (1988) and the like.

An animal cell can be transformed, for example, in accordance with the method described in Saibo Kogaku (Cell Engineering), extra issue 8, Shin Saibo Kogaku Jikken Protocol (New Cell Engineering Experimental Protocol), 263-267 (1995), published by Shujunsha, or Virology, Vol. 52, 456 (1973).

An animal can be transformed, for example, in accordance with the method described in Development of Transgenic Animal (CMC Publishing) (2001).

[0037]

A transformant can be cultured in accordance with a known method according to the kind of the host.

For culture of a transformant with the genus *Bacillus* as a host, for example, the medium to be used for the culture is preferably a liquid medium. In addition, the medium preferably contains a carbon source, a nitrogen source, an inorganic substance and the like, which are necessary for the growth of a transformant. Examples of the carbon source include glucose, dextrin, soluble starch, sucrose and the like; examples of the nitrogen source include inorganic or organic substances such as ammonium salts, nitrate salts, corn steep liquor, peptone, casein, meat extract, soybean cake, potato extract and the like; and examples of the inorganic substance include calcium chloride, sodium dihydrogen phosphate, magnesium chloride and the like. In addition, the medium may contain a yeast extract, vitamins, a growth promoting factor and the like. The medium has a pH of preferably about 5 - about 8. A transformant is cultured generally at about 30 - about 40°C for about 6 - about 24 hr. Where necessary, aeration and stirring may also be performed.

Examples of the medium for culture of a transformant with yeast as a host include Burkholder minimum medium, SD medium containing 0.5% casamino acid and the like. The medium has a pH of preferably about 5 - about 8. Culture is performed generally at about 20 - about 35°C for about 24 - about 72 hr.

Where necessary, aeration and stirring may also be performed.

Examples of the medium for culture of a transformant with an insect cell or insect as a host include Grace's Insect Medium appropriately supplemented with an additive such as inactivated 10% bovine serum and the like, and the like. The medium has a pH of preferably about 6.2 - about 6.4. Culture is performed generally at about 27°C for about 3 - about 5 days. Where necessary, aeration and stirring may also be performed.

Examples of the medium for culture of a transformant with an animal cell as a host include minimum essential medium (MEM), Dulbecco's modified Eagle medium (DMEM), RPMI 1640 medium, 199 medium and the like, containing about 5 - about 20% of fetal bovine serum. The medium has a pH of preferably about 6 - about 8. Culture is performed generally at about 30°C - about 40°C for about 15 - about 60 hr. Where necessary, aeration and stirring may also be performed.

When the host is an animal, a transgenic animal is obtained from a transgenic fertilized egg according to a conventional method and bred under general breeding conditions, and mammalian milk or chicken egg is obtained.

In the above manner, WNV VLP can be secreted in the medium of a transformant or extracellularly.

[0038]

In the present invention, moreover, a transformant may be cultured in a medium containing fetal bovine serum as mentioned above. However, from the aspects of removal of a risk factor such as bovine viral diarrhea virus (BVDV) that enters fetal bovine serum to contaminate cells and the like, removal of impurity such as bovine-derived protein and the like in an attempt to simplify a WNV VLP purification step, and economical aspect, a method including culturing a transformant with serum-free acclimation is also preferably used. As described in the below-mentioned Examples, it is preferable to obtain a non-adherent cell from an adherent cell by acclimation in a serum-free medium, since it facilitates

maintenance, passage and mass culture of the transformant.

[0039]

VLP secreted in a medium (extracellularly) can be purified by a method known per se. For example, VLP can be prepared by filtering a recovered medium (extracellular liquid) to remove low molecular weight proteins, and applying the medium to sucrose density gradient centrifugation.

[0040]

The WNV vaccine of the present invention contains WNV VLP as an active ingredient in an amount sufficient to provide immunity. Specifically, for example, WNV vaccine can be produced as in the following, though not limited thereto.

VLP of the present invention purified as an antigen are suspended in a solvent such as an isotoic salt solution, buffer, tissue culture liquid and the like, for example, PBS (phosphate buffered saline) to give a vaccine stock solution. Where necessary, a vaccine antigen may be immobilized with a conventional immobilizer to also immobilize the steric structure thereof. Examples of the immobilizer include formalin, phenol, glutardialdehyde, β-propiolactone and the like. The immobilizer may be add to an antigen before preparation of a vaccine stock solution, or added to a vaccine stock solution.

[0041]

Then, a vaccine stock solution is diluted to give a vaccine solution. A vaccine stock solution is diluted, for example, with PBS so that the amount of the antigen in the vaccine will be sufficient to induce antibody production to afford immunity, for example, 1 - 20 μg/ml, preferably 10 μg/ml, in a protein content. For preparation of a vaccine solution, a stabilizer that potentiates heat resistance of vaccine and an adjuvant as an auxiliary to enhance immunogenicity may be added. As the stabilizer, saccharides and amino acids can be mentioned. As the adjuvant, mineral oil, vegetable oil, alum, aluminum compound, bentonite, silica,

muramyl dipeptide derivative, thymosin, interleukin and the like can be mentioned.

[0042]

A vaccine solution is dispensed in a container with a suitable volume, for example, about 0.5 - 20 ml vial, and the container is tightly sealed and used as a vaccine. Such vaccine may be a liquid or freeze-dried after dispensing and used as a dry preparation.

[0043]

The vaccine of the present invention may be inoculated to a subject in the same manner as with general vaccines. For example, about 0.2 - 0.5 ml of a vaccine for one dose can be subcutaneously inoculated 1 - 3 times at about 1 - 4 weeks intervals. A dry preparation is dissolved in sterile distilled water and the like before inoculation to the original volume and thereafter used.

[0044]

Since an expression vector (non-virus vector and virus vector) containing the above-mentioned signal peptide of the present invention, and a nucleic acid comprising a base sequence encoding prM protein and E protein of WNV can efficiently secrete WNV VLP in the human body after administration thereof to human, it can be used as what is called a DNA vaccine.

Preferable production methods, administration methods and the like of these non-virus vector and virus vector are known to those of ordinary skill in the art and, for example, Experimental Medicine, extra issue, basic technique of gene therapy, YODOSHA, 1996; Experimental Medicine, extra issue, Transgene & Expression Analysis Experiment Method, YODOSHA, 1997; ed. Japan Society of Gene Therapy, Gene Therapy Development Research Handbook, NTS, 1999 and the like are used for reference. Examples of the non-virus vector suitable for DNA vaccines include, but are not limited to, pcDNA3.1, pZeoSV, pBK-CMV (Invitrogen, Stratagene), pCAGGS (Gene 108, 193-200

(1991)) and the like. Representative virus vector includes virus vectors such as recombinant adenovirus, retrovirus and the like. Specifically, for example, DNA viruses such as detoxicated retrovirus, adenovirus, adeno-associated virus, herpes virus, vaccinia virus, pox virus, polio virus, sindbis virus, Hemagglutinating Virus of Japan, SV40, human immunodeficiency virus (HIV) and the like and RNA virus can be mentioned.

[0045]

As a method for introducing the DNA vaccine of the present invention into human, an in vivo method including directly introducing a DNA vaccine into the body, an ex vivo method including taking out a certain cell from human, extracellularly introducing a DNA vaccine into the cell and returning the cell into the body and the like can be mentioned (ed. Japan Society of Gene Therapy, Gene Therapy Development Research Handbook, NTS, 1999). According to the in vivo method, the DNA vaccine of the present invention is dissolved in a suitable solvent (buffer such as PBS and the like, saline, sterile water etc.), sterilized by filtration with a filter etc. as necessary, filled in an aseptic container to give an injection, which is administered to human by injection. Injection may contain a conventional carrier and the like as necessary. In addition, a DNA vaccine may also be enclosed in a liposome made of a lipid bilayer membrane, and administered as an HVJ-liposome obtained by fusion of the liposome and inactivated Hemagglutinating Virus of Japan (HVJ) (Experimental Medicine, extra issue, basic technique of gene therapy, YODOSHA, 1996; Transgene & Expression Analysis Experiment Method, YODOSHA, 1994). The DNA vaccine of the present invention can be administered to the muscle, skin, nasal cavity and the like. As the ex vivo method, a lipofection method, a phosphoric acid-calcium coprecipitation method, a DEAE-dextran method, a method including direct injection of DNA vaccine intracellularly using a tiny glass

tube and the like, and the like can be mentioned.

[0046]

While the dose of the DNA vaccine of the present invention varies depending on the administration target, administration method, administration manner and the like, it is generally about 500 μg - about 50 mg, preferably about 500 μg - about 1 mg, per adult based on the gene.

[0047]

The present invention is explained in more detail in the following by referring to Examples, which are not to be construed as limitative.

Examples

[0048]

Example 1 Production of recombinant expression vector expressing WNV-derived mutated signal peptide sequence, prM protein and E protein

Polyadenylated RNA was extracted from WNV (WN-NY99), and using same as a template and 15 nucleotide poly(dT)primer, a reverse transcription reaction was performed with SuperScript RNase H^- reverse transcriptase (Invitrogen Life Technologies, Carlsbad, CA). The coding region of the obtained WNV cDNA was amplified by PCR by using a complementary primer set (base sequences shown by SEQ ID NO: 6 and SEQ ID NO: 7). The obtained PCR product was digested with BglII and XhoI, and subcloned to pBluescript plasmid. Using the obtained plasmid as a template and a primer set complementary to the base sequence encoding a signal sequence defective in one amino acid from the amino terminal side (one base sequence selected from the group consisting of SEQ ID NOs: 8 - 21 and one base sequence shown by SEQ ID NO: 22), DNA fragment was amplified by PCR. The amplified fragment was subcloned to a mammal expression vector (pCAGGS) to give 14 kinds of recombinant expression vectors expressing signal sequence having a particular length, prM protein and E protein (pWPME1 - pWPME14). The structure of polypeptide encoded by the obtained

PCR product is shown in Fig. 1.

[0049]

Example 2 Consideration of signal peptide sequence having length within range suitable for secretion

Various recombinant expression vectors (0.5 μg) obtained in Example 1 were introduced into HEK293T cells (2×10^5) using Fugene 6 (Roche Diagnostic, Basel, Switzerland). After 48 hr, the cells were washed twice with phosphate buffered saline (PBS), and solubilized with a solubilizing solution containing 50 mM Tris-HCl (pH 6.8), 100 mM ditiothreitol, 2% sodium dodecyl sulfate(SDS), 0.1% bromophenol blue and 10% glycerol. On the other hand, the culture supernatant was filtered (0.45 μm pore size, Millipore, Bedford, MA), and subjected to an ultracentrifugation treatment at 40,000×g for 30 min. The residue was solubilized with the aforementioned solubilizing solution.

[0050]

According to a conventional method, the WNV-like particles obtained in the above were subjected to SDS-PAGE, membrane transcription and Western blot using an antibody. Specifically, protein was separated by 10% PAGE, and the separated protein was transcribed to a polyvinyl difluoride (PDVF) membrane filter (Millipore). As primary antibodies, an anti-Japanese encephalitis virus (JEV) antibody that cross reacts with WNV and an anti-M protein polyclonal antibody (IMGENEX, San Diego, CA) were used, and as a secondary antibody, anti-rabbit IgG (Pierce, Rockford, IL) conjugated with horseradish peroxidase was used. For luminescence, ECL reagent (Amersham Pharmacia Biotech, Piscataway, NJ) was used, and the luminescence was detected by a LAS1000 imaging system (Fujifilm, Tokyo, Japan). The expression of the WNV-like particles between various recombinant expression vectors is shown in Fig. 2.

[0051]

Moreover, according to the method described in Kojima, A.,

A. Yasuda, H. Asanuma, T. Ishikawa, K. Yasui, T. Kurata. J. Viorl 77: 8745-55 (2003), E protein antigens secreted in the culture media was compared between the above-mentioned respective recombinant expression vectors by sandwich ELISA method using a monoclonal antibody (402) to E protein antigen.

Specifically, E protein antigens were bound using a microplate (Corning Incorporated Life Sciences, Acton, MA) coated with 402 antibody. The bound E protein antigens were detected using 402 antibody (TMB; DAKO Corp., Carpinteria, CA) conjugated with horseradish peroxidase as a substrate. Comparison of secretion amounts of the WNV-like particles between various recombinant expression vectors is shown in Fig. 3.

[0052]

In addition, recombinant expression vectors (pWPME388A, pWPME406A) obtained by adding alanine to immediately after starting amino acid of recombinant expression vectors pWPME1 and pWPME7 were produced, and the expression level was compared between various recombinant expression vectors by the above-mentioned method. The results are shown in Figs. 4 and 5. From these results, it was found that the length of signal sequence greatly affects the expression and secretion of WNV-like particles.

[0053]

Example 3 Observation image of WNV-like particles by electron microscope

The recombinant expression vector pWPME12 obtained in Example 1 was introduced into HEK293T cells, and the medium was collected 48 hr later. The collected medium was filtered (0.45 μm, Millipore) and purified (Biomax-100 membrane filter, Millipore). WNV-like particles obtained by removing protein small molecules having a molecular weight of less than 100 kDa were applied to 10-45% (w/w) sucrose density gradient centrifugation (150,000×g, 14 hr). The density and E protein antigen amount of sample fractions 1 - 17 were measured

according to the above-mentioned ELISA method. As the result, a peak of E protein antigen was observed in sample fraction 9 having a density of 1.11 and sample fraction 4 having a density of 1.16 (Fig. 6).

[0054]

A low molecular weight mixture was removed from E protein antigen contained in the two sample fractions 4 and 9 obtained in the above by Sephadex G-25 column (Hiprep; Amersham Pharmacia Bioteck) chromatography, and E protein antigen was eluted with phosphate buffered saline and preserved at 4°C. The obtained purified antigen was subjected to negative staining with sodium phosphotungstate using a grid coated with copper formvar. The specimen was observed with a JOEL 1200 Ex electron microscope (Hitachi, Tokyo, Japan). As a result, sample fraction 9 was confirmed to have many particles with diameter of about 30 nm and having a spherical structure (Fig. 7, right side). On the other hand, in sample fraction 4, particles (diameter about 25 nm) having various forms and coated with a staining fluid were occasionally seen (Fig. 7, left side).

[0055]

Example 4 Epitope analysis for inducing WNV-like particle neutralization antibody

The recombinant expression vector pWPME12 obtained in Example 1 was introduced into HEK293T cells, and the obtained E protein antigen was subjected to sucrose density gradient centrifugation to give sample fractions 1 - 11 (Fig. 8E). According to the above-mentioned ELISA method, sample fractions with high content of E protein antigen having neutralization epitope were found by using WNY-11 antibody (WNV neutralization antibody) and 402 antibody (non-neutralization antibody) as antibody for capture or antibody for detection. As a result, when the WNY-11 antibody was used as an antibody for capture and an antibody for detection, a peak could be confirmed in sample fraction 3 (density 1.16)

and sample fraction 5 (density 1.11) (Fig. 8A). On the other hand, when the WNY-11 antibody and 402 antibody were used as antibody for capture and/or antibody for detection, the content of E protein antigen in sample fraction 5 decreased as compared to the use only of WNY-11 antibody (Fig. 8B-D). The neutralization epitope recognized by WNY-11 in sample fraction 3 seems to be less than that in sample fraction 5.

[0056]

Example 5 Consideration of induction property of neutralization antibody

Using plural fractions including sample fractions 3 and 5 prepared in Example 4, whether or not induction of neutralization antibody in mouse was possible was considered. To be specific, BALB/c mouse was intraperitoneally immunized twice with adjuvant-conjugated WNV-like particles (about 300 ng, 1/16 of weight of formulation of Japanese encephalitis vaccine to human) at 7 day intervals. At 3 weeks from the second immunization, neutralization antibody titer of the immunized mouse was determined by plaque decrease assay. As a result, the mouse immunized with purified WNV-like particles expressed neutralization antibody at a high level (Fig. 9).

[0057]

Example 6 Establishment of transformed cell

[0058]

(1) Establishment of adherent transformed cell

Single layer CHO-K1 cells after 7 passages were spread the day before transfection on φ60 mm dish at 1:5, 1:10 or 1:20 split, and the next day, pWPME12 (5 μg) was introduced into 30 - 40% confluent cells. After culture for 2 days before formation of a single layer, the cells were split at 1:5, 1:10 or 1:20, and cultured in a selection medium containing Blasticidin S (BS) (10 μg/ml) in a φ100 mm dish for 8 days (medium was changed once). 24 colonies grown in the selection medium were taken by a penicillin cup method, transferred to a 24 well plate and further cultured in a selection medium

(hereafter selection medium containing BS). Thereafter, the cells were gradually grown by passaging in a 24-well plate, a 6-well plate and a φ100 mm dish, and the obtained cells were cryopreserved. The amount of WNV E protein in the medium cultured in the 24-well plate and 6-well plate was measured according to ELISA, and colony #22 was selected, which showed the maximum production amount. Then, the colony #22 cells were subcloned by a limit dilution method according to a conventional method, and clone #22.6 cells and sub-clone #22.6.6 cells were selected, which showed the maximum WNV E protein production levels according to ELISA. Fig. 10 shows comparison results of expression and secretion of E protein antigen between parent cells and each clone by Western blot.

[0059]

(2) Acclimation of adherent transformed cell in serum-free medium

The adherent clone #22.6 cells or sub-clone #22.6.6 cells cryopreserved in the above-mentioned (1) were placed back in F12K medium (GIBCO) containing 10 μg/ml BS and 10% FBS, and the growth was confirmed by maintaining for 3 - 4 passages. F12K medium containing 10 μg/ml BS and 10% FBS and CHO-S-SFM II medium containing 10 μg/ml BS were mixed in the same amount and the cells were cultured in the resulting medium (finally containing 5% FBS) for two passages to initiate acclimation in a serum-free medium. Thereafter, the cells were cultured in a medium containing the same amount of CHO-S-SFM II medium (GIBCO) containing 10 μg/ml BS for two or more passages, and the FBS concentration and ratio of F12K medium were gradually decreased by half.

When the cells were detached from the culture flask by pipetting with PBS(-) even without addition of a trypsin-EDTA solution, they were continuously passaged only in CHO-S-SFM II medium. When the growth of the cells was fine (growth to about 5 - 10 × 10^5/ml with viable cells of not less than 95% by passage of 1 - 2 x 10^5/ml) and the cells were capable of

passage in the medium by pipetting, they were judged to have acclimated in a serum-free medium. The cells were further passaged, and complete acclimation (Full-Adapt) cells that showed growth of 1 x 10^6/ml consecutively for 3 or more times and best acclimation (Best-Adapt) cells that showed growth of not less than 1 x 10^6/ml for 10 consecutive times were cryopreserved (#22.6S and #22.6.6S cells).

In addition, since the adherent CHO cells showed radical property change of becoming non-adherent cells during the acclimation process in a serum-free medium, they had a possibility of decreasing the production amount of antigen protein in this process. Thus, acclimated non-adherent cells #22.6S were cloned and re-cloned. The cloning method followed a conventional method. However, since the growth of the acclimation non-adherent cells showed high dependency on the cell density, the number of cells sown per well was set to 3, 10, 30 and 100. Respective growing clones were preferentially selected from the wells with less number of cells sown, and the production amount of protein was examined by ELISA and IFA. However, there was not found any noticeable difference between the selected plural clones and sub-clones.

Industrial Applicability

[0060]

The present invention can efficiently secrete and produce VLP of WNV, and can stably supply subunit vaccines of WNV containing the VLP as an active ingredient at a low cost and with safety. In addition, the WNV VLP secretion expression vector of the present invention itself can be used as a DNA vaccine of WNV.

This application is based on a patent application No. 2007-290169 filed in Japan (filing date: November 7, 2007), the contents of which are incorporated in full herein.

Claims

1. A signal peptide comprising an amino acid sequence the same or substantially the same as the amino acid sequence of the following (a) or (b)

(a) the amino acid sequence shown by SEQ ID NO: 1

(b) a partial amino acid sequence of the amino acid sequence of said (a), comprising at least the amino acid sequence shown by amino acid NOs 11 - 25.

2. An isolated nucleic acid substantially consisting of a base sequence encoding the signal peptide of claim 1.

3. A West Nile virus-like particle expression vector comprising a nucleic acid comprising a base sequence encoding the signal peptide of claim 1, and prM protein and E protein derived from West Nile virus.

4. A transformant obtained by transformation with the vector of claim 3.

5. A method of producing a West Nile virus-like particle, comprising culturing the transformant of claim 4, and recovering a virus-like particle secreted in a medium.

6. A West Nile virus-like particle obtained by the method of claim 5.

7. A West Nile virus vaccine comprising the particle of claim 6 as an active ingredient.

8. A West Nile virus vaccine comprising the vector of claim 3 as an active ingredient.

9. A method of preventing West Nile fever, comprising

administering an effective amount of the vector of claim 3 to a subject.

10. A method of preventing West Nile fever, comprising administering an effective amount of the virus-like particles of claim 6 to a subject.

SEQUENCE LISTING

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The Research Foundation for Microbial Diseases of Osaka University

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ctg ggg gct ctc ctg ttg tgg atg ggc atc aat gct cgt gat agg tcc     1440
Leu Gly Ala Leu Leu Leu Trp Met Gly Ile Asn Ala Arg Asp Arg Ser
465                 470                 475                 480

ata gct ctc acg ttt ctc gca gtt gga gga gtt ctg ctc ttc ctc tcc     1488
Ile Ala Leu Thr Phe Leu Ala Val Gly Gly Val Leu Leu Phe Leu Ser
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gtg aac gtg cac gct                                                1503
Val Asn Val His Ala
            500
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Gly Ala Thr Trp Val Asp Leu Val Leu Glu Gly Asp Ser Cys Val Thr
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Ile Met Ser Lys Asp Lys Pro Thr Ile Asp Val Lys Met Met Asn Met
        35                  40                  45
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Glu Ala Ala Asn Leu Ala Glu Val Arg Ser Tyr Cys Tyr Leu Ala Thr
50 55 60

Val Ser Asp Leu Ser Thr Lys Ala Ala Cys Pro Thr Met Gly Glu Ala
65 70 75 80

His Asn Asp Lys Arg Ala Asp Pro Ala Phe Val Cys Arg Gln Gly Val
85 90 95

Val Asp Arg Gly Trp Gly Asn Gly Cys Gly Leu Phe Gly Lys Gly Ser
100 105 110

Ile Asp Thr Cys Ala Lys Phe Ala Cys Ser Thr Lys Ala Ile Gly Arg
115 120 125

Thr Ile Leu Lys Glu Asn Ile Lys Tyr Glu Val Ala Ile Phe Val His
130 135 140

Gly Pro Thr Thr Val Glu Ser His Gly Asn Tyr Ser Thr Gln Val Gly
145 150 155 160

Ala Thr Gln Ala Gly Arg Phe Ser Ile Thr Pro Ala Ala Pro Ser Tyr
165 170 175

Thr Leu Lys Leu Gly Glu Tyr Gly Glu Val Thr Val Asp Cys Glu Pro
180 185 190

Arg Ser Gly Ile Asp Thr Asn Ala Tyr Tyr Val Met Thr Val Gly Thr
195 200 205

Lys Thr Phe Leu Val His Arg Glu Trp Phe Met Asp Leu Asn Leu Pro
210 215 220

Trp Ser Ser Ala Gly Ser Thr Val Trp Arg Asn Arg Glu Thr Leu Met
225 230 235 240

Glu Phe Glu Glu Pro His Ala Thr Lys Gln Ser Val Ile Ala Leu Gly
245 250 255

Ser Gln Glu Gly Ala Leu His Gln Ala Leu Ala Gly Ala Ile Pro Val
260 265 270

Glu Phe Ser Ser Asn Thr Val Lys Leu Thr Ser Gly His Leu Lys Cys
275 280 285

Arg Val Lys Met Glu Lys Leu Gln Leu Lys Gly Thr Thr Tyr Gly Val
290 295 300

Cys Ser Lys Ala Phe Lys Phe Leu Gly Thr Pro Ala Asp Thr Gly His
305 310 315 320

Gly Thr Val Val Leu Glu Leu Gln Tyr Thr Gly Thr Asp Gly Pro Cys
325 330 335

Lys Val Pro Ile Ser Ser Val Ala Ser Leu Asn Asp Leu Thr Pro Val

340 345 350

Gly Arg Leu Val Thr Val Asn Pro Phe Val Ser Val Ala Thr Ala Asn
355 360 365

Ala Lys Val Leu Ile Glu Leu Glu Pro Pro Phe Gly Asp Ser Tyr Ile
370 375 380

Val Val Gly Arg Gly Glu Gln Gln Ile Asn His His Trp His Lys Ser
385 390 395 400

Gly Ser Ser Ile Gly Lys Ala Phe Thr Thr Thr Leu Lys Gly Ala Gln
405 410 415

Arg Leu Ala Ala Leu Gly Asp Thr Ala Trp Asp Phe Gly Ser Val Gly
420 425 430

Gly Val Phe Thr Ser Val Gly Lys Ala Val His Gln Val Phe Gly Gly
435 440 445

Ala Phe Arg Ser Leu Phe Gly Gly Met Ser Trp Ile Thr Gln Gly Leu
450 455 460

Leu Gly Ala Leu Leu Leu Trp Met Gly Ile Asn Ala Arg Asp Arg Ser
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Val Asn Val His Ala
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