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(54) Title: METHOD FOR GENERATING A PARVOVIRUS B19 VIRUS-LIKE PARTICLE

(57) Abstract: The invention provides a process for generating parvovirus VP1/VP2 virus like particles (VLPs). The invention further provides methods for purification of the parvovirus VLPs and immunogenic compositions that contain the VLPs. The invention also includes recombinant nucleic acid molecules that encode parvovirus VP1 and VP2, and host cells that contain the recombinant nucleic acids.



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**METHOD FOR GENERATING A PARVOVIRUS B19 VIRUS-LIKE PARTICLE**

**[0001]** This application claims the benefit of United States provisional patent application 61/321856, filed April 7, 2010, the complete contents of which are incorporated herein by reference.

**[0002]** Parvovirus B19 produces mild, self-limiting illness in immunocompetent, hematologically normal individuals, however, it may cause serious illness in people with sickle-cell disease or other types of chronic anemia. In such people, parvovirus B19 can cause an acute, severe anemia. The ill person may be pale, weak, and tired. Furthermore, people who have problems with their immune systems may also develop a chronic anemia with parvovirus B19 infection that requires medical treatment. People who have leukemia or cancer, those born with immune deficiencies, who have received an organ transplant, or who have human immunodeficiency virus (HIV) infection are at risk for serious illness due to parvovirus B19 infection. Hematologically normal individuals who contract parvovirus B19 initially as an adult can present with arthralgias that last for weeks. Parvovirus B19 infection in pregnant women is also associated with hydrops fetalis due to severe fetal anemia, sometimes leading to miscarriage or stillbirth. Acquisition of parvovirus B19 by blood transfusion in those with acute anemias due to blood loss or other causes can prevent correction of the anemia by hematopoiesis.

**[0003]** Parvovirus B19 most commonly causes a childhood exanthem called fifth disease or erythema infectiosum, commonly known as slapped cheek syndrome. A person infected with parvovirus B19 is contagious during the early part of the illness, before the rash appears. Parvovirus B19 has been found in the respiratory secretions (e.g., saliva, sputum or nasal mucus) of infected persons before the onset of rash, when they appear to "just have a cold." The virus is probably spread from person to person by direct contact with those secretions. Parvovirus B19 is now considered a serious disease because of the complications that can occur, especially for pregnant women and their fetuses, or for people with unhealthy immune systems.

**[0004]** The symptoms of parvovirus B19 infection include a mild nonspecific prodromal illness that may consist of fever, malaise, headache, myalgia, nausea, and rhinorrhea. Symptoms typically begin 5-7 days after initial infection. These symptoms correspond to the initial viremia and dissipate in 2-3 days. Approximately one week later, a

bright red macular exanthema appears on the cheeks and is often associated with circumoral pallor. A diffuse maculopapular rash can appear 1-4 days later and fades to a lacy erythematous rash, which may be pruritic and may spread gradually toward the distal extremities.

**[0005]** Parvovirus B19 belongs to the Erythroviruses genus of the Parvoviridae family of small DNA viruses. It is a non-enveloped, icosahedral virus that contains a single-stranded linear DNA genome. The parvovirus B19 virion is 20-25 nm in diameter and has a genome of 5.6 kb (Clewley, 1984, Cotmore & Tattersall, 1984). The parvovirus B19 capsid consists of an 83 kDa minor structural protein, VP1, and a 58 kDa major structural protein, VP2. It has a non-segmented single stranded DNA genome surrounded by a protein shell containing two structural proteins, VP1 and VP2 in a ~5% to ~95% ratio (Ozawa et al., 1987). The sequences of the two proteins are co-linear, with VP2 being identical to the carboxyl-terminal region of VP1; however, VP1 comprises an additional 227 amino acids at the amino-terminus. Long lasting antibody responses are directed to both VP1 and VP2 proteins and thus these proteins alone are expected to raise a significant immune response.

**[0006]** The expression and purification in insect cells of parvovirus B19 VP1/VP2 VLPs has been previously described (U.S. Patent No. 5,508,186). Despite what is known about parvovirus B19, there is no vaccine or practical medicine that prevents parvovirus B19 infection. Human immunoglobulin-containing preparations are sometimes used to treat chronic parvovirus B19 infections in immunodeficient individuals. However, immunoglobulin injections are not sufficiently practical, long lasting, or affordable for widespread or routine prophylactic use. Excluding those with fifth disease from social interaction, e.g., at work, child care centers, or schools, is not an effective way to prevent the spread of the virus, because individuals are contagious well before they develop the signatory rash. Thus, a need exists for an improved parvovirus B19 vaccine development approach that allows for efficient preparation of parvovirus B19 immunogenic materials.

## SUMMARY OF THE INVENTION

**[0007]** The present invention, in part, provides methods for generating parvovirus virus-like particles, such as parvovirus B19 virus-like particles. In some aspects, the methods of the invention comprise providing a host cell with a recombinant nucleic acid, such as an expression vector, comprising both parvovirus VP1 and parvovirus VP2 and maintaining the

host cell under conditions whereby the VP1 and VP2 proteins are expressed and assembled into parvovirus VLPs. In another aspect, the method of the invention comprises providing a bicistronic nucleic acid vaccine that leads to expression of parvovirus VP2 and lower levels of parvovirus VP1 in a host, such that VLPs form in the immunized host.

**[0008]** The invention relates to a method of producing a parvovirus virus like particle (VLP), comprising: (a) providing a host cell that contains a recombinant nucleic acid molecule that comprises a nucleotide sequence encoding parvovirus VP1 that is operably linked to a first control element, and a nucleotide sequence encoding parvovirus VP2 protein that is operably linked to a second control element, and (b) maintaining the host cell under conditions whereby the VP1 and VP2 proteins are expressed and assembled into VLPs. In a preferred method, VP1 is produced in lower abundance than VP2. Preferably, soluble VP1 is produced in lower abundance than soluble VP2. In a particular aspect, VLPs that contain VP1 and VP2 are produced. Preferably, the recombinant nucleic acid molecule is a bicistronic vector. In a particular aspect, the first control element is a variant of the second control element, and the variant comprises a modification that transcriptionally, translationally or transcriptionally and translationally decreases production of VP1 relative to VP2. For example, the first control element can comprise a deletion of at least a portion of the TATA box upstream of the nucleic acid encoding VP1, deletion of a junction site upstream of the nucleic acid encoding VP1, introduction of a transcription start or stop site upstream of the nucleic acid encoding VP1, or combinations thereof. In a particular aspect, the second control element is the ADH2/GAPDH promoter (SEQ ID NO: 19).

**[0009]** The host cell can be a yeast cell, insect cell, mammalian cell, avian cell, bacterium, Tetrahymena cell or combinations thereof. In one aspect, the host cells are maintained under culture conditions suitable for production of VP1 and VP2 and assembly of the proteins into VLPs. The method can further comprising isolating the VLPs. The VLPs can be isolated from host cell conditioned media, host cell lysate, host cell homogenate, or combinations thereof. Optionally, the isolated VLPs can be further purified. The isolated VLPs may be purified using a purification method selected from the group consisting of sucrose cushion, sucrose gradient centrifugation, chromatographic methods and combinations thereof.

**[0010]** The invention also relates to a parvovirus virus like particle (VLP) that contains VP1 and VP2 produced according to any of the methods described herein.

[0011] The invention also relates to an immunogenic composition comprising a parvovirus VLP.

[0012] The invention also relates to a recombinant nucleic acid comprising a nucleotide sequence encoding parvovirus VP1 that is operably linked to a first control element, and a nucleotide sequence encoding parvovirus VP2 protein that is operably linked to a second control element, wherein when equivalent amounts of the sequence that encodes VP1 and the sequence that encodes VP2 are present in a suitable host cell, VP1 is expressed in lower abundance relative to VP2. Preferably, soluble VP1 is produced in lower abundance than soluble VP2. In one aspect, the first control element is a weak promoter and the second control element is a strong promoter. The first control element can be a variant of the second control element, and the variant can comprise a modification that transcriptionally, translationally or transcriptionally and translationally decreases production of VP1 relative to VP2. The modification can be deletion of at least a portion of the TATA box upstream of the nucleic acid encoding VP1, deletion of a junction site upstream of the nucleic acid encoding VP1, introduction of a transcription start or stop site upstream of the nucleic acid encoding VP1, or combinations thereof. The second control element can be the ADH2/GAPDH promoter (SEQ ID NO: 19). The nucleic acid can be in the form of DNA, or RNA, and can be either single or double stranded. The recombinant nucleic acid can be a plasmid. The plasmid can include a detectable marker.

[0013] The invention also relates to a host cell comprising a recombinant nucleic acid as described herein. The recombinant nucleic acid can be integrated into the genome of the host cell or can be carried on an extra-chromosomal element. The host cell can be a yeast cell, insect cell, mammalian cell, avian cell, bacteria, Tetrahymena cells or combinations thereof.

[0014] The invention also relates to a method of using a parvovirus virus like particle (VLP) that contains VP1 and VP2, produced according to the method described herein for therapeutic and diagnostic purposes, for example as an antigen in a diagnostic kit.

#### **BRIEF DESCRIPTION OF THE DRAWINGS**

[0015] FIG. 1 is a diagram of yeast expression vector pBS24.1. This vector is suitable to produce VLPs in *S. cerevisiae*.

**[0016]** FIG. 2 is a diagram of a bicistronic yeast expression vector pCDC.7, based on pBS24.1. This vector is suitable for co-expression of parvovirus B19 VP1 and VP2 VLPs in *S. cerevisiae*.

**[0017]** FIGS. 3A-3D show the nucleotide sequence (SEQ ID NO: 16) of pCDC.7 used for co-expression of parvovirus B19 VP1 and VP2 VLP's in *S. cerevisiae*.

**[0018]** FIG. 4 shows the nucleotide sequence (SEQ ID NO: 17) of the optimized parvovirus B19 VP1.

**[0019]** FIG. 5 shows the nucleotide sequence (SEQ ID NO: 18) of the optimized parvovirus B19 VP2.

**[0020]** FIG. 6 shows the nucleotide sequence (SEQ ID NO: 19) of the ADH2/GAPDH promoter (as BamHI/HindIII fragment) used in the expression of parvovirus B19 VP2.

**[0021]** FIG. 7 shows the nucleotide sequence (SEQ ID NO: 20) of the "delta As" version of the ADH2/GAPDH promoter in which ten bases of GAPDH sequence has been replaced with 14 bases of parvovirus B19 native viral sequence (underlined) as it naturally occurs immediately upstream from the initiating methionine of VP1 (bold).

**[0022]** FIG. 8 shows the nucleotide sequence (SEQ ID NO: 21) of the ADH2/GAPDH promoter (as BamHI/MluI fragment) with TATA deletion used in the expression of parvovirus B19 VP1.

**[0023]** FIG. 9 shows the nucleotide sequence (SEQ ID NO: 22) of parvovirus B19 VP1 with 1SS (synthetic start-stop sequence, underlined) upstream from the junction between the ADH2/GAPDH promoter and the initiating methionine of VP1 (bold).

**[0024]** FIG. 10 shows the variation in relative amounts of soluble VP1 and soluble VP2 produced in yeast on a Western blot using antibodies to VP2. The upper band represents VP1 and the lower, darker band represents VP2. The labels describe what is in the plasmid or the type of modification that has been made. For example, lane 7 shows that one start/stop decreases the level of VP1 relative to VP2, and lane 8 shows two start/stops decrease the level even more.

[0025] FIG. 11 shows (A) Coomassie gels representing purified parvovirus B19 VLP and (B) electron micrographs of parvovirus B19 VLPs.

[0026] FIG. 12 shows three graphs that represent ELISA data which demonstrate the immunogenicity of parvovirus B19 VLPs. Highest titers were demonstrated with the MF59 formulations and were similar for each dose concentration.

[0027] FIG. 13 is a diagram of a VP1 delta TATA/VP2 construct. This construct is suitable for use in the protein purification and experiments described in this application.

[0028] FIG. 14 shows the amino acid sequences of parvovirus B19 VP1 (SEQ ID NO: 23) and VP2 (SEQ ID NO: 24). VP1 is 781 amino acids long, and has a molecular weight of 84 kDa. VP2 is 554 amino acids long and has a molecular weight of 58 kDa.

[0029] FIG. 15 shows (A) SDS-PAGE and (B) Western of the sucrose shelf fractions from VP1/VP2 VLP expression.

[0030] FIG. 16 shows a (A) Coomassie stained SDS-PAGE and (B)  $\alpha$ -VP1/VP2 antibody probed Western (B) analysis of fractions from a Capto Q elution of parvovirus B19 VP1/VP2 VLPs.

[0031] FIG. 17 shows a size exclusion chromatography (SEC) (S-200) chromatogram of purified VP1/VP2 VLPs.

## DETAILED DESCRIPTION OF THE INVENTION

[0032] The present invention is, in part, based on the discovery that although parvovirus B19 VP1 and VP2 can be expressed in the same host cell to produce VLPs that contain both proteins, production of VLPs is inefficient. It has now been discovered that when VP1 and VP2 are expressed in the same host cell (e.g., a yeast cell), the expression of VP1 suppresses the expression of VP2 and the formation of VLPs. This results in inefficient VLP formation, and production of increased amount of insoluble material.

[0033] The present invention provides a solution to the problem of inefficient production of VLPs that contain parvovirus VP1 and VP2 proteins. The method provides a better way to produce VLPs that contain parvovirus VP1 and VP2 proteins by producing VP1 and VP2 in a host cell, such that VP1 is produced in lower abundance relative to VP2.

Preferably, soluble VP1 is produced in lower abundance than soluble VP2. The invention provides recombinant nucleic acids that encode VP1 and VP2 and drive expression of the proteins in host cells in desired amounts. For example, the expression of VP1 and VP2 can be controlled by separate promoters, allowing for controlled expression of each protein at desired ratios. The invention also provides methods for producing parvovirus VLPs that contain VP1 and VP2 using host cells that contain the recombinant nucleic acids. These methods provide improved VLP production efficiency and reduce the variability and quality control issues associated with the prior methods that used separate vectors for expression of VP1 and VP2, and mixed them together to produce host cells, including variability in transfection efficiency (e.g., cell to cell variability and batch to batch variability), variability in the expression of VP1 relative to VP2, and suppression of VP2 expression by VP1.

[0034] Other aspects of the present invention include recombinant nucleic acids, host cells, methods for producing VLPs and immunogenic compositions that contain VLPs.

[0035] **I. Definitions**

[0036] All scientific and technical terms used in this application have meanings commonly used in the art unless otherwise specified. As used in this application, the following words or phrases have the meanings specified.

[0037] It must be noted that, as used in this specification and the appended claims, the singular forms "a", "an" and "the" include plural references unless the content clearly dictates otherwise. Thus, for example, reference to "a polynucleotide" includes a mixture of two or more such polynucleotides, and the like.

[0038] The term "about" in relation to a numerical value x means, for example, x+10%.

[0039] As used herein, the terms "parvovirus" refers to all parvoviruses associated with mammalian species (e.g., human, canine, chicken, feline, murine, porcine, raccoon, mink, kilham rat, lapine) and broadly to all genus of the *Parvoviridae* family (i.e., *Parvovirus* (e.g., canine parvovirus), *Dependovirus* (e.g., adeno-associated virus), *Erythrovirus* (e.g., parvovirus B19) and *Bocavirus*). Preferably, the parvovirus infects humans, i.e., is of the *Dependovirus*, *Erythrovirus*, or *Bocavirus* genus. Most preferably, the parvovirus is

parvovirus B19. In some embodiments, the parvovirus is from the *Parvovirus* genus. The term parvovirus also includes isolates not characterized at the time of filing.

**[0040]** The parvovirus B19 species is subdivided into three distinct genotypes (Gallinella et al., 2003; Hokynar et al., 2002; Nguyen et al. 2002; Servant et al. 2002). The nucleotide divergency between the genotypes is approximately 10% and in the promoter region more than 20%. All viruses previously known as B19V were classified as genotype 1. Genotype 2 is found relatively infrequently. When genotype 2 is found, it is identified at a much higher frequency in individuals older than approximately 40 years of age. Genotype 3 viruses cluster into two subtypes represented by the prototype strains V9 (Genbank accession no. AX003421) and D91.1 (Genbank accession no. AY083234) (Parsyan et al., 2007). Genotype 3 virus has been shown to be endemic in Ghana West Africa (Candotti et al., 2004) and may be present in a certain region of Brazil.

**[0041]** The terms "polypeptide" and "protein" refer to a polymer of amino acid residues and are not limited to a minimum length of the product. Thus, peptides, oligopeptides, dimers, multimers, and the like, are included within the definition. Both full-length proteins and fragments thereof are encompassed by the definition. The terms also include postexpression modifications of the polypeptide, for example, glycosylation, acetylation, phosphorylation and the like. Furthermore, for purposes of the present invention, a "polypeptide" refers to a protein which includes modifications, such as deletions, additions and substitutions (generally conservative in nature), to the native sequence, so long as the protein maintains the desired activity. These modifications may be deliberate, as through site-directed mutagenesis, or may be accidental, such as through mutations of hosts which produce the proteins or errors due to PCR amplification.

**[0042]** "Substantially purified" generally refers to isolation of a substance (compound, polynucleotide, protein, polypeptide, polypeptide composition) such that the substance comprises the majority percent of the sample in which it resides. Typically in a sample, a substantially purified component comprises 50%, preferably 80%-85%, more preferably 90-95% of the sample. Techniques for purifying polynucleotides and polypeptides of interest are well-known in the art and include, for example, ion-exchange chromatography, affinity chromatography and sedimentation according to density.

[0043] By "isolated" is meant, when referring to a polypeptide, that the indicated molecule is separate and discrete from the whole organism with which the molecule is found in nature or is present in the substantial absence of other biological macro-molecules of the same type. The term "isolated" with respect to a polynucleotide is a nucleic acid molecule devoid, in whole or part, of sequences normally associated with it in nature; or a sequence, as it exists in nature, but having heterologous sequences in association therewith; or a molecule disassociated from the chromosome.

[0044] "Recombinant" as used herein to describe a nucleic acid molecule means a polynucleotide of genomic, cDNA, viral, semisynthetic, or synthetic origin which, by virtue of its origin or manipulation, is not associated with all or a portion of the polynucleotide with which it is associated in nature. The term "recombinant" as used with respect to a protein or polypeptide means a polypeptide produced by expression of a recombinant polynucleotide. In general, the gene of interest is cloned and then expressed in transformed organisms, as described further below. The host organism expresses the foreign gene to produce the protein under expression conditions.

[0045] The term "transformation" refers to the insertion of an exogenous polynucleotide into a host cell, irrespective of the method used for the insertion. For example, direct uptake, transfection, transduction or f-mating are included. The exogenous polynucleotide may be maintained as a non-integrated vector, for example, a plasmid, or alternatively, may be integrated into the host genome.

[0046] "Recombinant host cells", "host cells," "cells", "cell lines," "cell cultures", and other such terms denoting microorganisms or higher eukaryotic cell lines cultured as unicellular entities refer to cells which can be, or have been, used as recipients for recombinant vector or other transferred DNA, and include the original progeny of the original cell which has been transfected.

[0047] A "coding sequence" or a sequence which "encodes" a selected polypeptide, is a nucleic acid molecule which is transcribed (in the case of DNA) and translated (in the case of mRNA) into a polypeptide *in vivo* when placed under the control of appropriate regulatory sequences (or "control elements"). The boundaries of the coding sequence can be determined by a start codon at the 5' (amino) terminus and a translation stop codon at the 3' (carboxy) terminus. A coding sequence can include, but is not limited to,

cDNA from viral, prokaryotic or eukaryotic mRNA, genomic DNA sequences from viral or prokaryotic DNA or RNA, and even synthetic DNA sequences. A transcription termination sequence may be located 3' to the coding sequence.

**[0048]** Typical “control elements,” include, but are not limited to, transcription promoters, transcription enhancer elements, transcription termination signals, polyadenylation sequences (located 3' to the translation stop codon), sequences for optimization of initiation of translation (located 5' to the coding sequence), and translation termination sequences.

**[0049]** The term “nucleic acid” includes DNA and RNA, and also their analogs, such as those containing modified backbones (e.g. phosphorothioates, etc.), and also peptide nucleic acids (PNA), etc. The invention includes nucleic acids comprising sequences complementary to those described above (e.g. for antisense or probing purposes).

**[0050]** “Operably linked” refers to an arrangement of elements wherein the components so described are configured so as to perform their usual function. Thus, a given promoter operably linked to a coding sequence is capable of effecting the expression of the coding sequence when the proper enzymes are present. The promoter need not be contiguous with the coding sequence, so long as it functions to direct the expression thereof. Thus, for example, intervening untranslated yet transcribed sequences can be present between the promoter sequence and the coding sequence and the promoter sequence can still be considered “operably linked” to the coding sequence.

**[0051]** “Encoded by” refers to a nucleic acid sequence which codes for a polypeptide sequence, wherein the polypeptide sequence or a portion thereof contains an amino acid sequence of at least 3 amino acids.

**[0052]** “Expression cassette” or “expression construct” refers to an assembly which is capable of directing the expression of the sequence(s) or gene(s) of interest. An expression cassette generally includes control elements, as described above, such as a promoter which is operably linked to (so as to direct transcription of) the sequence(s) or gene(s) of interest, and often includes a polyadenylation sequence as well. Within certain embodiments of the invention, the expression cassette described herein may be contained within a plasmid construct. In addition to the components of the expression cassette, the plasmid construct may also include, one or more selectable markers, a signal which allows the plasmid construct to exist as single-stranded DNA (e.g., a M13 origin of replication), at least

one multiple cloning site, and a “mammalian” origin of replication (e.g., a SV40 or adenovirus origin of replication).

**[0053]** “Purified polynucleotide” refers to a polynucleotide of interest or fragment thereof which is essentially free, e.g., contains less than about 50%, preferably less than about 70%, and more preferably less than about at least 90%, of the protein with which the polynucleotide is naturally associated. Techniques for purifying polynucleotides of interest are well-known in the art and include, for example, disruption of the cell containing the polynucleotide with a chaotropic agent and separation of the polynucleotide(s) and proteins by ion-exchange chromatography, affinity chromatography and sedimentation according to density.

**[0054]** A “vector” is capable of transferring nucleic acid sequences to target cells (e.g., viral vectors, non-viral vectors, particulate carriers, and liposomes). Typically, “vector construct,” “expression vector,” and “gene transfer vector,” mean any nucleic acid construct capable of directing the expression of a nucleic acid of interest and which can transfer nucleic acid sequences to target cells. Thus, the term includes cloning and expression vehicles, as well as viral vectors.

**[0055]** As used herein, the term “epitope” generally refers to the site on an antigen which is recognised by a T-cell receptor and/or an antibody. Preferably it is a short peptide derived from or as part of a protein antigen. However the term is also intended to include peptides with glycopeptides and carbohydrate epitopes and protein surfaces comprised of peptides (with or without modifications, such as glycosylation) that are discontinuous in primary sequence but which are located in proximity to each other in the final folded structure. Several different epitopes may be carried by a single antigenic molecule. An epitope can be made from more than one molecule that associates covalently or non-covalently with one or more other molecules. The term “epitope” also includes modified sequences of amino acids or carbohydrates which stimulate responses which recognise the whole organism. It is advantageous if the selected epitope is an epitope of an infectious agent, which causes the infectious disease.

**[0056]** As used herein, the term “T cell epitope” refers generally to those features of a peptide structure which are capable of inducing a T cell response and a “B cell epitope”

refers generally to those features of a peptide structure which are capable of inducing a B cell response.

**[0057]** An “immunological response” to an antigen or composition is the development in a subject of a humoral and/or a cellular immune response to an antigen present in the composition of interest. For purposes of the present invention, a “humoral immune response” refers to an immune response mediated by antibody molecules, while a “cellular immune response” is one mediated by T-lymphocytes and/or other white blood cells. One important aspect of cellular immunity involves an antigen-specific response by cytolytic T-cells (“CTL”s). CTLs have specificity for peptide antigens that are presented in association with proteins encoded by the major histocompatibility complex (MHC) and expressed on the surfaces of cells. CTLs help induce and promote the destruction of intracellular microbes, or the lysis of cells infected with such microbes. Another aspect of cellular immunity involves an antigen-specific response by helper T-cells. Helper T-cells act to help stimulate the function, and focus the activity of, nonspecific effector cells against cells displaying peptide antigens in association with MHC molecules on their surface. A “cellular immune response” also refers to the production of cytokines, chemokines and other such molecules produced by activated T-cells and/or other white blood cells, including those derived from CD4+ and CD8+ T-cells.

**[0058]** An immunogenic composition or vaccine that elicits a cellular immune response may serve to sensitize a vertebrate subject by the presentation of antigen in association with MHC molecules at the cell surface. The cell-mediated immune response is directed at, or near, cells presenting antigen at their surface. In addition, antigen-specific T-lymphocytes can be generated to allow for the future protection of an immunized host.

**[0059]** The ability of a particular antigen to stimulate a cell-mediated immunological response may be determined by a number of assays, such as by lymphoproliferation (lymphocyte activation) assays, CTL cytotoxic cell assays, or by assaying for T-lymphocytes specific for the antigen in a sensitized subject. Such assays are well known in the art. See, e.g., Erickson et al., *J. Immunol.* (1993) 151:4189-4199; Doe et al., *Eur. J. Immunol.* (1994) 24:2369-2376. Recent methods of measuring cell-mediated immune response include measurement of intracellular cytokines or cytokine secretion by T-cell populations, or by measurement of epitope specific T-cells (e.g., by the tetramer technique)(reviewed by McMichael, A. J., and O'Callaghan, C. A., *J. Exp. Med.* 187(9)1367-

1371, 1998; Mcheyzer-Williams, M. G., et al., *Immunol. Rev.* 150:5-21, 1996; Lalvani, A., et al., *J. Exp. Med.* 186:859-865, 1997).

**[0060]** Thus, an immunological response as used herein may be one that stimulates the production of antibodies (e.g., neutralizing antibodies that block bacterial toxins and pathogens such as viruses entering cells and replicating by binding to toxins and pathogens, typically protecting cells from infection and destruction). The antigen of interest may also elicit production of CTLs. Hence, an immunological response may include one or more of the following effects: the production of antibodies by B-cells; and/or the activation of suppressor T-cells and/or memory/effector T-cells directed specifically to an antigen or antigens present in the composition or vaccine of interest. These responses may serve to neutralize infectivity, and/or mediate antibody-complement, or antibody dependent cell cytotoxicity (ADCC) to provide protection to an immunized host. Such responses can be determined using standard immunoassays and neutralization assays, well known in the art. (See, e.g., Montefiori et al. (1988) *J Clin Microbiol.* 26:231-235; Dreyer et al. (1999) *AIDS Res Hum Retroviruses* (1999) 15(17):1563-1571). The innate immune system of mammals also recognizes and responds to molecular features of pathogenic organisms via activation of Toll-like receptors and similar receptor molecules on immune cells. Upon activation of the innate immune system, various non-adaptive immune response cells. are activated to, e.g., produce various cytokines, lymphokines and chemokines. Cells activated by an innate immune response include immature and mature dendritic cells of the monocyte and plasmacytoid lineage (MDC, PDC), as well as gamma, delta, alpha and beta T cells and B cells and the like. Thus, the present invention also contemplates an immune response wherein the immune response involves both an innate and adaptive response.

**[0061]** An “immunogenic composition” is a composition that comprises an antigenic molecule where administration of the composition to a subject results in the development in the subject of a humoral and/or a cellular immune response to the antigenic molecule of interest.

**[0062]** The terms “immunogenic” protein or polypeptide refer to an amino acid sequence which elicits an immunological response as described above. An “immunogenic” protein or polypeptide, as used herein, includes the full-length natural or recombinant sequence of the protein in question, including the precursor and mature forms, analogs thereof, or immunogenic fragments thereof.

**[0063]** A parvovirus polynucleotide, oligonucleotide, nucleic acid, protein, polypeptide, or peptide, as defined above, is a molecule derived from a parvovirus, respectively, including, without limitation, any of the various isolates of parvovirus. The molecule need not be physically derived from the particular isolate in question, but may be synthetically or recombinantly produced.

**[0064]** In particular, the parvovirus B19 genome contains three open reading frames: a non-structural 77 kDa protein, NS1, is encoded by nucleotides 436-2451; the minor structural protein, VP1 is encoded by nucleotides 2444-4787, and the major structural protein, VP2, is encoded by nucleotides 3125-4787 (Corcoran *et al.*, *J. Med. Microb.*, 2004). Parvovirus B19 uses a single promoter, p6, which is capable of expressing structural and non-structural genes differentially (Blundell *et al.*, 1987, Ozawa *et al.*, 1987). Although the foregoing numbering is relative to the nucleotide sequence of the parvovirus B19 genome, it is to be understood that the corresponding positions in sequences obtained from other genotypes and isolates of parvovirus are also intended to be encompassed by the present invention. Any one of the nucleic acids encoding VP1 or VP2, as well as variants thereof, such as immunogenic fragments thereof, and polypeptides encoded by such nucleic acids can be used in the practice of the invention.

**[0065]** As used herein, the terms “minor structural protein” or “minor structural polypeptide” or “minor capsid protein” or “minor capsid polypeptide” or “VP1” in reference to a parvovirus refer to a polypeptide comprising a sequence homologous or identical to the ORF2-encoded polypeptide of a parvovirus, and includes sequences displaying at least about 80-100% sequence identity thereto, including any percent identity within these ranges, such as 81, 82, 83, 84, 85, 86, 87, 88, 89, 90, 91, 92, 93, 94, 95, 96, 97, 98, 99, 100% sequence identity thereto.

**[0066]** As used herein, the terms “major structural protein” or “major structural polypeptide” or “major capsid protein” or “major capsid polypeptide” or “VP2” in reference to a Parvovirus refer to a polypeptide comprising a sequence homologous or identical to the ORF3-encoded polypeptide of a Parvovirus, and include sequences displaying at least about 80-100% sequence identity thereto, including any percent identity within these ranges, such as 81, 82, 83, 84, 85, 86, 87, 88, 89, 90, 91, 92, 93, 94, 95, 96, 97, 98, 99, 100% sequence identity thereto.

**[0067]** As used herein, the term “virus-like particle” or “VLP” refers to a nonreplicating, noninfectious viral shell that contains a viral capsid but lacks all or part of the viral genome, in particular, the replicative components of the viral genome. VLPs are generally composed of one or more viral proteins, such as, but not limited to those proteins referred to as capsid, coat, shell, surface, structural proteins (e.g., VP1, VP2). Parvovirus VLPs can form spontaneously upon recombinant expression of VP2 in an appropriate expression system. Methods for producing particular VLPs are known in the art and discussed more fully below. The presence of VLPs following recombinant expression of viral proteins can be detected using conventional techniques known in the art, such as by electron microscopy, biophysical characterization, and the like. For example, VLPs can be isolated by density gradient centrifugation and/or identified by characteristic density banding. Alternatively, cryoelectron microscopy can be performed on vitrified aqueous samples of the VLP preparation in question, and images recorded under appropriate exposure conditions.

**[0068]** As used herein, a “biological sample” refers to a sample of tissue or fluid isolated from a subject, including but not limited to, for example, blood, plasma, serum, fecal matter, urine, bone marrow, bile, spinal fluid, lymph fluid, samples of the skin, external secretions of the skin, respiratory, intestinal, and genitourinary tracts, tears, saliva, milk, blood cells, organs, biopsies and also samples of in vitro cell culture constituents including but not limited to conditioned media resulting from the growth of cells and tissues in culture medium, e.g., recombinant cells, and cell components. In particular, parvovirus may be obtained from biological samples such as aerosol or respiratory secretions or blood from individuals infected with the viruses.

**[0069]** By “subject” is meant any member of the subphylum chordata, including, without limitation, humans and other primates, including non-human primates such as chimpanzees and other apes and monkey species; farm animals such as cattle, sheep, pigs, goats and horses; domestic mammals such as dogs and cats; laboratory animals including rodents such as mice, rats and guinea pigs; birds, including domestic, wild and game birds such as chickens, turkeys and other gallinaceous birds, ducks, geese, and the like. The term does not denote a particular age. Thus, both adult and newborn individuals are intended to be covered.

**[0070]** By “therapeutically effective amount” in the context of the immunogenic compositions is meant an amount of an immunogen (e.g., immunogenic polypeptide, fusion

protein, polyprotein, VLP, or nucleic acid encoding an antigen) which will induce an immunological response, either for antibody production or for treatment or prevention of parvovirus infection. Such a response will generally result in the development in the subject of an antibody-mediated and/or a secretory or cellular immune response to the composition. Usually, such a response includes but is not limited to one or more of the following effects; the production of antibodies from any of the immunological classes, such as immunoglobulins A, D, E, G or M; the proliferation of B and T lymphocytes; the provision of activation, growth and differentiation signals to immunological cells; expansion of helper T cell, suppressor T cell, and/or cytotoxic T cell and/or  $\gamma\delta$ T cell populations.

[0071] For purposes of the present invention, an “effective amount” of an adjuvant will be that amount which enhances an immunological response to a coadministered antigen or nucleic acid encoding an antigen.

[0072] As used herein, “treatment” refers to any of (i) the prevention of infection or reinfection, as in a traditional vaccine, (ii) the reduction or elimination of symptoms, and (iii) the substantial or complete elimination of the pathogen in question. Treatment may be effected prophylactically (prior to infection) or therapeutically (following infection).

## II. MODES OF CARRYING OUT THE INVENTION

[0073] Before describing the present invention in detail, it is to be understood that this invention is not limited to particularly exemplified molecules or process parameters as such may, of course, vary. It is also to be understood that the terminology used herein is for the purpose of describing particular embodiments of the invention only, and is not intended to be limiting. In addition, the practice of the present invention will employ, unless otherwise indicated, conventional methods of virology, microbiology, molecular biology, recombinant DNA techniques and immunology all of which are within the ordinary skill of the art. Such techniques are explained fully in the literature. See, e.g., Sambrook, et al., *Molecular Cloning: A Laboratory Manual* (3rd Edition, 2000); *DNA Cloning: A Practical Approach*, vol. I & II (D. Glover, ed.); *Oligonucleotide Synthesis* (N. Gait, ed., 1984); *A Practical Guide to Molecular Cloning* (1984); and *Fundamental Virology*, 4th Edition, 2001 (B. N. Fields and D. M. Knipe, eds.); *Handbook of Experimental Immunology*, Vols. I-IV (D. M. Weir and C. C. Blackwell eds., Blackwell Scientific Publications); A. L. Lehninger, *Biochemistry* (Worth Publishers, Inc., current edition); *Methods In Enzymology* (S. Colowick and N. Kaplan eds.,

Academic Press, Inc.). Although a number of methods and materials similar or equivalent to those described herein can be used in the practice of the present invention, the preferred materials and methods are described herein. In addition, all publications, patents and patent applications cited herein, whether supra or infra, are hereby incorporated by reference in their entireties.

### A. Nucleic Acids

[0074] The invention relates to recombinant nucleic acid molecules that encode parvovirus VP1 and parvovirus VP2. The encoded VP1 and VP2 can be from any desired parvovirus, or any desired combination. Preferably, the encoded VP1 and VP2 are from a parvovirus that infects humans, *i.e.*, a parvovirus of the *Dependovirus*, *Erythrovirus*, or *Bocavirus* genus. Most preferably, the parvovirus is parvovirus B19. In some aspects, the sequence that encodes VP1 and the sequence that encodes VP2, and their respective control elements, are on separate nucleic acid molecules, for example, on separate vectors. Preferably, the sequence that encodes VP1 and the sequence that encodes VP2, and their respective control elements, are components of a single nucleic acid molecule, such as a bicistronic vector.

[0075] The parvovirus VP1 and VP2 proteins can have the same or substantially the same amino acid sequence as the naturally occurring protein, such as the amino acid sequence of naturally occurring VP1 of parvovirus B19. If desired, in some embodiments, the amino acid sequence can differ from a naturally occurring sequence by one or more amino acid substitutions, additions and/or deletions. Preferably, less than about 10% or less than about 5% of the amino acid residues in VP1 or VP2 are substituted, added or deleted. Preferably, the amino acid sequence of the VP1 or VP2 protein is at least about 80%, at least about 85%, at least about 90%, at least about 95%, at least about 96%, at least about 97%, at least about 98%, or at least about 99% identical to a naturally occurring parvovirus VP1 or VP2 protein. Amino acid sequence variation is generally permitted in areas that are not conserved among VP1 or VP2 proteins, and therefore does not interfere with the ability of the proteins to form VLPs. If desired, mutations (e.g., amino acid substitutions, additions or deletions) can be introduced to alter certain functions, for example, VP1 can be mutated to inactivate its phospholipase activity. For example, the amino acid sequence of VP1 may contain a point mutation (e.g., His153Ala), or any of the mutations described in WO 06/032697, EP 1791858 or US 20070286870.

[0076] The recombinant nucleic acids of the invention provide independent regulation of transcription and/or translation of VP1 and VP2, so that when equivalent amounts of the sequence that encodes VP1 and the sequence that encodes VP2 are present in a suitable host cell, VP1 is expressed in lower abundance relative to VP2. Preferably, soluble VP1 is produced in lower abundance than soluble VP2. In some embodiments, the codon usage in the nucleotide sequence that encodes VP1 and in the nucleotide sequence that encodes VP2 are selected so that when equivalent amounts of the sequence that encodes VP1 and the sequence that encodes VP2 are present in a suitable host cell, VP1 is expressed in lower abundance relative to VP2. Preferably, soluble VP1 is produced in lower abundance than soluble VP2. This can be accomplished, for example, using a codon optimized nucleotide sequence that encodes VP2 and a nucleotide sequence that is not codon optimized, or that is codon deoptimized, that encodes VP1.

[0077] In other embodiments, the nucleotide sequence that encodes VP1 and the nucleotide sequence that encodes VP2 are operably linked to different control elements, so that when equivalent amounts of the sequence that encodes VP1 and the sequence that encodes VP2 are present in a suitable host cell, VP1 is expressed in lower abundance relative to VP2. Preferably, soluble VP1 is produced in lower abundance than soluble VP2. The control elements for VP1 and VP2 expression are selected so that VP1 is produced by the desired host cell in lower abundance than VP2 when equivalent amounts of sequences that encode VP1 and VP2 are present in the desired host cell. This is generally accomplished using control elements that affect transcription, translation, or transcription and translation of VP1 and VP2. For example, the nucleotide sequence encoding VP2 can be operably linked to a strong promoter, and the nucleotide sequence encoding VP1 can be operably linked to a weak promoter. The promoters used may be any promoters that allows for independent regulation of VP1 and VP2 expression. An exemplary strong promoter for expression in yeast is the ADH2/GAPDH promoter, which is incorporated into pBS24.1 (FIG. 1). An exemplary weak promoter for expression in yeast is the YPTI constitutively active promoter. *See, Sears et al., Yeast* 14:783-790 (1998). Other suitable strong and weak promoters for use in yeasts or other host cells are known in the art. For example, a strong promoter for expression in mammalian cells is the CMV promoter. An exemplary strong promoter for expression in bacteria is the recA promoter. An exemplary weak promoter for expression in bacteria is the araBAD promoter.

**[0078]** In some embodiments, a nucleotide sequence encoding VP2 is operably linked to a promoter suitable for expression in a desired host cell, and a nucleotide sequence encoding VP1 is operably linked to a variant of that promoter, such as an expression de-optimized variant. The variant promoter can be modified relative to the parental promoter in a variety of ways to reduce the expression of VP1 relative to VP2 (e.g., soluble VP1 is produced in lower abundance than soluble VP2). For example, the variant promoter can be modified by alteration of the sequence of, or deletion of a portion of, transcriptional elements upstream of the nucleic acid encoding VP1, such as alteration or deletion of all or a portion of the TATA box or the junction site, by introduction of one or more transcription start sites upstream of the nucleic acid encoding VP1, by introduction of one or more transcription stop sites upstream of the nucleic acid encoding VP1, or by any combination thereof. These functional portions of promoters are well-known in the art and can be readily identified and modified in any desired promoter. The recombinant nucleic acids can contain other regulatory elements, such as enhancer binding sites, ribosome binding sites, polyadenylation sites, inducible promoters, repressors, elements that affect RNA stability, siRNA, splice sites, and the like and deletions and alterations thereof, if desired, so that VP1 is produced by the desired host cell in lower abundance than VP2 when equivalent amounts of sequences that encode VP1 and VP2 are present in the desired host cell.

**[0079]** In some embodiments, the recombinant nucleic acid contains a nucleotide sequence that encodes VP2 that is operably linked to the ADH2/GAPDH promoter (FIG. 6, SEQ ID NO: 19), and a nucleotide sequence that encodes VP1 that is operably linked to a variant of the ADH2/GAPDH promoter that results in decreased expression of VP1 relative to VP2 (e.g., soluble VP1 is produced in lower abundance than soluble VP2). Examples of variant ADH2/GAPDH promoters that are suitable for use in the invention include variants in which one or more portions of the promoter (e.g., portions of about 1 to about 20 nt) are replaced with portions of the native parvovirus sequence, such as the "delta As" variant (FIG. 7, SEQ ID NO: 20), variants in which the TATA box is deleted or altered in whole or in part, such as the TATA deletion variant (FIG. 8, SEQ ID NO: 21), variants that contain one or more transcription start and/or stop sequences, such as the 1SS variant (FIG. 9, SEQ ID NO: 22), and promoters that contain any combination of these modifications. In other embodiments, a nucleotide sequence that encodes VP2 is codon optimized for expression in a desired host cell, and a nucleotide sequence that encodes VP1 is not codon optimized for

expression in a desired host cell or is codon deoptimized, for example, through the use of non-preferred codons.

**[0080]** The expression control elements that control expression of VP1 and VP2 can be selected to produce any desired VP1:VP2 expression ratio (%:%) such as about 49:51, about 40:60, about 30:70, about 25:75, about 20:80, about 15:85, about 10:90, about 9:91, about 8:92, about 7:93, about 6:94, about 5:95, about 4:96, about 3:97, about 2:98, about 1:99, about 2:98 to about 20:80, about 5:95 to about 20:80, about 5:95 to about 15:85, or about 7:93 to about 12:88. Preferably, the expression ratio is the ratio of soluble VP1:soluble VP2.

**[0081]** Without wishing to be bound by any particular theory, it is believed that lowered VP1 expression relative to VP2 expression reduces VP1-mediated suppression of VP2 expression, assembly and/or stability and results in more efficient production and higher yield of VLPs that contain VP1 and VP2.

**[0082]** A nucleic acid encoding a parvovirus protein can be constructed using any suitable method (e.g. by chemical synthesis, using recombinant DNA technology) and can take various forms (e.g. single-stranded, double-stranded, vectors, etc.). Many suitable methods for producing recombinant constructs are well-known and conventional in the art. For example, the recombinant nucleic acids can be produced from two or more oligonucleotides comprising sequences encoding portions of the VP1 or VP2 protein or by ligating oligonucleotides to form a coding sequence for the full-length VP1 or VP2 protein using standard molecular biology techniques. See, e.g., U.S. Pat. No. 6,632,601 and U.S. Pat. No. 6,630,298. Preferably, nucleic acids are prepared in substantially pure form (i.e. substantially free from other host cell or non host cell nucleic acids).

**[0083]** Polynucleotides that encode VP1 and/or VP2 proteins of interest can be isolated from a genomic library derived from viral RNA, present in, for example, respiratory secretions or blood from an infected individual. Alternatively, parvovirus nucleic acids can be isolated from infected humans or other mammals or from respiratory secretions or blood collected from infected individuals. An amplification method such as PCR can be used to amplify polynucleotides from either parvovirus genomic RNA or cDNA encoding parvovirus. Alternatively, polynucleotides can be chemically synthesized. The nucleotide sequence can be designed with the appropriate codons for the particular amino acid sequence desired. Preferably, synthetic constructs will contain codons optimized for expression in the intended host cell in which the VP2 or VP1/VP2 protein will be produced. The complete sequence of

the polynucleotide of interest can be assembled from overlapping oligonucleotides prepared by standard methods and assembled into a complete coding sequence. See, e.g., Edge (1981) *Nature* 292:756; Nambair et al. (1984) *Science* 223:1299; Jay et al. (1984) *J. Biol. Chem.* 259:6311; Stemmer et al. (1995) *Gene* 164:49-53. The polynucleotides can be RNA or single- or double-stranded DNA. Preferably, the polynucleotides are isolated free of other components, such as proteins and lipids.

**[0084]** Alternatively, particular nucleotide sequences can be obtained from vectors harboring the desired sequences or synthesized completely or in part using various oligonucleotide synthesis techniques known in the art, such as site-directed mutagenesis and polymerase chain reaction (PCR) techniques where appropriate. See, e.g., Sambrook, *supra*. In particular, one method of obtaining nucleotide sequences encoding the desired amino acid sequences is by annealing complementary sets of overlapping synthetic oligonucleotides, followed by ligation with an appropriate DNA ligase and amplification of the ligated nucleotide sequence via PCR. See, e.g., Jayaraman et al. (1991) *Proc. Natl. Acad. Sci. USA* 88:4084-4088. Additionally, oligonucleotide directed synthesis (Jones et al. (1986) *Nature* 54:75-82), oligonucleotide directed mutagenesis of pre-existing nucleotide regions (Riechmann et al. (1988) *Nature* 332:323-327 and Verhoeyen et al. (1988) *Science* 239:1534-1536), and enzymatic filling-in of gapped oligonucleotides using T4 DNA polymerase (Queen et al. (1989) *Proc. Natl. Acad. Sci. USA* 86:10029-10033) can be used.

**[0085]** Recombinant constructs encoding VP1 and/or VP2 proteins can be prepared in suitable vectors, such as expression vectors, using conventional methods. Preferred recombinant constructs, such as an expression vector, include a nucleic acid sequence which encodes a parvovirus VP1 protein and a nucleic acid sequence which encodes a parvovirus VP2 protein. The recombinant construct can be in the form of DNA, RNA, or a combination of RNA and DNA and can be either single or double stranded. For example, the construct can be in the form of a plasmid, linear DNA or RNA, mRNA, self replicating RNA (e.g., an alpha virus-based replicon), and the like.

**[0086]** A number of suitable vectors for expression of recombinant proteins in a desired host cell are well-known and conventional in the art. Suitable vectors can contain a number of components, including, but not limited to one or more of the following: an origin of replication; a selectable marker gene; one or more expression control elements, such as a transcriptional control element (e.g., a promoter, an enhancer, a terminator), and/or one or more translation signals; and a signal sequence or leader sequence for targeting to the

secretory pathway in a selected host cell (e.g., of mammalian origin or from a heterologous mammalian or non-mammalian species). For example, for expression in insect cells a suitable baculovirus expression vector, such as pFastBac (Invitrogen), can be used to produce recombinant baculovirus particles. The baculovirus particles are amplified and used to infect insect cells to express recombinant protein. For expression in mammalian cells, a vector that will drive expression of the construct in the desired mammalian host cell (e.g., Chinese hamster ovary cells) is used. Similarly, for expression in yeast, a vector that will drive expression in the desired yeast host cell (e.g., *Saccharomyces cerevisiae*, *Candida albicans*, *Candida maltosa*, *Hansenula polymorpha*, *Kluyveromyces fragilis*, *Kluyveromyces lactis*, *Pichia guillermondii*, *Pichia pastoris*, *Schizosaccharomyces pombe* and *Yarrowia lipolytica*) is used.

**[0087]** Viral vectors can be used for the production of VLPs of the invention in eukaryotic cells, such as those derived from the pox family of viruses, including vaccinia virus and avian poxvirus. Additionally, a vaccinia based infection/transfection system, as described in Tomei et al., *J. Virol.* (1993) 67:4017-4026 and Selby et al., *J. Gen. Virol.* (1993) 74:1103-1113, will also find use with the present invention. In this system, cells are first infected *in vitro* with a recombinant vaccinia virus that encodes the bacteriophage T7 RNA polymerase. This polymerase displays exquisite specificity in that it only transcribes templates bearing T7 promoters. Following infection, cells are transfected with the DNA of interest, driven by a T7 promoter. The polymerase expressed in the cytoplasm from the recombinant vaccinia virus transcribes the transfected DNA into RNA which is then translated into protein by the host translational machinery. Alternately, T7 can be added as a purified protein or enzyme as in the "progenitor" system (Studier and Moffatt, *J. Mol. Biol.* (1986) 189:113-130). The method provides for high level, transient, cytoplasmic production of large quantities of RNA and its translation product(s).

**[0088]** A recombinant nucleic acid may comprise a first nucleic acid sequence encoding parvovirus B19 VP1 protein and a second nucleic acid sequence encoding parvovirus B19 VP2 protein, wherein each nucleic acid sequence is controlled by a separate promoter. The parvovirus B19 VP1 and VP2 proteins and their respective control sequences can appear in any desired orientation or order in the recombinant nucleic acid (e.g., plasmid), and are not limited by the language "first nucleic acid" and "second nucleic acid" as used herein.

[0089] The recombinant nucleic acid may be a plasmid that comprises a modification that transcriptionally and/or translationally decrease expression of parvovirus B19 VP1 relative to Parvovirus B19 VP2. A modification may be deletion or alteration of the sequence of all or a portion of the TATA box, deletion or alteration of the sequence of a junction site upstream of the nucleic acid encoding parvovirus B19 VP1, introduction of one or more transcription start and/or stop sites upstream of the nucleic acid encoding parvovirus B19 VP1, or any combination of these modifications. For example, the modification may be deletion of a portion of the TATA box and deletion of a junction site upstream of the nucleic acid encoding parvovirus B19 VP1, or the modification may be introduction of two or more transcription start and/or stop sites upstream of the nucleic acid encoding parvovirus B19 VP1.

[0090] If desired, the recombinant nucleic acid may be a vector that can include a detectable marker. For example, the detectable marker can be a polypeptide that confers resistance to one or more antibiotics.

## **B. Host Cells**

[0091] The invention provides recombinant host cells that contain a recombinant nucleic acid molecule, as described herein, that encode parvovirus VP1 and parvovirus VP2. In some embodiments, the nucleotide sequence that encodes VP1 and the nucleotide sequence that encodes VP2, and their respective control elements, are on separate nucleic acid molecules, for example, on separate vectors. In these embodiments, the host cell contains two different recombinant nucleic acid molecules. Preferably, the host cell contains substantially equal amounts of the nucleotide sequence that encodes VP1 and the nucleotide sequence that encodes VP2 (e.g., substantially equal amounts of two different recombinant nucleic acids). More preferably, the nucleotide sequence that encodes VP1, the nucleotide sequence that encodes VP2, and their respective control elements, are components of a single nucleic acid molecule, such as a bicistronic vector. In these embodiments, the host cell contains a single recombinant nucleic acid molecule, which may be present in multiple copies.

[0092] The recombinant host cells can contain any of the nucleic acid molecules described herein. Suitable host cells include, for example, yeast cells (e.g., *Saccharomyces cerevisiae*, *Candida albicans*, *Candida maltosa*, *Hansenula polymorpha*, *Kluyveromyces fragilis*, *Kluyveromyces lactis*, *Pichia guillermondii*, *Pichia pastoris*, *Schizosaccharomyces*

*pombe* and *Yarrowia lipolytica*), insect cells (e.g., *Aedes aegypti*, *Autographa californica*, *Bombyx mori*, *Drosophila melanogaster*, *Spodoptera frugiperda*, and *Trichoplusia ni*), mammalian cells (e.g., human, non-human primate, horse, cow, sheep, dog, cat, and rodent (e.g., hamster), avian cells (e.g., chicken, duck, and geese, bacteria (e.g., *E. coli*, *Bacillus subtilis*, and *Streptococcus spp.*), Tetrahymena cells (e.g., *Tetrahymena thermophila*) or combinations thereof. Suitable yeast strains include, for example, the AD2, JSC310, AD3 and AD4 strains of *S. cerevisiae*.

**[0093]** Many suitable insect cells and mammalian cells are well-known in the art. Suitable insect cells include, for example, Sf9 cells, Sf21 cells, Tn5 cells, Schneider S2 cells, and High Five cells (a clonal isolate derived from the parental *Trichoplusia ni* BTI-TN-5B1-4 cell line (Invitrogen)). Suitable mammalian cells include, for example, Chinese hamster ovary (CHO) cells, human embryonic kidney cells (HEK293 cells, typically transformed by sheared adenovirus type 5 DNA), NIH-3T3 cells, 293-T cells, Vero cells, HeLa cells, PERC.6 cells (ECACC deposit number 96022940), Hep G2 cells, MRC-5 (ATCC CCL-171), WI-38 (ATCC CCL-75), fetal rhesus lung cells (ATCC CL-160), Madin-Darby bovine kidney (“MDBK”) cells, Madin-Darby canine kidney (“MDCK”) cells (e.g., MDCK (NBL2), ATCC CCL34; or MDCK 33016, DSM ACC 2219), baby hamster kidney (BHK) cells, such as BHK21-F, HKCC cells, and the like. Suitable avian cells include, for example, chicken embryonic stem cells (e.g., EBx® cells), chicken embryonic fibroblasts, chicken embryonic germ cells, duck cells (e.g., AGE1.CR and AGE1.CR.pIX cell lines (ProBioGen) which are described, for example, in *Vaccine* 27:4975-4982 (2009) and WO2005/042728), EB66 cells, and the like.

**[0094]** Suitable insect cell expression systems, such as baculovirus systems, are known to those of skill in the art and described in, e.g., Summers and Smith, *Texas Agricultural Experiment Station Bulletin* No. 1555 (1987). Materials and methods for baculovirus/insect cell expression systems are commercially available in kit form from, *inter alia*, Invitrogen, San Diego CA. Avian cell expression systems are also known to those of skill in the art and described in, e.g., U.S. Patent Nos. 5,340,740; 5,656,479; 5,830,510; 6,114,168; and 6,500,668; European Patent No. EP 0787180B; European Patent Application No. EP03291813.8; WO 03/043415; and WO 03/076601. Similarly, yeast, bacterial and mammalian cell expression systems are also known in the art and described in, e.g., *Yeast Genetic Engineering* (Barr *et al.*, eds., 1989) Butterworths, London.

### C. Production of Virus-like Particles (VLPs)

[0095] The invention provides a process for producing a parvovirus VLP that contains VP1 and VP2. The VLP can contain VP1 and VP2 from any desired parvovirus, or any desired combination. As described herein, the VP1 and VP2 proteins can have an amino acid sequence that is the same as or substantially the same as a naturally occurring parvovirus VP1 or VP2, or can contain one or more amino acid substitutions, deletions or additions. For example, VP1 can be mutated to inactivate its phospholipase activity. For example, the amino acid sequence of VP1 may contain a point mutation (e.g., His153Ala), or any of the mutations described in WO 06/032697, EP 1791858 or US 20070286870. Preferably, the VLP contains VP1 and VP2 are from a parvovirus that infects humans, *i.e.*, a parvovirus of the *Dependovirus*, *Erythrovirus*, or *Bocavirus* genus. Most preferably, the VLP contains parvovirus B19 VP1 and parvovirus B19 VP2. The process includes maintaining a host cell that produces VP1 and VP2, as described herein, under conditions whereby the VP1 and VP2 proteins encoded by the recombinant nucleic acid are produced and assembled into VLPs that contain VP1 and VP2. In the method of the invention, VP1 is produced by the host cell in lower abundance relative to VP2. Preferably, soluble VP1 is produced in lower abundance than soluble VP2. Optionally, the process further includes isolating or purifying the VLPs from the culture media, from a cell lysate or homogenate, or from any combination thereof.

[0096] In some aspects, the method comprises culturing a host cell that contains a recombinant nucleic acid that encodes VP1 and VP2 protein under conditions suitable for expression of VP1 and VP2 and self-assembly of VP1 and VP2 to form VLPs. Conditions suitable for the formation of VLPs are well-known and can be easily determined by a person of ordinary skill in the art. *See, e.g.*, U.S. Pat. No. 7,527,801, which describes the production of viral particles in yeast cells (*Saccharomyces cerevisiae*) and insect cells (SF9), and Taube, S. *et al*, *Archives of Virology*, 150:1425-1431 (2005), which describes the production of VLPs in HEK293T cells. If desired, the method can further include the step of isolating or purifying the parvovirus VLP from the culture media, cells (e.g., from a cell lysate or homogenate) or a combination thereof. In some preferred embodiments, the host cell used to produce the VLPs are yeast cells, mammalian cells, insect cell, or combinations thereof, and the host cells produce less VP1 relative to VP2.

[0097] As described herein, the host cells produce VP1 in lower abundance relative to VP2 (e.g., soluble VP1 is produced in lower abundance than soluble VP2), as a

result of the individual control elements that are operably linked to the nucleic acids that encode VP1 and VP2 and/or as a result of other features of the recombinant nucleic acids that encode VP1 and VP2, such as optimized codon usage and deoptimized codon usage. Such control elements (e.g., promoters) and features (e.g., codon usage) allow for the relative production of VP1 and VP2 to be controlled. Any suitable host cell that permits VP1 to be produced in lower abundance than VP2 can be used in the method. For example, yeast cells (e.g., *Saccharomyces cerevisiae*, *Candida albicans*, *Candida maltosa*, *Hansenula polymorpha*, *Kluyveromyces fragilis*, *Kluyveromyces lactis*, *Pichia guilliermondii*, *Pichia pastoris*, *Schizosaccharomyces pombe* and *Yarrowia lipolytica*), insect cells (e.g., *Aedes aegypti*, *Autographa californica*, *Bombyx mori*, *Drosophila melanogaster*, *Spodoptera frugiperda*, and *Trichoplusia ni*), mammalian cells (e.g., human, non-human primate, horse, cow, sheep, dog, cat, and rodent (e.g., hamster), avian cells (e.g., chicken, duck, and geese, bacteria (e.g., *E. coli*, *Bacillus subtilis*, and *Streptococcus spp.*), Tetrahymena cells (e.g., *Tetrahymena thermophila*) or combinations thereof, can be used. Suitable methods for maintaining such host cells to permit expression of recombinant nucleic acids, including small scale and large scale culture conditions, are known in the art. In some embodiments, the host cell is a yeast cell, mammalian cell or insect cell. In more particular embodiments, the host cell is a yeast cell, such as the *S. cerevisiae* strain AD2, JSC310, AD3, AD4 or combinations thereof.

[0098] In some embodiments, the provided host cell contains two different recombinant nucleic acid molecules, one that encodes VP1 and one that encodes VP2. In other embodiments the provided host cell contains a single recombinant nucleic acid molecule (which may be present in multiple copies) that includes a sequence that encodes VP1, a sequence that encodes VP2, and their respective control elements. For example, in these embodiments, the recombinant nucleic acid can be a bicistronic vector in which a nucleotide sequence that encodes VP1 is operably linked to a first control sequence, and a nucleotide sequence that encodes VP2 is operably linked to a second control sequence. For example, the nucleotide sequence encoding VP2 can be operably linked to a strong promoter, and the nucleotide sequence encoding VP1 can be operably linked to a weak promoter. An exemplary strong promoter for expression in yeast is the ADH2/GAPDH promoter, which is incorporated into pBS24.1 (FIG. 1). An exemplary weak promoter for expression in yeast is the YPTI constitutively active promoter. See, Sears *et al.*, *Yeast* 14:783-790 (1998). Other suitable strong and weak promoters for use in yeasts or other host cells are known in the art.

The promoters used may be any promoters that allow for independent regulation of VP1 and VP2 expression.

**[0099]** In some embodiments, the host cell contains a recombinant nucleic acid in which a nucleotide sequence encoding VP2 is operably linked to a promoter suitable for expression in a desired host cell, and a nucleotide sequence encoding VP1 is operably linked to a variant of that promoter, such as an expression de-optimized variant. The variant promoter can be modified relative to the parental promoter in a variety of ways to reduce the expression of VP1 relative to VP2 (e.g, soluble VP1 is produced in lower abundance than soluble VP2). For example, the variant promoter can be modified by deletion of a portion of transcriptional elements upstream of the nucleic acid encoding VP1, such as deletion or alteration of the sequence of all or a portion of the TATA box, the junction site, by introduction of one or more transcription start sites upstream of the nucleic acid encoding VP1, by introduction of one or more transcription stop sites upstream of the nucleic acid encoding VP1, by any combination thereof. These functional portions of promoters are well-known in the art and can be readily identified and modified in any desired promoter. The recombinant nucleic acids can contain other regulatory elements, such as enhancer binding sites, ribosome binding sites, polyadenylation sites, and the like and deletions and alterations thereof, if desired, so that VP1 is produced by the desired host cell in lower abundance than VP2 when equivalent amounts of sequences that encode VP1 and VP2 are present in the desired host cell.

**[00100]** In some embodiments, the host cell comprises a recombinant nucleic acid that contains a nucleotide sequence that encodes VP2 that is operably linked to the ADH2/GAPDH promoter (FIG. 6, SEQ ID NO: 19), and a nucleotide sequence that encodes VP1 that is operably linked to a variant of the ADH2/GAPDH promoter that results in decreased expression of VP1 relative to VP2. Examples of variant ADH2/GAPDH promoters that are suitable for use in the invention include variants in which one or more portions of the promoter (e.g., portions of about 1 to about 20 bases) are replaced with portions of the native parvovirus sequence, such as the "delta As" variant (FIG. 7, SEQ ID NO: 20), variants in which the TATA box is deleted or altered in whole or in part, such as the TATA deletion variant (FIG. 8, SEQ ID NO: 21), variants that contain one or more transcription start and/or stop sequences, such as the 1SS variant (FIG. 9, SEQ ID NO: 22), and promoters that contain any combination of these modifications. In other embodiments, the host cell comprises a

recombinant nucleic acid that contains a nucleotide sequence encoding VP2 that is codon optimized for expression is a desired host cell, and a nucleotide sequence encoding VP1 that is not codon optimized for expression is a desired host cell or is codon deoptimized, for example, through the use of non-preferred codons.

**[00101]** The host cell provided can contain a recombinant nucleic acid in which the expression control elements that control expression of VP1 and VP2 are selected to produce any desired VP1:VP2 expression ratio. For example, the host cell can produce VP1 and VP2 with a VP1:VP2 expression ratio (%:%) of about 49:51, about 40:60, about 30:70, about 25:75, about 20:80, about 15:85, about 10:90, about 9:91, about 8:92, about 7:93, about 6:94, about 5:95, about 4:96, about 3:97, about 2:98, about 1:99, about 2:98 to about 20:80, about 5:95 to about 20:80, about 5:95 to about 15:85, or about 7:93 to about 12:88. Preferably, the expression ratio is the ratio of soluble VP1:soluble VP2.

**[00102]** In particular embodiments of the method, the host cell comprises an expression vector that encodes VP1 and VP2, preferably parvovirus B19 VP1 and parvovirus B19 VP2. For example, the expression vector can be a plasmid comprising a regulatory element that is operably linked to a nucleotide sequence encoding VP2, and a variant of that regulatory element that is operably linked to a nucleotide sequence encoding VP1. The variant regulatory element transcriptionally, translationally or transcriptionally and translationally decreases expression of VP1 relative to VP2. The variant regulatory element may, for example, have a portion of the TATA box sequence altered or deleted, a portion of the junction site upstream of the VP1 gene deleted, or a combination thereof. The variant regulatory element may alternatively or in addition contain one or more introduced transcription start/stop sites upstream of the VP1 coding sequence. Such variant regulatory elements, and others described herein, allow for the efficient production of VP1/VP2 VLPs. When a variant regulatory element is operably linked to the nucleotide sequence that encodes VP1, the variant regulatory element can contain any one or more of the particular modifications described herein in any desired combination, (e.g., deletion or alteration of all or a portion of the TATA box sequence, deletion or alteration of all or a portion of the A-rich junction region, introduction of one or more transcription start sites and/or transcription stop sites upstream of the VP1 coding sequence, and any combinations thereof). For example, the variant regulatory element is operably linked to the nucleotide sequence that encodes VP1 can include i) deletion or alteration of all or a portion of the TATA box sequence and deletion or

alteration of all or a portion of the A-rich junction region; ii) deletion or alteration of all or a portion of the TATA box sequence and one or more introduced transcription start sites and/or transcription stop sites upstream of the VP1 coding sequence; or iii) deletion or alteration of all or a portion of the A-rich junction region and one or more introduced transcription start sites and/or transcription stop sites upstream of the VP1 coding sequence.

**[00103]** In some aspects, the method further comprises isolating or purifying the VLPs from host cell culture media (e.g., conditioned media), host cells (e.g., cell lysate, cell homogenate) or a combination thereof. Preferably, the VLPs are isolated or purified directly from the host cell culture media, i.e., conditioned culture media. However, if desired, the host cells can be recovered, for example by centrifugation, a host cell homogenate or lysate can be formed using any suitable methods, and VLPs can be isolated from the homogenate or lysate. Suitable chemical, physical or mechanical methods that can be used to lyse cells but keep VLPs substantially intact are known in the art and are described in, e.g., *Protein Purification Applications: A Practical Approach*, (E. L. V. Harris and S. Angal, Eds., 1990).

**[00104]** VLPs can be isolated from culture media or host cells, e.g., a cell lysate or homogenate, using any suitable method. Suitable methods that maintain the integrity of VLPs, such as, density gradient centrifugation, e.g., sucrose gradients, PEG-precipitation, pelleting, and the like (see, e.g., Kirnbauer *et al. J. Virol.* (1993) 67:6929-6936), as well as standard purification techniques including chromatography, e.g., ion exchange and gel filtration chromatography, can be used. Centrifugation, on a sucrose cushion or sucrose gradient is a convenient method for isolating VLPs from VP1 and VP2 monomers or oligomers and from other cellular components. These methods can be used singly, consecutively or integrated into a larger purification scheme. For example, VLPs that are purified on a sucrose cushion or gradient can then be further purified if desired, for example, using ion exchange chromatography, size exclusion chromatography, filtration techniques or any other suitable method.

**[00105]** In one embodiment, the invention provides a method of isolating parvovirus VLPs from host cells, comprising preparing a host cell lysate or homogenate; separating said VLPs from the host cell lysate or homogenate using any suitable method; and further purifying said VLPs. The purification of the VLPs according to this aspect of the invention can, for example, include purification through a sucrose gradient.

[00106] The present invention further provides VLPs made using the methods described herein. In some embodiments, the method comprises culturing a host cell that contains a recombinant nucleic acid that encodes VP1 and VP2 protein under conditions suitable for expression of VP1 and VP2 and self-assembly of VP1 and VP2 to form VLPs. If desired, the method can further comprise isolating or purifying the VLPs from host cell culture media (e.g., conditioned media), host cells (e.g., cell lysate, cell homogenate) or a combination thereof.

#### **D. Immunogenic Composition**

[00107] The invention also provides immunogenic compositions comprising a parvovirus VLP that contains VP1 and VP2 produced by the methods described herein, and immunogenic compositions that contain a recombinant nucleic acid that encodes parvovirus VP1 and VP2 as described herein.

[00108] The immunogenic compositions may comprise a single type of VLP or a mixture of two or more different VLPs and one or more additional antigens, if desired. Antigens may be administered individually or in combination, in e.g., prophylactic (i.e., to prevent infection) or therapeutic (to treat infection) immunogenic compositions. The immunogenic composition may be given more than once (e.g., a “prime” administration followed by one or more “boosts”) to achieve the desired effects. The same composition can be administered in one or more priming and one or more boosting steps. Alternatively, different compositions can be used for priming and boosting. For example, a nucleic acid composition can be administered for priming and a VLP composition can be administered for boosting, or vice versa.

[00109] In the immunogenic compositions that contain a recombinant nucleic acid that encodes parvovirus VP1 and VP2, the recombinant nucleic acid can be any recombinant nucleic acid described herein, for example, linear DNA or RNA, plasmid DNA, mRNA, self replicating RNA and the like. If desired, the nucleic acid can contain one or more modified bases to improve stability and/or resistance to nuclease degradation. If desired, the immunogenic nucleic acid can also include one or more components to facilitate uptake of the nucleic acid by cells and/or reduce nuclease degradation, such as, cationic microparticles or nanoparticles, cationic lipids and the like.

[00110] The immunogenic compositions generally include one or more “pharmaceutically acceptable excipients or vehicles” such as water, saline, glycerol, ethanol, and the like singly or in combination. Immunogenic compositions will typically, in addition to the components mentioned above, comprise one or more “pharmaceutically acceptable carriers.” These include any carrier which does not itself induce the production of antibodies harmful to the individual receiving the composition. Suitable carriers typically are large, slowly metabolized macromolecules such as proteins, polysaccharides, polylactic acids, polyglycolic acids, polymeric amino acids, amino acid copolymers, and lipid aggregates (such as oil droplets or liposomes). Such carriers are well known to those of ordinary skill in the art. Additionally, an auxiliary substance, such as a wetting or emulsifying agent, pH buffering substance, and the like, may be present. A thorough discussion of pharmaceutically acceptable components is available in Gennaro (2000) Remington: *The Science and Practice of Pharmacy*. 20th Ed., ISBN: 0683306472.

[00111] Pharmaceutically acceptable salts can also be used in immunogenic compositions of the invention, for example, mineral salts such as hydrochlorides, hydrobromides, phosphates, or sulfates, as well as salts of organic acids such as acetates, propionates, malonates, or benzoates.

[00112] If desired, antigens can be adsorbed to, entrapped within or otherwise associated with liposomes and particulate carriers such as poly(D,L-lactide co-glycolide) (PLG) microparticles or nanoparticles. Antigens can be conjugated to a carrier protein in order to enhance immunogenicity. See Ramsay et al. (2001) *Lancet* 357(9251):195-196; Lindberg (1999) *Vaccine* 17 Suppl 2:S28-36; Buttery & Moxon (2000) *J R Coll Physicians Lond* 34:163-168; Ahmad & Chapnick (1999) *Infect Dis Clin North Am* 13:113-133, vii; Goldblatt (1998) *J. Med. Microbiol.* 47:563-567; European patent 0 477 508; U.S. Pat. No. 5,306,492; WO98/42721; *Conjugate Vaccines* (eds. Cruse *et al.*) ISBN 3805549326, particularly vol. 10:48-114; Hermanson (1996) *Bioconjugate Techniques* ISBN: 0123423368 or 012342335X.

[00113] Immunogenic compositions of the present invention may be administered in conjunction with other immunoregulatory agents. For example, an immunogenic composition of the invention can include an adjuvant. Preferred adjuvants include, but are not limited to, one or more of the following types of adjuvants described below.

Immunogenic compositions of the present invention may also be pre-mixed with an adjuvant before administration.

**[00114]** In one embodiment, an immunogenic composition comprises a parvovirus B19 VLP (e.g., 0.05, 0.5, 5 µg protein/ml) and a MF59C.1 adjuvant, wherein 50% of the final volume is MF59C.1 adjuvant. In a specific embodiment, an immunogenic composition comprises 0.8 mg/ml of parvovirus B19 VLP and MF59C.1 as an adjuvant, at a final volume of 1:1.

**[00115] Alum**

**[00116]** In one embodiment, the adjuvant for use in the present invention is alum (aluminum potassium sulfate ( $\text{AlK}(\text{SO}_4)_2$ )), or an alum derivative, such as that formed in-situ by mixing an antigen in phosphate buffer with alum, followed by titration and precipitation with a base such as ammonium hydroxide or sodium hydroxide.

**[00117] Retinoic acid**

**[00118]** In one embodiment, the adjuvant for use in the present invention is retinoic acid, the oxidized form of Vitamin A, with only partial vitamin A function.

**[00119] MF59C.1**

**[00120]** In one embodiment, the adjuvant for use in the present invention is MF59C.1, an oil-in-water emulsion (squalene) in citrate buffer. MF59C.1 has been shown to be an effective adjuvant and enhance the production of high titers of neutralizing antibodies against parvovirus B19 (Ballou *et al.*, JID, 187:675-678 (2003)).

**[00121] Mineral-containing compositions**

**[00122]** Mineral containing compositions suitable for use as adjuvants in the invention include mineral salts, such as aluminum salts and calcium salts. The invention includes mineral salts such as hydroxides (e.g. oxyhydroxides), phosphates (e.g. hydroxyphosphates, orthophosphates), sulphates, *etc.* [e.g. see chapters 8 & 9 of *Vaccine Design...* (1995) eds. Powell & Newman. ISBN: 030644867X. Plenum.], or mixtures of different mineral compounds, with the compounds taking any suitable form (e.g. gel,

crystalline, amorphous, *etc.*), and with adsorption being preferred. The mineral containing compositions may also be formulated as a particle of metal salt.

**[00123]** The adjuvants known as “aluminium hydroxide” are typically aluminium oxyhydroxide salts, which are usually at least partially crystalline. Aluminium oxyhydroxide, which can be represented by the formula  $\text{AlO}(\text{OH})$ , can be distinguished from other aluminium compounds, such as aluminium hydroxide  $\text{Al}(\text{OH})_3$ , by infrared (IR) spectroscopy, in particular by the presence of an adsorption band at  $1070\text{cm}^{-1}$  and a strong shoulder at  $3090\text{--}3100\text{cm}^{-1}$  [chapter 9 of *Vaccine Design...* (1995) eds. Powell & Newman. ISBN: 030644867X. Plenum.] The degree of crystallinity of an aluminum hydroxide adjuvant is reflected by the width of the diffraction band at half height (WHH), with poorly-crystalline particles showing greater line broadening due to smaller crystallite sizes. The surface area increases as WHH increases, and adjuvants with higher WHH values have been seen to have greater capacity for antigen adsorption. A fibrous morphology (*e.g.* as seen in transmission electron micrographs) is typical for aluminum hydroxide adjuvants. The pI of aluminium hydroxide adjuvants is typically about 11 *i.e.* the adjuvant itself has a positive surface charge at physiological pH. Adsorptive capacities of between 1.8-2.6 mg protein per mg  $\text{Al}^{+++}$  at pH 7.4 have been reported for aluminium hydroxide adjuvants.

**[00124]** The adjuvants known as “aluminium phosphate” are typically aluminium hydroxyphosphates, often also containing a small amount of sulfate (*i.e.* aluminium hydroxyphosphate sulfate). They may be obtained by precipitation, and the reaction conditions and concentrations during precipitation influence the degree of substitution of phosphate for hydroxyl in the salt. Hydroxyphosphates generally have a  $\text{PO}_4/\text{Al}$  molar ratio between 0.3 and 1.2. Hydroxyphosphates can be distinguished from strict  $\text{AlPO}_4$  by the presence of hydroxyl groups. For example, an IR spectrum band at  $3164\text{cm}^{-1}$  (*e.g.* when heated to  $200^\circ\text{C}$ ) indicates the presence of structural hydroxyls [ch. 9 of *Vaccine Design...* (1995) eds. Powell & Newman. ISBN: 030644867X. Plenum.].

**[00125]** The  $\text{PO}_4/\text{Al}^{3+}$  molar ratio of an aluminium phosphate adjuvant will generally be between 0.3 and 1.2, preferably between 0.8 and 1.2, and more preferably  $0.95 \pm 0.1$ . The aluminium phosphate will generally be amorphous, particularly for hydroxyphosphate salts. A typical adjuvant is amorphous aluminium hydroxyphosphate with  $\text{PO}_4/\text{Al}$  molar ratio between 0.84 and 0.92, included at  $0.6\text{mg Al}^{3+}/\text{ml}$ . The aluminium phosphate will generally be particulate (*e.g.* plate-like morphology as seen in transmission

electron micrographs). Typical diameters of the particles are in the range 0.5-20 $\mu$ m (*e.g.* about 5-10 $\mu$ m) after any antigen adsorption. Adsorptive capacities of between 0.7-1.5 mg protein per mg Al<sup>+++</sup> at pH 7.4 have been reported for aluminium phosphate adjuvants.

**[00126]** The point of zero charge (PZC) of aluminium phosphate is inversely related to the degree of substitution of phosphate for hydroxyl, and this degree of substitution can vary depending on reaction conditions and concentration of reactants used for preparing the salt by precipitation. PZC is also altered by changing the concentration of free phosphate ions in solution (more phosphate = more acidic PZC) or by adding a buffer such as a histidine buffer (makes PZC more basic). Aluminium phosphates used according to the invention will generally have a PZC of between 4.0 and 7.0, more preferably between 5.0 and 6.5 *e.g.* about 5.7.

**[00127]** Suspensions of aluminium salts used to prepare compositions of the invention may contain a buffer (*e.g.* a phosphate or a histidine or a Tris buffer), but this is not always necessary. The suspensions are preferably sterile and pyrogen-free. A suspension may include free aqueous phosphate ions *e.g.* present at a concentration between 1.0 and 20 mM, preferably between 5 and 15 mM, and more preferably about 10 mM. The suspensions may also comprise sodium chloride.

**[00128]** In one embodiment, an adjuvant component includes a mixture of both an aluminium hydroxide and an aluminium phosphate. In this case there may be more aluminium phosphate than hydroxide *e.g.* a weight ratio of at least 2:1 *e.g.*  $\geq 5:1$ ,  $\geq 6:1$ ,  $\geq 7:1$ ,  $\geq 8:1$ ,  $\geq 9:1$ , *etc.*

**[00129]** The concentration of Al<sup>+++</sup> in a composition for administration to a patient is preferably less than 10mg/ml *e.g.*  $\leq 5$  mg/ml,  $\leq 4$  mg/ml,  $\leq 3$  mg/ml,  $\leq 2$  mg/ml,  $\leq 1$  mg/ml, *etc.* A preferred range is between 0.3 and 1mg/ml. A maximum of <0.85mg/dose is preferred.

#### **[00130] Oil Emulsions**

**[00131]** Oil emulsion compositions suitable for use as adjuvants in the invention include squalene-water emulsions, such as MF59 [Chapter 10 of *Vaccine Design...* (1995) eds. Powell & Newman. ISBN: 030644867X. Plenum.] (5% Squalene, 0.5% Tween 80, and 0.5% Span 85, formulated into submicron particles using a microfluidizer). Complete Freund's adjuvant (CFA) and incomplete Freund's adjuvant (IFA) may also be used.

**[00132]** Various suitable oil-in-water emulsions are known, and they typically include at least one oil and at least one surfactant, with the oil(s) and surfactant(s) being biodegradable (metabolisable) and biocompatible. The oil droplets in the emulsion are generally less than 5µm in diameter, and advantageously the emulsion comprises oil droplets with a sub-micron diameter, with these small sizes being achieved with a microfluidiser to provide stable emulsions. Droplets with a size less than 220nm are preferred as they can be subjected to filter sterilization.

**[00133]** The invention can be used with oils such as those from an animal (such as fish) or vegetable source. Sources for vegetable oils include nuts, seeds and grains. Peanut oil, soybean oil, coconut oil, and olive oil, the most commonly available, exemplify the nut oils. Jojoba oil can be used *e.g.* obtained from the jojoba bean. Seed oils include safflower oil, cottonseed oil, sunflower seed oil, sesame seed oil and the like. In the grain group, corn oil is the most readily available, but the oil of other cereal grains such as wheat, oats, rye, rice, teff, triticale and the like may also be used. 6-10 carbon fatty acid esters of glycerol and 1,2-propanediol, while not occurring naturally in seed oils, may be prepared by hydrolysis, separation and esterification of the appropriate materials starting from the nut and seed oils. Fats and oils from mammalian milk are metabolizable and may therefore be used in the practice of this invention. The procedures for separation, purification, saponification and other means necessary for obtaining pure oils from animal sources are well known in the art. Most fish contain metabolizable oils which may be readily recovered. For example, cod liver oil, shark liver oils, and whale oil such as spermaceti exemplify several of the fish oils which may be used herein. A number of branched chain oils are synthesized biochemically in 5-carbon isoprene units and are generally referred to as terpenoids. Shark liver oil contains a branched, unsaturated terpenoid known as squalene, 2,6,10,15,19,23-hexamethyl-2,6,10,14,18,22-tetracosahexaene. Other preferred oils are the tocopherols (see below). Oil in water emulsions comprising squalene are particularly preferred. Mixtures of oils can be used.

**[00134]** Surfactants can be classified by their 'HLB' (hydrophile/lipophile balance). Preferred surfactants of the invention have a HLB of at least 10, preferably at least 15, and more preferably at least 16. The invention can be used with surfactants including, but not limited to: the polyoxyethylene sorbitan esters surfactants (commonly referred to as the Tweens), especially polysorbate 20 and polysorbate 80; copolymers of ethylene oxide (EO),

propylene oxide (PO), and/or butylene oxide (BO), sold under the DOWFAX™ tradename, such as linear EO/PO block copolymers; octoxynols, which can vary in the number of repeating ethoxy (oxy-1,2-ethanediyl) groups, with octoxynol-9 (Triton X-100, or t-octylphenoxypolyethoxyethanol) being of particular interest; (octylphenoxy)polyethoxyethanol (IGEPAL CA-630/NP-40); phospholipids such as phosphatidylcholine (lecithin); polyoxyethylene fatty ethers derived from lauryl, cetyl, stearyl and oleyl alcohols (known as Brij surfactants), such as triethyleneglycol monolauryl ether (Brij 30); and sorbitan esters (commonly known as the SPANs), such as sorbitan trioleate (Span 85) and sorbitan monolaurate. Preferred surfactants for including in the emulsion are Tween 80 (polyoxyethylene sorbitan monooleate), Span 85 (sorbitan trioleate), lecithin and Triton X-100. As mentioned above, detergents such as Tween 80 may contribute to the thermal stability seen in the examples below.

**[00135]** Mixtures of surfactants can be used *e.g.* Tween 80/Span 85 mixtures. A combination of a polyoxyethylene sorbitan ester such as polyoxyethylene sorbitan monooleate (Tween 80) and an octoxynol such as t-octylphenoxypolyethoxyethanol (Triton X-100) is also suitable. Another useful combination comprises laureth 9 plus a polyoxyethylene sorbitan ester and/or an octoxynol.

**[00136]** Preferred amounts of surfactants (% by weight) are: polyoxyethylene sorbitan esters (such as Tween 80) 0.01 to 1%, in particular about 0.1 %; octyl- or nonylphenoxy polyoxyethanols (such as Triton X-100, or other detergents in the Triton series) 0.001 to 0.1 %, in particular 0.005 to 0.02%; polyoxyethylene ethers (such as laureth 9) 0.1 to 20 %, preferably 0.1 to 10 % and in particular 0.1 to 1 % or about 0.5%.

**[00137]** Specific oil-in-water emulsion adjuvants useful with the invention include, but are not limited to:

- A submicron emulsion of squalene, Tween 80, and Span 85. The composition of the emulsion by volume can be about 5% squalene, about 0.5% polysorbate 80 and about 0.5% Span 85. In weight terms, these ratios become 4.3% squalene, 0.5% polysorbate 80 and 0.48% Span 85. This adjuvant is known as 'MF59'. The MF59 emulsion advantageously includes citrate ions *e.g.* 10mM sodium citrate buffer.
- An emulsion comprising squalene, an  $\alpha$ -tocopherol, and polysorbate 80. These emulsions may have from 2 to 10% squalene, from 2 to 10% tocopherol and from 0.3 to

3% Tween 80, and the weight ratio of squalene:tocopherol is preferably  $\leq 1$  (*e.g.* 0.90) as this provides a more stable emulsion. Squalene and Tween 80 may be present volume ratio of about 5:2, or at a weight ratio of about 11:5. One such emulsion can be made by dissolving Tween 80 in PBS to give a 2% solution, then mixing 90ml of this solution with a mixture of (5g of DL- $\alpha$ -tocopherol and 5ml squalene), then microfluidising the mixture. The resulting emulsion may have submicron oil droplets *e.g.* with an average diameter of between 100 and 250nm, preferably about 180nm.

- An emulsion of squalene, a tocopherol, and a Triton detergent (*e.g.* Triton X-100). The emulsion may also include a 3d-MPL (see below). The emulsion may contain a phosphate buffer.
- An emulsion comprising a polysorbate (*e.g.* polysorbate 80), a Triton detergent (*e.g.* Triton X-100) and a tocopherol (*e.g.* an  $\alpha$ -tocopherol succinate). The emulsion may include these three components at a mass ratio of about 75:11:10 (*e.g.* 750 $\mu$ g/ml polysorbate 80, 110 $\mu$ g/ml Triton X-100 and 100 $\mu$ g/ml  $\alpha$ -tocopherol succinate), and these concentrations should include any contribution of these components from antigens. The emulsion may also include squalene. The emulsion may also include a 3d-MPL (see below). The aqueous phase may contain a phosphate buffer.
- An emulsion of squalane, polysorbate 80 and poloxamer 401 (“Pluronic™ L121”). The emulsion can be formulated in phosphate buffered saline, pH 7.4. This emulsion is a useful delivery vehicle for muramyl dipeptides, and has been used with threonyl-MDP in the “SAF-1” adjuvant (0.05-1% Thr-MDP, 5% squalane, 2.5% Pluronic L121 and 0.2% polysorbate 80). It can also be used without the Thr-MDP, as in the “AF” adjuvant (5% squalane, 1.25% Pluronic L121 and 0.2% polysorbate 80). Microfluidisation is preferred.
- An emulsion comprising squalene, an aqueous solvent, a polyoxyethylene alkyl ether hydrophilic nonionic surfactant (*e.g.* polyoxyethylene (12) cetostearyl ether) and a hydrophobic nonionic surfactant (*e.g.* a sorbitan ester or mannide ester, such as sorbitan monoleate or ‘Span 80’). The emulsion is preferably thermoreversible and/or has at least 90% of the oil droplets (by volume) with a size less than 200 nm. The emulsion may also include one or more of: alditol; a cryoprotective agent (*e.g.* a sugar, such as dodecylmaltoside and/or sucrose); and/or an alkylpolyglycoside. Such emulsions may be lyophilized.

- An emulsion having from 0.5-50% of an oil, 0.1-10% of a phospholipid, and 0.05-5% of a non-ionic surfactant. Preferred phospholipid components are phosphatidylcholine, phosphatidylethanolamine, phosphatidylserine, phosphatidylinositol, phosphatidylglycerol, phosphatidic acid, sphingomyelin and cardiolipin. Submicron droplet sizes are advantageous.
- A submicron oil-in-water emulsion of a non-metabolisable oil (such as light mineral oil) and at least one surfactant (such as lecithin, Tween 80 or Span 80). Additives may be included, such as QuilA saponin, cholesterol, a saponin-lipophile conjugate (such as GPI-0100, produced by addition of aliphatic amine to desacylsaponin via the carboxyl group of glucuronic acid), dimethyldioctadecylammonium bromide and/or N,N-dioctadecyl-N,N-bis (2-hydroxyethyl)propanediamine.
- An emulsion comprising a mineral oil, a non-ionic lipophilic ethoxylated fatty alcohol, and a non-ionic hydrophilic surfactant (*e.g.* an ethoxylated fatty alcohol and/or polyoxyethylene-polyoxypropylene block copolymer) (see WO2006/113373).
- An emulsion comprising a mineral oil, a non-ionic hydrophilic ethoxylated fatty alcohol, and a non-ionic lipophilic surfactant (*e.g.* an ethoxylated fatty alcohol and/or polyoxyethylene-polyoxypropylene block copolymer) (see WO2006/113373).
- An emulsion in which a saponin (*e.g.* QuilA or QS21) and a sterol (*e.g.* a cholesterol) are associated as helical micelles.

[00132] Antigens (VLPs) and adjuvants in a composition will typically be in admixture at the time of delivery to a patient. The emulsions may be mixed with antigen (VLP) during manufacture, or extemporaneously, at the time of delivery. Thus the adjuvant and antigen (VLP) may be kept separately in a packaged or distributed vaccine, ready for final formulation at the time of use. The antigen (VLP) will generally be in an aqueous form, such that the vaccine is finally prepared by mixing two liquids. The volume ratio of the two liquids for mixing can vary (*e.g.* between 5:1 and 1:5) but is generally about 1:1.

[00133] **Saponin formulations** (see chapter 22 of *Vaccine Design...* (1995) eds. Powell & Newman. ISBN: 030644867X. Plenum.)

[00134] Saponin formulations may also be used as adjuvants in the invention. Saponins are a heterogeneous group of sterol glycosides and triterpenoid glycosides that are found in the bark, leaves, stems, roots and even flowers of a wide range of plant species.

Saponin from the bark of the *Quillaia saponaria* Molina tree have been widely studied as adjuvants. Saponin can also be commercially obtained from *Smilax ornata* (sarsapilla), *Gypsophilla paniculata* (brides veil), and *Saponaria officianalis* (soap root). Saponin adjuvant formulations include purified formulations, such as QS21, as well as lipid formulations, such as ISCOMs. QS21 is marketed as Stimulon™.

**[00135]** Saponin compositions have been purified using HPLC and RP-HPLC. Specific purified fractions using these techniques have been identified, including QS7, QS17, QS18, QS21, QH-A, QH-B and QH-C. Preferably, the saponin is QS21. A method of production of QS21 is disclosed in US 5,057,540. Saponin formulations may also comprise a sterol, such as cholesterol.

**[00136]** Combinations of saponins and cholesterol can be used to form unique particles called immunostimulating complexes (ISCOMs) (chapter 23 of *Vaccine Design...* (1995) eds. Powell & Newman. ISBN: 030644867X. Plenum.). ISCOMs typically also include a phospholipid such as phosphatidylethanolamine or phosphatidylcholine. Any known saponin can be used in ISCOMs. Preferably, the ISCOM includes one or more of QuilA, QHA & QHC. ISCOMs are further described in WO96/33739. Optionally, the ISCOMS may be devoid of additional detergent.

**[00137] Virosomes and virus-like particles**

**[00138]** Virosomes and virus-like particles (VLPs) can also be used as adjuvants in the invention. These structures generally contain one or more proteins from a virus optionally combined or formulated with a phospholipid. They are generally non-pathogenic, non-replicating and generally do not contain any of the native viral genome. The viral proteins may be recombinantly produced or isolated from whole viruses. These viral proteins suitable for use in virosomes or VLPs include proteins derived from influenza virus (such as HA or NA), Hepatitis B virus (such as core or capsid proteins), Hepatitis E virus, measles virus, Sindbis virus, Rotavirus, Foot-and-Mouth Disease virus, Retrovirus, Norwalk virus, human Papilloma virus, HIV, RNA-phages, Q $\beta$ -phage (such as coat proteins), GA-phage, fr-phage, AP205 phage, and Ty (such as retrotransposon Ty protein p1).

**[00139] Bacterial or microbial derivatives**

**[00140]** Adjuvants suitable for use in the invention include bacterial or microbial derivatives such as non-toxic derivatives of enterobacterial lipopolysaccharide (LPS), Lipid

A derivatives, immunostimulatory oligonucleotides and ADP-ribosylating toxins and detoxified derivatives thereof.

**[00141]** Non-toxic derivatives of LPS include monophosphoryl lipid A (MPL) and 3-O-deacylated MPL (3dMPL). 3dMPL is a mixture of 3 de-O-acylated monophosphoryl lipid A with 4, 5 or 6 acylated chains. A preferred “small particle” form of 3 De-O-acylated monophosphoryl lipid A is disclosed in EP-A-0689454. Such “small particles” of 3dMPL are small enough to be sterile filtered through a 0.22µm membrane (EP-A-0689454). Other non-toxic LPS derivatives include monophosphoryl lipid A mimics, such as aminoalkyl glucosaminide phosphate derivatives *e.g.* RC-529.

**[00142]** Lipid A derivatives include derivatives of lipid A from *Escherichia coli* such as OM-174. OM-174 is described for example in Meraldi *et al.* (2003) *Vaccine* 21:2485-2491 and Pajak *et al.* (2003) *Vaccine* 21:836-842.

**[00143]** Immunostimulatory oligonucleotides suitable for use as adjuvants in the invention include nucleotide sequences containing a CpG motif (a dinucleotide sequence containing an unmethylated cytosine linked by a phosphate bond to a guanosine). Double-stranded RNAs and oligonucleotides containing palindromic or poly(dG) sequences have also been shown to be immunostimulatory.

**[00144]** The CpG's can include nucleotide modifications/analogues such as phosphorothioate modifications and can be double-stranded or single-stranded. References Kandimalla *et al.* (2003) *Nucleic Acids Research* 31:2393-2400; WO02/26757, and WO99/62923 disclose possible analog substitutions *e.g.* replacement of guanosine with 2'-deoxy-7-deazaguanosine. The adjuvant effect of CpG oligonucleotides is further discussed in Krieg (2003) *Nature Medicine* 9:831-835; McCluskie *et al.* (2002) *FEMS Immunology and Medical Microbiology* 32:179-185; WO98/40100; US 6,207,646; US 6,239,116; and US 6,429,199.

**[00145]** The CpG sequence may be directed to TLR9, such as the motif GTCGTT or TTCGTT (Kandimalla *et al.* (2003) *Biochemical Society Transactions* 31 (part 3):654-658). The CpG sequence may be specific for inducing a Th1 immune response, such as a CpG-A ODN, or it may be more specific for inducing a B cell response, such as a CpG-B ODN. CpG-A and CpG-B ODNs are discussed in Blackwell *et al.* (2003) *J Immunol* 170:4061-4068; Krieg (2002) *Trends Immunol* 23:64-65 and WO01/95935. Preferably, the CpG is a CpG-A ODN.

[00146] Preferably, the CpG oligonucleotide is constructed so that the 5' end is accessible for receptor recognition. Optionally, two CpG oligonucleotide sequences may be attached at their 3' ends to form "immunomers". See, for example, Kandimalla *et al.* (2003) *Biochemical Society Transactions* 31 (part 3):654-658 & Kandimalla *et al.* (2003) BBRC 306:948-953; Bhagat *et al.* (2003) BBRC 300:853-861 and WO03/035836

[00147] A particularly useful adjuvant based around immunostimulatory oligonucleotides is known as IC-31<sup>TM</sup> (Schellack *et al.* (2006) *Vaccine* 24:5461-72 ). Thus an adjuvant used with the invention may comprise a mixture of (i) an oligonucleotide (*e.g.* between 15-40 nucleotides) including at least one (and preferably multiple) CpI motifs (*i.e.* a cytosine linked to an inosine to form a dinucleotide), and (ii) a polycationic polymer, such as an oligopeptide (*e.g.* between 5-20 amino acids) including at least one (and preferably multiple) Lys-Arg-Lys tripeptide sequence(s). The oligonucleotide may be a deoxynucleotide comprising 26-mer sequence 5'-(IC)<sub>13</sub>-3' (SEQ ID NO: 1). The polycationic polymer may be a peptide comprising 11-mer amino acid sequence KKLKLLKLLK (SEQ ID NO: 2).

[00148] Bacterial ADP-ribosylating toxins and detoxified derivatives thereof may be used as adjuvants in the invention. Preferably, the protein is derived from *E.coli* (*E.coli* heat labile enterotoxin "LT"), cholera ("CT"), or pertussis ("PT"). The use of detoxified ADP-ribosylating toxins as mucosal adjuvants is described in WO95/17211 and as parenteral adjuvants in WO98/42375. The toxin or toxoid is preferably in the form of a holotoxin, comprising both A and B subunits. Preferably, the A subunit contains a detoxifying mutation; preferably the B subunit is not mutated. Preferably, the adjuvant is a detoxified LT mutant such as LT-K63, LT-R72, and LT-G192. The use of ADP-ribosylating toxins and detoxified derivatives thereof, particularly LT-K63 and LT-R72, as adjuvants can be found in Beignon *et al.* (2002) *Infect Immun* 70:3012-3019; Pizza *et al.* (2001) *Vaccine* 19:2534-2541; Pizza *et al.* (2000) *Int J Med Microbiol* 290:455-461; Schariton-Kersten *et al.* (2000) *Infect Immun* 68:5306-5313; Ryan *et al.* (1999) *Infect Immun* 67:6270-6280; Partidos *et al.* (1999) *Immunol Lett* 67:209-216; Peppoloni *et al.* (2003) *Expert Rev Vaccines* 2:285-293; Pine *et al.* (2002) *J Control Release* 85:263-270 and Tebbey *et al.* (2000) *Vaccine* 18:2723-34. A useful CT mutant is or CT-E29H. Numerical reference for amino acid substitutions is preferably based on the alignments of the A and B subunits of ADP-ribosylating toxins set forth in Domenighini *et al.* (1995) *Mol Microbiol* 15:1165-1167, specifically incorporated herein by reference in its entirety.

**[00149] Human immunomodulators**

**[00150]** Human immunomodulators suitable for use as adjuvants in the invention include cytokines, such as interleukins (*e.g.* IL-1, IL-2, IL-4, IL-5, IL-6, IL-7, IL-12, *etc.*) (WO99/40936 and WO99/44636), interferons (*e.g.* interferon- $\gamma$ ), macrophage colony stimulating factor, and tumor necrosis factor. A preferred immunomodulator is IL-12.

**[00151] Bioadhesives and Mucoadhesives**

**[00152]** Bioadhesives and mucoadhesives may also be used as adjuvants in the invention. Suitable bioadhesives include esterified hyaluronic acid microspheres (Singh *et al*] (2001) *J Cont Release* 70:267-276) or mucoadhesives such as cross-linked derivatives of poly(acrylic acid), polyvinyl alcohol, polyvinyl pyrrolidone, polysaccharides and carboxymethylcellulose. Chitosan and derivatives thereof may also be used as adjuvants in the invention (WO99/27960).

**[00153] Microparticles**

**[00154]** Microparticles may also be used as adjuvants in the invention. Microparticles (*i.e.* a particle of ~100nm to ~150 $\mu$ m in diameter, more preferably ~200nm to ~30 $\mu$ m in diameter, and most preferably ~500nm to ~10 $\mu$ m in diameter) formed from materials that are biodegradable and non-toxic (*e.g.* a poly( $\alpha$ -hydroxy acid), a polyhydroxybutyric acid, a polyorthoester, a polyanhydride, a polycaprolactone, *etc.*), with poly(lactide-co-glycolide) are preferred, optionally treated to have a negatively-charged surface (*e.g.* with SDS) or a positively-charged surface (*e.g.* with a cationic detergent, such as CTAB).

**[00155] Liposomes** (Chapters 13 & 14 of *Vaccine Design...* (1995) eds. Powell & Newman. ISBN: 030644867X. Plenum.)

**[00156]** Examples of liposome formulations suitable for use as adjuvants are described in US 6,090,406, US 5,916,588 and EP A 0626169.

**[00157] Polyoxyethylene ether and polyoxyethylene ester formulations**

**[00158]** Adjuvants suitable for use in the invention include polyoxyethylene ethers and polyoxyethylene esters (WO99/52549). Such formulations further include polyoxyethylene sorbitan ester surfactants in combination with an octoxynol (WO01/21207) as well as polyoxyethylene alkyl ethers or ester surfactants in combination with at least one

additional non-ionic surfactant such as an octoxynol (WO01/21152). Preferred polyoxyethylene ethers are selected from the following group: polyoxyethylene-9-lauryl ether (laureth 9), polyoxyethylene-9-stearyl ether, polyoxyethylene-8-stearyl ether, polyoxyethylene-4-lauryl ether, polyoxyethylene-35-lauryl ether, and polyoxyethylene-23-lauryl ether.

**[00159] Polyphosphazene (PCPP)**

**[00160]** PCPP formulations are described, for example, in Andrianov *et al.* (1998) *Biomaterials* 19:109-115 and Payne *et al.* (1998) *Adv Drug Delivery Review* 31:185-196.

**[00161] Muramyl peptides**

**[00162]** Examples of muramyl peptides suitable for use as adjuvants in the invention include N-acetyl-muramyl-L-threonyl-D-isoglutamine (thr-MDP), N-acetyl-normuramyl-L-alanyl-D-isoglutamine (nor-MDP), and N-acetylmuramyl-L-alanyl-D-isoglutaminyl-L-alanine-2-(1'-2'-dipalmitoyl-*sn*-glycero-3-hydroxyphosphoryloxy)-ethylamine MTP-PE).

**[00163] Imidazoquinolone Compounds**

**[00164]** Examples of imidazoquinolone compounds suitable for use adjuvants in the invention include Imiquimod and its homologues (*e.g.* "Resiquimod 3M"), described further in Stanley (2002) *Clin Exp Dermatol* 27:571-577 and Jones (2003) *Curr Opin Investig Drugs* 4:214-218.

**[00165]** The invention may also comprise combinations of aspects of one or more of the adjuvants identified above. For example, the following adjuvant compositions may be used in the invention: (1) a saponin and an oil-in-water emulsion (WO99/11241); (2) a saponin (*e.g.* QS21) + a non-toxic LPS derivative (*e.g.* 3dMPL) (WO94/00153); (3) a saponin (*e.g.* QS21) + a non-toxic LPS derivative (*e.g.* 3dMPL) + a cholesterol; (4) a saponin (*e.g.* QS21) + 3dMPL + IL-12 (optionally + a sterol) (WO98/57659); (5) combinations of 3dMPL with, for example, QS21 and/or oil-in-water emulsions (European patent applications 0835318, 0735898 and 0761231); (6) SAF, containing 10% squalane, 0.4% Tween 80™, 5% pluronic-block polymer L121, and thr-MDP, either microfluidized into a submicron emulsion or vortexed to generate a larger particle size emulsion; (7) Ribi™ adjuvant system (RAS), (Ribi Immunochem) containing 2% squalene, 0.2% Tween 80, and one or more bacterial cell wall components from the group consisting of monophosphorylipid A (MPL), trehalose

dimycolate (TDM), and cell wall skeleton (CWS), preferably MPL + CWS (Detox™); and (8) one or more mineral salts (such as an aluminum salt) + a non-toxic derivative of LPS (such as 3dMPL).

[00166] Other substances that act as immunostimulating agents are disclosed in chapter 7 of *Vaccine Design...* (1995) eds. Powell & Newman. ISBN: 030644867X. Plenum.

The use of an aluminium hydroxide and/or aluminium phosphate adjuvant is useful, particularly in children, and antigens are generally adsorbed to these salts. Squalene-in-water emulsions are also preferred, particularly in the elderly. Useful adjuvant combinations include combinations of Th1 and Th2 adjuvants such as CpG and alum or resiquimod and alum. A combination of aluminium phosphate and 3dMPL may be used.

[00167] In some embodiments, the invention is an immunogenic composition that contains a parvovirus VLP that contains VP1 and VP2, as described herein, and an adjuvant, such as MF59. The VLP and the adjuvant (e.g., MF59) can be premixed and provided as a single composition, or can be provided as separate components that are to be mixed prior to administration.

## **E. Administration**

[00168] Compositions of the invention will generally be administered directly to a patient. Direct delivery may be accomplished by parenteral injection (e.g. subcutaneously, intraperitoneally, intravenously, intramuscularly, or to the interstitial space of a tissue) for example using a syringe and hypodermic needle, or mucosally, such as by rectal, oral (e.g. tablet, spray), vaginal, topical, transdermal (See e.g. WO99/27961) or transcutaneous (See e.g. WO02/074244 and WO02/064162), intranasal (See e.g. WO03/028760), ocular, aural, pulmonary or other mucosal administration. Immunogenic compositions can also be administered topically by direct transfer to the surface of the skin. Topical administration can be accomplished without utilizing any devices, or by contacting naked skin with the immunogenic composition utilizing a bandage or a bandage-like device (see, e.g., U.S. Pat. No. 6,348,450).

[00169] Preferably the mode of administration is parenteral, mucosal or a combination of mucosal and parenteral immunizations. Even more preferably, the mode of administration is parenteral, mucosal or a combination of mucosal and parenteral immunizations in a total of 1-2 vaccinations 1-3 weeks apart. Preferably the route of administration includes but is not limited to oral delivery, intra-muscular delivery and a

combination of oral and intramuscular delivery. A particularly preferred mode of administration is by intramuscular injection, for example using a syringe and hypodermic needle. Preferably the immunogenic composition is administered in two doses intramuscularly.

**[00170]** Compositions of the invention may be administered concomitantly with routine childhood immunizations or with human papillomavirus and/or group B strep vaccinations delivered to adolescent females.

**[00171]** It has already been demonstrated that mucosal and systemic immune responses to antigens from mucosal pathogens, such as *Helicobacter pylori* antigens can be enhanced through mucosal priming followed by systemic boosting immunizations (see Vajdy et al. (2003) *Immunology* 110: 86-94). In some embodiments, the method for treating or preventing an infection by a parvovirus, comprises mucosally administering to a subject in need thereof a first immunogenic composition comprising one or more parvovirus antigens followed by parenterally administering a therapeutically effective amount of a second immunogenic composition comprising one or more parvovirus antigens.

**[00172]** The immunogenic composition may be used to elicit systemic and/or mucosal immunity, preferably to elicit an enhanced systemic and/or mucosal immunity.

**[00173]** Preferably the immune response is characterized by the induction of a serum IgG and/or intestinal IgA immune response.

**[00174]** As noted above, prime-boost methods are preferably employed where one or more gene delivery vectors and/or polypeptide antigens are delivered in a “priming” step and, subsequently, one or more second gene delivery vectors and/or polypeptide antigens are delivered in a “boosting” step. In certain embodiments, priming and boosting with one or more gene delivery vectors or polypeptide antigens described herein is followed by additional boosting with one or more polypeptide-containing compositions (e.g., polypeptides comprising parvovirus antigens). A VLP is a polypeptide-containing composition. VLPs may be formed in a host immunized with a suitable gene delivery vector.

**[00175]** In any method involving co-administration, the various compositions can be delivered in any order. Thus, in embodiments including delivery of multiple different compositions or molecules, the nucleic acids need not be all delivered before the

polypeptides. For example, the priming step may include delivery of one or more polypeptides and the boosting comprises delivery of one or more nucleic acids and/or one or more polypeptides. Multiple polypeptide administrations can be followed by multiple nucleic acid administrations or polypeptide and nucleic acid administrations can be performed in any order. Thus, one or more of the gene delivery vectors described herein and one or more of the polypeptides described herein can be co-administered in any order and via any administration route. Therefore, any combination of polynucleotides and polypeptides described herein can be used to elicit an immune reaction.

#### **(i). Dosage Regime**

[00176] Dosage treatment can be according to a single dose schedule or a multiple dose schedule. Multiple doses may be used in a primary immunization schedule and/or in a booster immunization schedule. In a multiple dose schedule, the various doses may be given by the same or different routes, e.g. a parenteral prime and mucosal boost, a mucosal prime and parenteral boost, etc. For example, a multiple dose schedule may comprise a prime, followed by two boosts. The prime and/or boost dose will preferably be co-administered with an adjuvant.

[00177] Preferably the dosage regime enhances the avidity of the antibody response leading to antibodies with a neutralizing characteristic. An *in-vitro* neutralization assay may be used to test for neutralizing antibodies (see for example Asanaka *et al.* (2005) *J of Virology* 102: 10327; Wobus *et al.* (2004) *PLOS Biology* 2(12); e432; and Dubekti *et al.* (2002) *J Medical Virology* 66: 400).

[00178] There is a strong case for a correlation between serum antibody levels and protection from disease caused by parvovirus.

[00179] Parvovirus VLPs as described above can be administered to a mammal, such as a mouse, baboon, chimpanzee, or human, to activate parvovirus-specific T cells *in vivo*. Administration can be by any means known in the art, including parenteral, intranasal, intramuscular or subcutaneous injection, including injection using a needle or syringe.

[00180] A composition of the invention comprising a parvovirus VLP is administered in a manner compatible with the particular composition used and in an amount

which is effective to induce an immune response (e.g., a T cell response and/or a humoral response), preferably a protective immune response.

[00181] Parvovirus-specific T cell responses can be measured by, *inter alia*, a 51Cr release assay, a lymphoproliferation assay, or by intracellular staining for IFN- $\gamma$ . The proteins can be administered either to a mammal which is not infected with a parvovirus or can be administered to a parvovirus-infected mammal. The particular dosages of the protein in a composition will depend on many factors including, but not limited to the species, age, and general condition of the mammal to which the composition is administered, and the mode of administration of the composition. An effective amount of the composition of the invention can be readily determined using only routine experimentation. *In vitro* and *in vivo* models can be employed to identify appropriate doses. Generally, 0.5, 0.75, 1.0, 1.5, 2.0, 2.5, 5, 10, 20 or 50 mg of a parvovirus polypeptide or VLP will be administered to a large mammal, such as a baboon, chimpanzee, or human. If desired, co-stimulatory molecules or adjuvants can also be provided before, after, or together with the compositions.

[00182] Immune responses of the mammal generated by the delivery of a composition of the invention, including activation of parvovirus-specific T cells, can be enhanced by varying the dosage, route of administration, or boosting regimens. Compositions of the invention may be given in a single dose schedule, or preferably in a multiple dose schedule in which a primary course of vaccination includes 1-10 separate doses, followed by other doses given at subsequent time intervals required to maintain and/or reinforce an immune response, for example, at 1-4 months for a second dose, and if needed, a subsequent dose or doses after several months.

#### **F. Tests to Determine the Efficacy of an Immune Response**

[00183] One way of assessing efficacy of therapeutic treatment involves monitoring infection after administration of a composition of the invention. One way of assessing efficacy of prophylactic treatment involves monitoring immune responses against the antigens in the compositions of the invention after administration of the composition.

[00184] Another way of assessing the immunogenicity of the component proteins of the immunogenic compositions of the present invention is to express the proteins recombinantly and to screen patient sera or mucosal secretions by immunoblot. A positive reaction between the protein and the patient serum indicates that the patient has previously

mounted an immune response to the protein in question—that is, the protein is an immunogen. This method may also be used to identify immunodominant proteins and/or epitopes.

**[00185]** Another way of checking efficacy of therapeutic treatment involves monitoring infection after administration of the compositions of the invention. One way of checking efficacy of prophylactic treatment involves monitoring immune responses systemically (such as monitoring the level of IgG1 and IgG2a production) and/or mucosally (such as monitoring the level of IgA production) against the antigens in the compositions of the invention after administration of the composition. Typically, serum specific antibody responses are determined post-immunization but pre-challenge whereas mucosal specific antibody body responses are determined post-immunization and post-challenge.

**[00186]** There is currently no accepted animal model for evaluating protective immunity induced by the human parvovirus (e.g., parvovirus B19) immunogenic compositions of the present invention. However, the capacity of such compositions to induce an immune response can be assessed in suitable animals, as described herein. Other, non-human parvovirus, immunogenic compositions of the present invention can be evaluated in *in vitro* and *in vivo* animal models prior to host, e.g., human, administration. Particularly useful mouse models include those in which intraperitoneal immunization is followed by either intraperitoneal challenge or intranasal challenge.

**[00187]** The efficacy of immunogenic compositions of the invention can also be determined *in vivo* by challenging animal models of infection, e.g., guinea pigs or mice or rhesus macaques, with the immunogenic compositions. The immunogenic compositions may or may not be derived from the same strains as the challenge strains. Preferably the immunogenic compositions are derivable from the same strains as the challenge strains.

**[00188]** The immune response may be one or both of a TH1 immune response and a TH2 response. The immune response may be an improved or an enhanced or an altered immune response. The immune response may be one or both of a systemic and a mucosal immune response. Preferably the immune response is an enhanced systemic and/or mucosal response.

**[00189]** An enhanced systemic and/or mucosal immunity is reflected in an enhanced TH1 and/or TH2 immune response. Preferably, the enhanced immune response

includes an increase in the production of IgG1 and/or IgG2a and/or IgA. Preferably the mucosal immune response is a TH2 immune response. Preferably, the mucosal immune response includes an increase in the production of IgA.

**[00190]** Activated TH2 cells enhance antibody production and are therefore of value in responding to extracellular infections. Activated TH2 cells may secrete one or more of IL-4, IL-5, IL-6, and IL-10. A TH2 immune response may result in the production of IgG1, IgE, IgA and memory B cells for future protection.

**[00191]** A TH2 immune response may include one or more of an increase in one or more of the cytokines associated with a TH2 immune response (such as IL-4, IL-5, IL-6 and IL-10), or an increase in the production of IgG1, IgE, IgA and memory B cells. Preferably, the enhanced TH2 immune response will include an increase in IgG1 production.

**[00192]** A TH1 immune response may include one or more of an increase in CTLs, an increase in one or more of the cytokines associated with a TH1 immune response (such as IL-2, IFN $\gamma$ , and TNF $\beta$ ), an increase in activated macrophages, an increase in NK activity, or an increase in the production of IgG2a. Preferably, the enhanced TH1 immune response will include an increase in IgG2a production.

**[00193]** Immunogenic compositions of the invention, in particular, immunogenic composition comprising one or more antigens of the present invention may be used either alone or in combination with other antigens optionally with an immunoregulatory agent capable of eliciting a Th1 and/or Th2 response.

**[00194]** The immune response may be an antibody-mediated immune mechanism. Preferably, the antibody is a serum antibody, more preferably, a neutralizing serum antibody. The immunogenic compositions of the present invention may be tested primarily by assessing antibody titers (e.g., total antibody and/or neutralizing antibody), using any suitable method, such as western blots or ELISA for evaluation of immunoreactivity against parvovirus VP1 and/or VP2 (Wong *et al.*, J. of Virology, 82(5):2470-2476 (2008); Bostic *et al.*, J. Infectious Disease, 179:619-626 (1999)).

**[00195]** The immunogenic compositions of the invention will preferably elicit both a cell mediated immune response as well as a humoral immune response in order to effectively address an infection. This immune response will preferably induce long lasting (e.g., neutralizing) antibodies and a cell mediated immunity that can quickly respond upon exposure to one or more infectious antigens. By way of example, evidence of neutralizing

antibodies in patients blood samples is considered as a surrogate parameter for protection since their formation is of decisive importance for virus elimination in parvovirus infections (see Young and Brown, NEJM 350:586-97, 2004).

**[00196]** The invention also comprises an immunogenic composition comprising one or more immunoregulatory agent, such as a mineral salt, such as an aluminium salt and an oligonucleotide containing a CpG motif. Most preferably, the immunogenic composition includes both an aluminium salt and an oligonucleotide containing a CpG motif. Alternatively, the immunogenic composition includes an ADP ribosylating toxin, such as a detoxified ADP ribosylating toxin and an oligonucleotide containing a CpG motif. Preferably, the one or more immunoregulatory agents include an adjuvant. The adjuvant may be selected from one or more of the group consisting of a TH1 adjuvant and TH2 adjuvant, further discussed above.

#### **G. Use of the Immunogenic Compositions as Medicaments**

**[00197]** The invention also provides a composition of the invention for use as a medicament. The medicament is preferably able to raise an immune response in a mammal (i.e. it is an immunogenic composition) and is more preferably a vaccine. The invention also provides the use of the compositions of the invention in the manufacture of a medicament for raising an immune response in a mammal. The medicament is preferably a vaccine. Preferably the vaccine is used to prevent and/or treat an erythema infectiosum in children, transient aplastic crisis in patients with red blood cell dyscrasias, usually sickle cell anemia, hemolytic anemia or hereditary spherocytosis, chronic red blood cell aplasia and anemia in immunodeficient patients, persistent infection in immunosuppressed individuals, persistent arthropathy in adults, persistent and sometimes lethal cytopenias in immunocompromised patients, exacerbation of anemia during co-infection with Plasmodium, exacerbated anemia in malaria and hydrops fetalis or spontaneous abortion in pregnant women.

**[00198]** The invention provides methods for inducing or increasing an immune response using the compositions described herein. The immune response is preferably protective and can include antibodies and/or cell-mediated immunity (including systemic and mucosal immunity). Immune responses include booster responses.

**[00199]** The invention also provides a method for raising an immune response in a mammal comprising the step of administering an effective amount of a composition of the

invention. The immune response is preferably protective and preferably involves antibodies and/or cell-mediated immunity. Preferably, the immune response includes one or both of a TH1 immune response and a TH2 immune response. The method may raise a booster response.

**[00200]** The mammal is preferably a human. Where the immunogenic composition, preferably a vaccine is for prophylactic use, the human is preferably a child (e.g. a toddler or infant, preferably pre-school, preferably one year or less or from three years (preferably 1-4 years) onwards) or a teenager (an adolescent); where the vaccine is for therapeutic use, the human is preferably a teenager or an adult. A vaccine intended for children may also be administered to adults, e.g. to assess safety, dosage, immunogenicity, etc. Preferably, the human is a teenager(an adolescent). More preferably, the human is a pre-adolescent teenager. Even more preferably, the human is a pre-adolescent female or male. Preferably the pre-adolescent male or female is around 9-12 years of age. Preferably the adolescent male or female is around 15-19 years of age. Preferably the male or female is around 20-49 years of age. Preferably the male or female is over 49 years of age. Preferably the human is elderly, preferably around 60-80 years of age.

**[00201]** Other populations who can benefit from the immunogenic compositions (e.g., vaccines) of the present invention include: children aged six to ten years; day care children; school-aged children and pediatric and/or elderly populations as discussed above; chronic anemia patients; children with sickle cell anemia; children with hereditary spherocytosis; female children, adolescent or adult females of child-bearing age, e.g., for prevention of congenital infection (fetal infection); children living in geographic areas at risk for malaria or hookworm (e.g. areas where malaria or hookworm are endemic); children or adults who will become immunosuppressed (e.g., cancer patients, patients awaiting transplantation); children or adults who are planning surgical procedures that are likely to involve blood loss and require transfusion; transplant recipients, and immunocompromised individuals.

## **H. Kits**

**[00202]** The invention also provides kits comprising one or more containers of immunogenic compositions of the invention. Compositions can be in liquid form or can be lyophilized, as can individual antigens. Suitable containers for the compositions include, for

example, bottles, vials, syringes, and test tubes. Containers can be formed from a variety of materials, including glass or plastic. A container may have a sterile access port (for example, the container may be an intravenous solution bag or a vial having a stopper pierceable by a hypodermic injection needle).

**[00203]** The kit can further comprise a second container comprising a pharmaceutically-acceptable buffer, such as phosphate-buffered saline, Ringer's solution, or dextrose solution. It can also contain other materials useful to the end-user, including other pharmaceutically acceptable formulating solutions such as buffers, diluents, filters, needles, and syringes or other delivery device. The kit may further include a third component comprising an adjuvant.

**[00204]** The kit can also comprise a package insert containing written instructions for methods of inducing immunity or for treating infections. The package insert can be an unapproved draft package insert or can be a package insert approved by the Food and Drug Administration (FDA) or other regulatory body.

**[00205]** The invention also provides a delivery device pre-filled with the immunogenic compositions of the invention, such as a syringe with or without a hypodermic needle.

**[00206]** The invention also provides diagnostic kits that use VLPs produced by the method described above as antigens to determine seropositivity for parvovirus. The diagnostic kit can contain a parvovirus VLP as described herein, and one or more ancillary reagents to determine whether an individual is seropositive for antibodies that bind parvovirus. Suitable ancillary reagents include, for example, buffers, secondary or detecting antibodies (e.g., a labeled anti-human Ig antibody, a labeled anti-dog Ig antibody), detection agents and the like. In one example, the kit contains a parvovirus VLP and ancillary reagents for performing an ELISA or other suitable immunoassay. The kit can further comprise instructions for performing the diagnostic assay.

### III. EXEMPLIFICATION

**[00207]** Below are examples of specific embodiments for carrying out the present invention. The examples are offered for illustrative purposes only, and are not intended to limit the scope of the present invention in any way.

**[00208]** The initial plasmid containing VP1 and VP2 under the control of two equal strength promoters was generated using traditional molecular biological techniques. The promoter preceding VP1 was then modified as mentioned above (e.g. TATA deletions, various deletions of the A-rich upstream sequence, introduction of start/stop sites upstream of the VP1 start codon).

Table 1: List of oligonucleotides used in the different cloning steps:

|             |                                                                                            |
|-------------|--------------------------------------------------------------------------------------------|
| C489G_A490C | 5'-CTGCTGTTGATTCTGCTGCTAGAAATTGCTGATTTCAGATACTCTCAATT-3'<br>(SEQ ID NO: 3)                 |
| VP1-MS1:    | 5'-AGATCTACGCGTACAAAACAAAATGTCTAAGAAATCTGGTAAATGG -3'<br>(SEQ ID NO: 4)                    |
| VP1-MS2     | 5'-AGATCTGTCGACTCATTACAATGGATGAACTCTAGATTTAGCAGTCC-3'<br>(SEQ ID NO: 5)                    |
| VP2-AN1:    | 5'-AGATCTCCTAGGACAAAACAAAATGACATCTGTAAATTCTGCTGAAGC-3'<br>(SEQ ID NO: 6)                   |
| VP2-AN2 :   | 5'-AGATCTGCGGCCGCTCATTACAATGGATGAACTCTAGATTTAGCAGTC-3'<br>(SEQ ID NO: 7)                   |
| dTATA.1 :   | 5'-AGATCTGGATCCTTCAATATGCGCACATACGCTG-3' (SEQ ID NO: 8)                                    |
| dTATA.2 :   | 5'-TCTAGACCTAGGGGAATAATTCAGGGAAGTGGTTCAACC-3' (SEQ ID NO: 9)                               |
| am-1 :      | 5'-CTAGCATTGTAATTCTGTAAATCTATTTCTTAACTTC-3' (SEQ ID NO: 10)                                |
| am-2 :      | 5'-TTAAATTCTACTTTTATAGTTAGTCTTTTTTTTAGTTTTTAAAC-3' (SEQ ID NO: 11)                         |
| am-3 :      | 5'-ACCAAGAACTTAGTTTCGAATAAACACACATAAACAAACA-3' (SEQ ID NO: 12)                             |
| am-4 :      | 5'-CGCGTGTTTGTTTATGTGTGTTTATTCGAACTAAGTTCTTGGTGTTTT<br>AAACTAAAAAAAAGAC-3' (SEQ ID NO: 13) |
| am-5 :      | 5'-TAACTATAAAAGTAGAATTTAAGAAGTTTAAGAAATAGATTTACAGAATTA<br>CAATG-3' (SEQ ID NO: 14)         |

**[00209]** The first step was to generate PCR products for VP1 with MluI-SalI restriction ends appropriate for cloning in the pCDC7 yeast expression vector (FIG. 2) and for VP2 with AvrII-NotI restriction ends appropriate for cloning into the second expression cassette of pCDC7. In addition, an ACAAACAAA (SEQ ID NO: 15) sequence was

introduced between the 5' end restriction site and the initiating Met of the VP1 or VP2 sequence.

**[00210]** For VP2, the PCR amplification reaction contained 100ng pMK-RG\_parvo\_shade\_vp1, 50 pmoles each of primers VP2-AN1 and VP2-AN2, 10ul of 10X Buffer with MgCl<sub>2</sub>, 16ul of dNTPs 1.25mM each, H<sub>2</sub>O up to 100ul and 0.75ul DNA polymerase from Roche Expand High Fidelity PCR kit. Amplification conditions : 5' 95°C, 35 cycles : 30'' 95°C, 1' 60°C, 2' 72°C, followed by a single 7' 72°C extension cycle and a 4°C hold.

**[00211]** The VP2 PCR reaction was purified using the Promega Wizard PCR Preps DNA Purification System and digested with AvrII and NotI for 3 hours at 37°C with 120u of each enzyme in a 300ul final volume. The 1685bp AvrII-NotI fragment was then purified from 1% agarose GTG (genetic technology grade) using the Qiagen MinElute Gel Extraction kit.

**[00212]** 15ng of the VP2 AvrII-NotI gel-purified PCR product was ligated to a dephosphorylated, gel-purified pT7Blue2 AvrII-NotI vector using the Roche Rapid DNA Ligation kit. After 20' at RT° the ligation reaction was used to transform competent HB101, then plated on LB agar plates containing 100ug/ml ampicillin. Miniprep plasmid DNA was prepared from single ampicillin resistant colonies grown in Terrific Broth containing 100ug/ml ampicillin; a portion of the DNA was digested with AvrII+NotI to confirm the presence of the VP2 insert. The plasmid from 3 positive clones was analyzed with sequencing.

**[00213]** For VP1, the PCR amplification reaction contained 100ng pMK-RG\_parvo\_shade\_vp1, 50 pmoles each of primers VP1-MS1 and VP1-MS2, 40ul of 5' Hot Master Mix (2.5X) and H<sub>2</sub>O up to 100ul. Amplification conditions : 5' 95°C, 35 cycles : 30'' 95°C, 1' 60°C, 2' 72°C, followed by a single 7' 72°C extension cycle and a 4°C hold.

**[00214]** The VP1 PCR reactions were purified using the Promega Wizard PCR Preps DNA Purification System and digested with MluI and Sall for 3 hours at 37°C with 120u of each enzyme in a 300ul final volume. The 2366bp MluI-Sall fragments were then purified from 1% agarose GTG (genetic technology grade) using the Qiagen MinElute Gel Extraction kit.

**[00215]** An aliquot of VP1 MluI-Sall gel-purified PCR product was ligated to a dephosphorylated, gel-purified pGEM4Z MluI-Sall vector using the Roche Rapid DNA

Ligation kit. After 20' at RT° the ligation reaction was used to transform competent HB101, then plated on LB agar plates containing 100ug/ml ampicillin. Miniprep plasmid DNA was prepared from single ampicillin resistant colonies grown in Terrific Broth containing 100ug/ml ampicillin; a portion of the DNA was digested with MluI+SalI to confirm the presence of the VP1 insert. The plasmid from 3 or 4 positive clones was analyzed with sequencing.

**[00216]** Next, the VP2 and the VP1 restriction fragments were prepared from sequence confirmed subclones. For VP2, 10ug of pT7Blue2.AvrII.NotI.Parvo.VP2 #11 were digested for 2.5 hours at 37°C with 80 units each of AvrII and NotI in a 200ul final volume. For VP1, 12ug of pGEM4Z.opti.Parvo.VP1 #3 was digested for 3 hours at 37°C with 100u each MluI and SalI in a 300ul final volume. PvuI enzyme was included in the VP1 prep digest in order to prevent contamination of the VP1 fragment with vector. The VP2 and VP1 fragments were purified from 1% agarose GTG using the Qiagen MinElute Gel Extraction kit.

**[00217]** The bicistronic yeast expression vector pCDC7 was digested with AvrII+NotI, dephosphorylated with calf intestine alkaline phosphatase and then purified from 1% agarose GTG using the Qiagen MinElute Gel Extraction kit. Approximately 90ng of the VP2 AvrII-NotI fragment was ligated to ~30ng of the AvrII-NotI pCDC7 vector in a 20ul reaction using the Roche Rapid DNA Ligation kit. After 20' at RT° the ligation reaction was used to transform competent HB101, then plated on LB agar plates containing 100ug/ml ampicillin. Miniprep plasmid DNA was prepared from single ampicillin resistant colonies grown in Terrific Broth containing 100ug/ml ampicillin; the plasmid DNA was digested with AvrII+NotI to confirm the presence of the VP2 insert.

**[00218]** To prepare a vector for the cloning of VP1 into the second expression cassette of pCDC7, approximately 10ug of pCDC7.ParvoB19.VP2 #11 plasmid DNA from were digested for 3 hours at 37°C with 100u each of MluI and SalI in a 200ul final volume. The digested plasmid was then dephosphorylated with 7u calf intestine alkaline phosphatase for 1 hour at 37°. Next, the vector was purified from 1% agarose GTG using the Qiagen MinElute Gel Extraction kit.

**[00219]** Finally, ~100ng of gel-purified VP1 MluI-SalI fragment was ligated to ~60ng of the pCDC7.ParvoB19.VP2 #11 MluI-SalI vector in two separate 20ul ligation reactions using the Roche Rapid Ligation kit. After 20' at RT° the ligation reaction was used to transform competent HB101, then plated on LB agar plates containing 100ug/ml

ampicillin. Miniprep plasmid DNA was prepared from single ampicillin resistant colonies grown in Terrific Broth containing 100ug/ml ampicillin; the plasmid DNA was digested with MluI+Sall to confirm the presence of the VP1insert. pCDC7.ParvoB19.VP2.VP1 #2 was selected for amplification in 100ml Terrific Broth containing 100ug/ml ampicillin and subsequent plasmid DNA purification using a Qiagen MaxiPrep kit.

[00220] Next, the  $\Delta$ TATA promoter restriction fragment was prepared from the sequence confirmed subclone. Approximately 8ug of pT7Blue2. $\Delta$ TATA #9 was digested for 4 hours at 37°C with 120 units each of BamHI and MluI in a 300ul final volume. A BamHI-MluI yeast expression vector was prepared by digesting 5ug pCDC7.ParvoB19.VP2.VP1#2 with 80 units each of BamHI and MluI in a 200ul reaction volume, followed by a 1 hour 37°C dephosphorylation with 10u calf intestine alkaline phosphatase. The promoter and the vector fragments were both purified from 1% agarose GTG using the Qiagen MinElute Gel Extraction kit.

[00221] Finally, ~20ng of the 1333bp BamHI-MluI  $\Delta$ TATA promoter fragment was ligated to ~20ng of BamHI-MluI pCDC7 vector (described in the previous paragraph) in a 20ul reaction using the Roche Rapid DNA Ligation kit. After 20' at RT° the ligation reaction was used to transform competent HB101, then plated on LB agar plates containing 100ug/ml ampicillin. Miniprep plasmid DNA was prepared from single ampicillin resistant colonies grown in Terrific Broth containing 100ug/ml ampicillin; the DNA was digested with BamHI + MluI to confirm the presence of the 1333bp  $\Delta$ TATA promoter. pCDC7.ParvoB19.VP2+ $\Delta$ TATA.VP1 #16 was selected for amplification in 100ml Terrific Broth containing 100ug/ml ampicillin and subsequent plasmid DNA purification using a Qiagen MaxiPrep kit.

### Example 1

#### Expression of Parvovirus B19 in Yeast

[00222] Parvovirus B19 VP2 and VP1 genes were co-expressed in *S.cerevisiae* AD3, characterized as follows : *MATa*, *leu2*, *ura3-52*, *prb1-1122*, *pep4-3*, *prc1-407*, *gal2*, [*cir*°] ::pDM15(pGAP/ADR1::G418R),::Yip5 $\Delta$ leuAD.

[00223] The yeast are streaked on YEPD plates before selecting a single colony for preparation of competent cells for transformation. Yeast transformation was performed using the Invitrogen S.C. EasyComp Transformation kit. Transformants were obtained on ura-8%

glucose selective plates. Several transformants were streaked for single colonies on ura-8% glucose selective plates. Single colonies were then patched on leu-8% glucose selective plates in order to increase plasmid copy number.

**[00224]** A small-scale expression check was done for several of the single colony patches. First, a 3 ml overnight culture in leu-7.1% glucose media was grown. Then, the leu-culture was diluted 1:20 into YEPD or Veggie YEPD. The culture was shaken at 30°C or 25°C for 48-72 hours. Induction occurred upon depletion of the glucose from the media.

**[00225]** The cell pellets were thawed, washed in water and pelleted in a graduated tube so that an estimate could be made for the packed cell volume. Each pellet was then resuspended in a volume of cold lysis buffer (10mM NaPO<sub>4</sub> pH7.5, 0.1% TritonX-100, 0.5M NaCl) equal to the packed cell volume, in order to normalize for differences in the density of different expression cultures. An aliquot of the cell / buffer suspension was lysed with washed glass beads (Sigma G8772) by vortexing at top speed for 1hr at 4°. The crude lysates were centrifuged at 13K for 30min in a 4°C microfuge to pellet the insoluble fraction. After the protein concentration of the supernatants was determined, 300ug samples were prepared in 150ul SDS sample buffer + 50mM DTT. Meanwhile, the insoluble pellet fraction (IP) was resuspended in 1ml SDS sample buffer + 50mM DTT. The samples for soluble and insoluble fractions were boiled before loading onto 4-20% Tris-Glycine gels. For the soluble fraction, 10 µl (20ug total protein) were loaded ; for the insoluble pellet, 5ul were loaded.

**[00226]** Gels were run in duplicate—one for Coomassie staining (Invitrogen SimplyBlue Safestain) and one for western blot which was probed with a mouse monoclonal antibody to parvovirus B19 (Santa Cruz sc-58178).

**[00227]** Using the same protocol, four different colonies were analyzed to get a sense of the variability from colony to colony. Four 72 hour shake-flask expression cultures were inoculated with cells from four single-colony Leu- patches. The western blot in Fig. 9b. shows the range of variability for soluble VP2 and VP1 from four colonies of each construct.

## **Example 2**

### **Parvovirus B19 Purification Protocol**

**[00228]** BCA Protein Assay (Pierce) using bovine serum albumin (BSA) was used as a standard reference for protein concentration determination. In some cases UV absorbance at 280 nm was used, assuming 1 absorbance unit = 1 mg/ml as indicated by calibrated with BCA results.

[00229] SDS-PAGE was performed with gels from Invitrogen (NuPAGE 4-12% Bis-Tris cat#NP0322BOX) and NuPAGE MES SDS Running Buffer (Cat#NP0002-02). Markers used for SDS-PAGE were the Full range rainbow molecular weight marker (GE Healthcare, cat# RPN800E). Native PAGE 3-12% Bis-Tris Gels were used (Invitrogen cat# BN1003BOX) with the NativePAGE Running buffer kit (Invitrogen cat# BN2007 and Native PAGE Markers (Invitrogen cat#151-1901). SDS PAGE gels were stained using Coomassie Blue R-250 Stain Solution from Teknova (cat# C1050).

[00230] The SDS-PAGE gel was blotted on a nitro-cellulose membrane by I-Blot System (Invitrogen) for purity and identity by western blot analysis.. For yeast contaminant determinations a Yeast Whole Cell ELISA kit was used (*S. cerevisiae* HCP ELISA Kit-CYGNUS Technologies, #F135).

[00231] Parvovirus B19 virus-like particles (VLPs) were purified from yeast cells expressing both VP1 and VP2. Yeast cells were resuspended in lysis buffer and the cells were broken open with multiple high pressure (25,000 psi) passes in a Microfluidizer. Following a low speed spin (15,000 x g) to separate cellular debris, the supernatant was subjected to a four hour high speed spin (100,000 x g) into a 20%/70% sucrose step cushion. The cushion was fractionated and the fractions containing VLPs were diluted with a buffer (20 mM Tris pH 7.5, 100 mM NaCl). The VLPs were loaded onto a capto Q column and eluted with high salt. The eluted fractions containing the VLPs were concentrated and buffer exchanged into a lower salt buffer (20 mM Tris pH 7.5, 100 mM NaCl) and stored at 4°C.

[00232] Parvovirus B19 VP1/VP2 VLPs were purified to greater than 90% purity and appeared to be identically sized (~26 nm) (FIG. 11A, 11B). The gels are based on the VP2/VP1 (delta TATA) VLPs. Additional VLPs were constructed that contained a site specific mutant.

#### Sucrose Shelf Cushion and Fractionation

[00233] A sucrose shelf was prepared in a 38 mL Polyallomer tube (Seton Scientific). The sucrose shelf has two layers : 5 mL of 20% sucrose and 26 mL of 70% sucrose. 6 mL of clarified lysate was added gently atop the sucrose. This step has been performed with up to 8 mL of clarified lysate with 2 mLs less 70% sucrose being used. The sucrose shelf was centrifuged for 4 hours at 100,000 xg in a JS-24 rotor at 4°C.

**[00234]** The sucrose shelf was fractionated from the top with a Piston Gradient Fractionator (BioComp) into 1 mL fractions. The VLPs of interest were found at the 20%:70% sucrose interface. The fractions that contained the VLPs were at a Brix value between 28.5 and 38.5 (generally between fractions 12 and 14). Westerns with a commercial antibody that recognizes both VP1 and VP2 (Santa Cruz BioTech cat# sc-71852) were performed to check for the location and concentration of the VLPs. This material was stored at 4°C for at least 1 week with no protein degradation.

#### Capto Q chromatography

**[00235]** A Western analysis with VP1/VP2 specific antibody (Santa Cruz BioTech cat# sc-71852) identified the fractions from the sucrose shelf that contained the greatest amount of protein. These fractions were either pooled or kept separate (based on a particular run). These fractions were diluted 1:4 in Buffer A (25 mM Tris, 100 mM NaCl pH 7.5) and loaded over a pre-washed and pre-equilibrated column at 6 cm/hr. The sucrose still present in the loaded protein partially prevented the protein from binding to the column at faster flow rates (i.e. 60 cm/hr). A NaCl gradient was used to elute the bound protein. Chromatography was run as follows:

**[00236]** Buffer A: 25 mM Tris, 100 mM NaCl pH 7.5; Buffer B: 25 mM Tris, 2 M NaCl pH 7.5; Washing: 5 CV with Buffer A; Gradient: 0-40%B (10 CV); Washing: 100%B 5CV; Flow: 0.2 ml/min. Fractions: 1 ml during elution. Pool from ~5% B to ~10% B. 11.6. Buffer Exchange and Concentration

**[00237]** The Capto Q fractions were selected such that there was no band of greater or equal intensity to the VP1 band. These fractions were pooled and buffer exchanged into 25 mM Tris, 100 mM NaCl through a concentration and dilution method using 50 kDa cutoff Ultra 4 (Millipore- regenerated cellulose) concentrator. The final concentration was ~0.8 mg/mL. This process resulted in material that is >95% pure with low to not detectable levels of endotoxin. This material could be stored for >1 month prior to use. General yields were approximately 1.5 mg of purified material for 20 grams of biomass (the result of a 1 L shake flask).

Table 2. VP1/VP2 VLP production inoculation ranging.

|                                             |      |      |
|---------------------------------------------|------|------|
| Lot                                         |      |      |
| Inoc OD                                     | .1   | .9   |
| Wet Weight per Liter (g)                    | 0.3  | 1.2  |
| Percent VP1 to VP2                          | 1    | 6    |
| Yield ( $\mu$ g protein/g Wet Weight cells) | 6    | 8    |
| Purity                                      | 95%  | 95%  |
| Endotoxin level (EU / 10 $\mu$ g dose)      | .264 | .734 |

## EXAMPLE 3

## Analysis of parvovirus B19 VLPs

## VLP Purity

[00238] VLP purity was shown by Coomassie and western analysis. An example of a Coomassie stained SDS-PAGE and corresponding western is shown in Fig. 16

## VLP identification

[00239] VLP identification was carried out using two antibodies. One antibody (Santa Cruz BioTech cat# sc-71852) recognizes the common VP2 region and hence can identify both VP1 and VP2 ( $\alpha$ -VP1/VP2 antibody). A second antibody (USBio P3113-81D) recognizes just the VP1 unique region and hence can identify VP1 only. A dimer of VP2 is still visible if the fractions are not fully boiled are reduced. As expected, the VP2 band is not present but the same VP1 band is visible when the  $\alpha$ -VP1 antibody is used to probe the membrane.

## VLP characteristics

[00240] VLPs were selected using a sucrose shelf to remove protein not assembled into VLPs. The purified VLPs were analyzed for monomer content using SEC, native PAGE,

DLS and electron microscopy. The results confirmed that the VLPs were >99% in a single large molecular weight species representative of a VLP (FIG. 17).

#### VLP Stability

**[00241]** The VLPs appeared stable in both MF59 and in 25 mM Tris, 100 mM NaCl buffer for at least 3 months at 4°C. This was examined with Coomassie stained SDS-PAGE, Western with parvovirus specific antibodies, native PAGE and electron microscopy. Furthermore, the VLPs could be disrupted into monomers using 0.05% SDS but not 0.5% Triton X-100.

#### Endotoxin examination

**[00242]** The protein was analyzed for endotoxin with the Charles River Laboratories Endosafe-PTS system. This is a cartridge based system (Product Code PTS20) using Limulus Amebocyte Lysate detection of endotoxin. The assay was performed with Endotoxin Specific Buffer (Product Code BG120) as directed by the manufacturer to decrease spurious background signal generated by  $\beta$ -glucan (a component of the yeast cell wall) through a secondary activation pathway. The results from multiple assays demonstrated the range per 5 ug dose of material would be from 0.036 EU/dose to below the detectable limit of 0.013 EU/dose.

### Example 4

#### Mouse Immunogenicity Data

**[00243]** Immunogenicity was studied in Balb/c mice (n=10). Pre-immune serum was taken on the day before the first immunization. Immunizations were administered by IM at various concentrations (0.05, 0.5 or 5  $\mu$ g) with different adjuvants (phosphate buffered saline, alum, or MF59). Mice were bled on day 20 and received a second immunization on day 21. The mice were bled again three weeks after the second immunization (3wp2, day 41) and administered a third immunization on day 42. Serum samples were taken again on days 56 (2wp3 – 2 weeks post second) and 84 (6wp3 – 6 weeks post third).

**[00244]** Sera for each vaccine group were pooled (n=5 for PBS vaccinated, n=10 for all other groups). IgG ELISAs were performed using plates coated with the parvovirus B19 VLPs (FIG. 12). Highest titers were demonstrated with the MF59 formulations and were similar for each dose concentration.

### Example 5

#### Cell Based Neutralization

**[00245]** The neutralization assay was based on qRT-PCR and used erythroid progenitor cells that were CD36<sup>+</sup> and globoside<sup>+</sup> (the primary cell receptor) as the viral substrate. The assay measured the presence or absence of an RNA sequence that would only be present during infection. To determine neutralizing titers, the sera was pre-incubated with virus prior to mixing with the cells. The cells were harvested after 48 hours and probed for infection. Highest neutralizing titers for immunized mice were  $6 \times 10^4$ . Convalescent human sera tested in this assay gave a neutralizing titer (IC<sub>50</sub>) of  $1 \times 10^4$ .

Table 3: Table of ELISA titers (serum IgG) corresponding neutralizing titers from animal sera two weeks post third immunization.

| VLP (MF59 ADJUVANTED) | Dose (ug) | ELISA titers (serum IgG) | Neutralizing titers (IC <sub>50</sub> ) |
|-----------------------|-----------|--------------------------|-----------------------------------------|
| VP2 alone             | 0.05      | $5 \times 10^5$          | <10                                     |
| VP2 alone             | 0.5       | $5 \times 10^5$          | <10                                     |
| VP2 alone             | 5.0       | $5 \times 10^5$          | <10                                     |
| VP1/VP2               | 5.0       | $5 \times 10^5$          | $6 \times 10^4$                         |

## CLAIMS

1. A method of producing a parvovirus virus like particle (VLP), comprising: (a) providing a host cell that contains a recombinant nucleic acid molecule that comprises a nucleotide sequence encoding parvovirus VP1 that is operably linked to a first control element, and a nucleotide sequence encoding parvovirus VP2 protein that is operably linked to a second control element, and (b) maintaining the host cell under conditions whereby the VP1 and VP2 proteins are expressed and assembled into VLPs.
2. The method of claim 1, wherein VP1 is produced in lower abundance than VP2.
3. The method of claim 2, wherein VLPs that contain VP1 and VP2 are produced.
4. The method of claim 1, wherein the recombinant nucleic acid molecule is a bicistronic vector.
5. The method of claim 1, wherein said first control element is a variant of said second control element, and the variant comprises a modification that transcriptionally, translationally or transcriptionally and translationally decreases production of VP1 relative to VP2.
6. The method of claim 4, wherein the first control element comprises deletion of at least a portion of the TATA box upstream of the nucleic acid encoding VP1, deletion of a junction site upstream of the nucleic acid encoding VP1, introduction of a transcription start or stop site upstream of the nucleic acid encoding VP1, or combinations thereof.
7. The method of claim 1 wherein the second control element is the ADH2/GAPDH promoter (SEQ ID NO: 19).
8. The method of claim 1, wherein the host cell is a yeast cell, insect cell, mammalian cell, avian cell, bacterium, Tetrahymena cell or combinations thereof.
9. The method of claim 1, wherein the host cells are maintained under culture conditions suitable for production of VP1 and VP2 and assembly of the proteins into VLPs.
10. The method of claim 1, further comprising isolating the VLPs.

11. The method of claim 10, wherein the VLPs are isolated from host cell conditioned media, host cell lysate, host cell homogenate, or combinations thereof.
12. The method of claim 11, wherein the isolated VLPs are further purified.
13. The method of claim 12, wherein the isolated VLPs are further purified using a purification method selected from the group consisting of sucrose cushion, sucrose gradient centrifugation, chromatographic methods and combinations thereof.
14. A parvovirus virus like particle (VLP) that contains VP1 and VP2 produced according to the method of any one of claims 1-13.
15. An immunogenic composition comprising a parvovirus VLP according to claim 14.
16. A recombinant nucleic acid comprising a nucleotide sequence encoding parvovirus VP1 that is operably linked to a first control element, and a nucleotide sequence encoding parvovirus VP2 protein that is operably linked to a second control element, wherein when equivalent amounts of the sequence that encodes VP1 and the sequence that encodes VP2 are present in a suitable host cell, VP1 is expressed in lower abundance relative to VP2.
17. The recombinant nucleic acid of claim 16, wherein the first control element is a weak promoter and the second control element is a strong promoter.
18. The recombinant nucleic acid of claim 17, wherein said first control element is a variant of said second control element, and the variant comprises a modification that transcriptionally, translationally or transcriptionally and translationally decreases production of VP1 relative to VP2.
19. The recombinant nucleic acid of claim 18, wherein the modification comprises deletion of at least a portion of the TATA box upstream of the nucleic acid encoding VP1, deletion of a junction site upstream of the nucleic acid encoding VP1, introduction of a transcription start or stop site upstream of the nucleic acid encoding VP1, or combinations thereof.
20. The recombinant nucleic acid of claim 16 wherein the second control element is the ADH2/GAPDH promoter (SEQ ID NO: 19).

21. The recombinant nucleic acid of claim 20, wherein said nucleic acid is in the form of DNA, or RNA, and is either single or double stranded.
22. The recombinant nucleic acid of claim 21, wherein the recombinant nucleic acid is a plasmid.
23. The recombinant nucleic acid of claim 22, wherein the plasmid includes a detectable marker.
24. A host cell comprising a recombinant nucleic acid according to any one of claims 16-23.
25. A host cell of claim 24, wherein said recombinant nucleic acid is integrated into the genome of the host cell or is carried on an extra-chromosomal element.
26. The host cell according to claim 25, wherein the host cell is a yeast cell, insect cell, mammalian cell, avian cell, bacteria, Tetrahymena cells or combinations thereof.
27. A method of using a parvovirus virus like particle (VLP) that contains VP1 and VP2 produced according to the method of any one of claims 1-13 as an antigen in a diagnostic kit.
28. A method of detecting anti-parvovirus antibodies in a biological sample, comprising providing a biological sample obtained from an individual, combining the sample with a parvovirus VLP according to any one of claims 1-13, and detecting antibody-VLP complexes, wherein the presence of antibody-VLP complexes indicates that anti- parvovirus antibodies are present in the biological sample.
29. The method of claim 28, wherein the biological sample is a blood or serum sample.
30. A method of using an immunogenic composition comprising the recombinant nucleic acid of claim 16 as a medicament, wherein said medicament is capable of raising an immune response in a mammal.
31. The method of claim 28, wherein said medicament is used to treat or prevent a condition selected from the group consisting of erythema infectiosum in children, sickle cell anemia, hemolytic anemia, hereditary spherocytosis, chronic red blood cell aplasia, anemia in

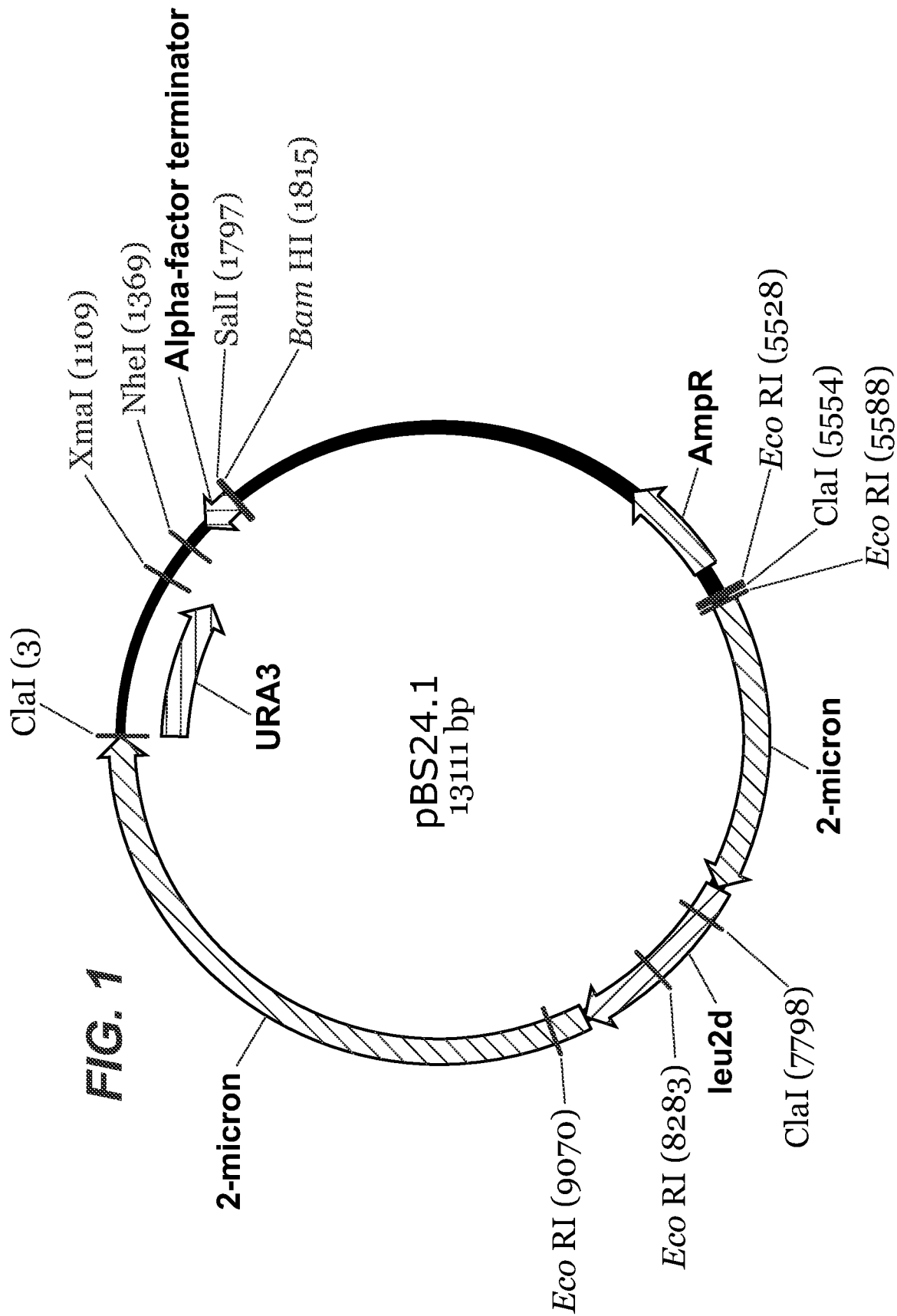
immunodeficient patients, persistent arthropathy in adults, cytopenias in immunocomprised patients, exacerbation of anemia during co-infection with Plasmodium, exacerbated anemia in malaria, hydrops fetalis and spontaneous abortion in pregnant women.

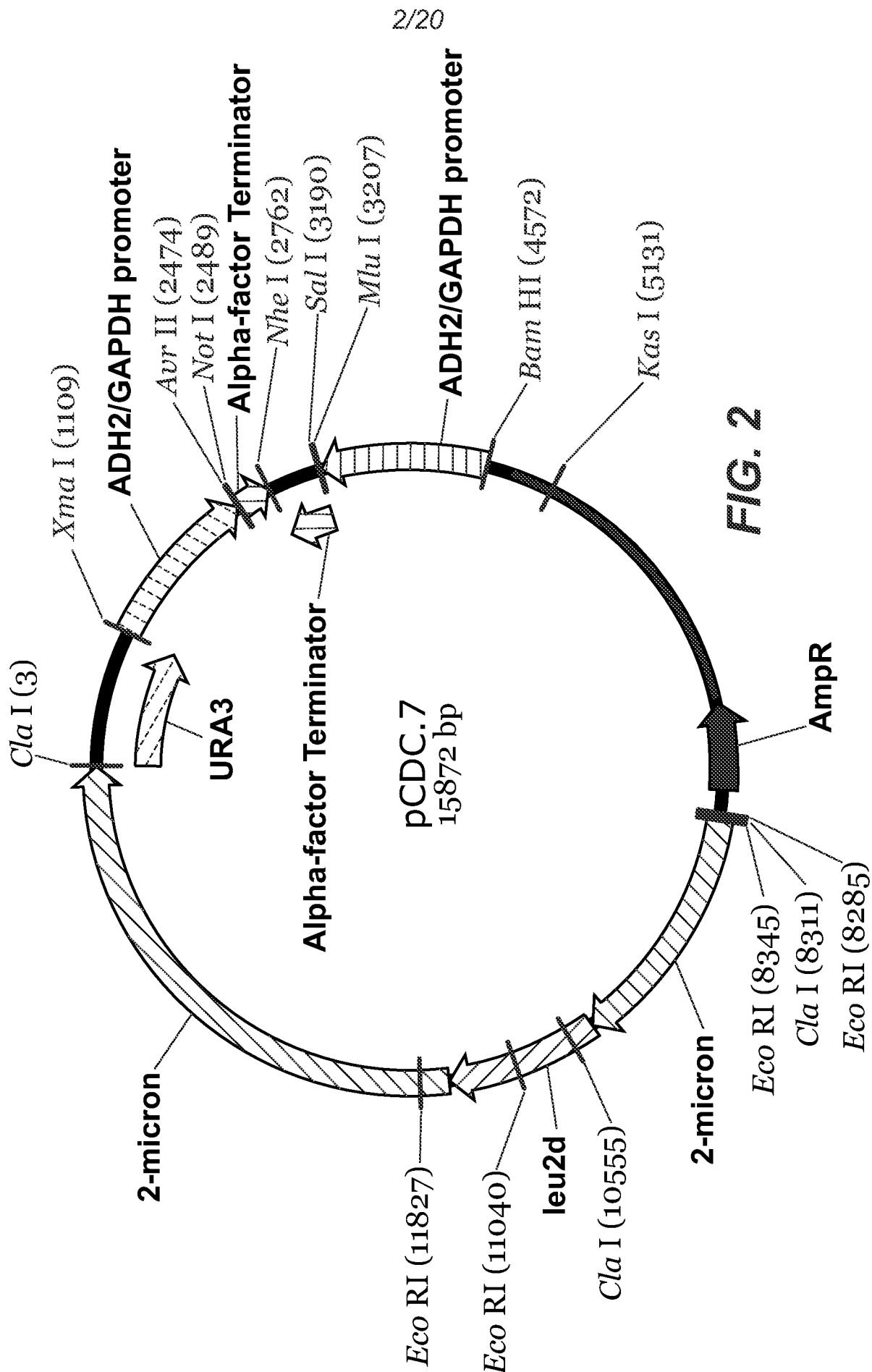
32. A method of inducing an immune response to parvovirus VP1 and/or VP2, comprising administering to an individual an effective amount of a recombinant nucleic acid according to any one of claims 16-23, whereby an immune response is induced.

33. The method of claim 32, wherein the recombinant nucleic acid is administered by intramuscular injection.

34. A method of inducing an immune response to parvovirus VP1 and/or VP2, comprising administering to an individual an effective amount of a virus like particle according to claim 14 or an immunogenic composition according to claim 15, whereby an immune response is induced.

35. The method of claim 34, wherein the virus like or an immunogenic composition is administered by intramuscular injection.





**FIG. 2**

pCDC7

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used for co-expression of ParvoB19 VP1 and VP2 VLPs in *S.cerevisiae*

ATCGATAAGCTTTTCAATTCATCATTTTTTTTTTTTATTCTTTTTTTTGATTTCGGTTTCCTTGAAATTTTTTTTGAT  
 TCGGTAATCTCCGAACAGAAGGAAGAACGAAGGAAGGAGCACAGACTTAGATTGGTATATATACGCATATGTAGT  
 GTTGAAGAAACATGAAATTGCCCAGTATTCTTAACCCAACTGCACAGAACAAAAACCTGCAGGAAACGAAGATAA  
 ATCATGTGCGAAAGCTACATATAAGGAACGTGCTGCTACTCATCCTAGTCCTGTTGCTGCCAAGCTATTTAATATC  
 ATGCACGAAAAGCAAACAACTTTGTGTGCTTCATTGGATGTTTCGTACCACCAAGGAATTACTGGAGTTAGTTGAA  
 GCATTAGGTCCCAAAATTTGTTTACTAAAAACACATGTGGATATCTTGACTGATTTTTCCATGGAGGGCACAGTT  
 AAGCCGCTAAAGGCATTATCCGCCAAGTACAATTTTTTACTCTTCGAAGACAGAAAATTTGCTGACATTGGTAAAT  
 ACAGTCAAAATGCAGTACTCTGCGGGTGTATACAGAATAGCAGAATGGGCAGACATTACGAATGCACACGGTGTG  
 GTGGGCCCCAGGTATTGTTAGCGGTTTGAAGCAGGCGGCAGAAGAAGTAACAAAGGAACCTAGAGGCCCTTTTGATG  
 TTAGCAGAATTGTCAATGCAAGGGCTCCCTATCTACTGGAGAATATACTAAGGGTACTGTTGACATTGCCAAGAGC  
 GACAAAGATTTTGTATCGGCTTTATTGCTCAAAGAGACATGGGTGGAAGAGATGAAGGTTACGATTGGTTGATT  
 ATGACACCCCGTGTGGGTTTAGATGACAAGGGAGACGCATTGGGTCAACAGTATAGAACCGTGCATGATGTGGTC  
 TCTACAGGATCTGACATTATTATTGTTGGAAGAGGACTATTTGCAAAGGGAAGGGATGCTAAGGTAGAGGGTGAA  
 CGTTACAGAAAAGCAGGCTGGGAAGCATATTTGAGAAGATGCGGCCAGCAAACTAAAAAACTGTATTATAAGTA  
 AATGCATGTATACTAACTCACAAATTAGAGCTTCAATTTAATTATATCAGTTATTACCCGGG

TTCAATATGCGCACATACGCTGTTATGTTCAAGGTCCCTTCGTTTAAAGAACGAA  
 AGCGGTCTTCCTTTTGAGGGATGTTTCAAGTTGTTCAAATCTATCAAATTTGCAAATCCC  
 CAGTCTGTATCTAGAGCGTTGAATCGGTGATGCGATTTGTTAATTAAATTTGATGGTGTCA  
 CCATTACCAGGTCTAGATATACCAATGGCAAACCTGAGCACAACAATACCAGTCCGGATCA  
 ACTGGCACCATCTCTCCCGTAGTCTCATCTAATTTTTCTTCCGGATGAGGTTCCAGATAT  
 ACCGCAACACCTTTATTATGGTTTCCCTGAGGGAATAATAGAATGTCCCATTGGAATCA  
 CCAATTCATAACCTGGGCGAATTGTATTTGCGGTTTGTAACTCGTTCCAGTCAGGAATG  
 TTCCACGTGAAGCTATCTTCCAGCAAAGTCTCCACTTCTTCATCAAATTTGTGGAGAATAC  
 TCCCAATGCTCTTATCTATGGGACTTCCGGGAAACACAGTACCGATACTTCCCAATTCGT  
 CTTGAGAGCTCATTGTTTGTGTTGAAGAGACTAATCAAAGAATCGTTTTCTCAAAAAAATTT  
 AATATCTTAACTGATAGTTTGTGATCAAAGGGGCAAACCGTAGGGGCAAACAAACGGAAAAA  
 TCGTTTCTCAAATTTCTGATGCCAAGAAGTCTAACCAGTCTTATCTAAAAATTTGCCCTTA  
 TGATCCGTCTCTCCGGTTACAGCCTGTGTAAGTGAATTAATCCTGCCTTTCTAATCACCAT  
 TCTAATGTTTTAATTAAGGGATTTGTCTTCATTAACGGCTTTGCTCATAAAAATGTTA  
 TGACGTTTTGCCCGCAGGCGGGAACCATCCACTTCACGAGACTGATCTCCTCTGCCGGA  
 ACACCGGGCATCTCCAACCTATAAGTTGGAGAAATAAGAGAATTTGAGATTGAGAGAATG  
 AAAAAAAAACC

CTTAGTTCATAGGTCCATTTCTCTTAGCGCAACTACAGAGAACAGGGGCACAAACAGG  
 CAAAAACGGGACACAACCTCAATGGAGTGATGCAACCTGCCTGGAGTAAATGATGACACAA  
 GGCAATTGACCCACGCATGTATCTATCTCATTTTTCTTACACCTTCTATTACCTTCTGCTCT  
 CTCTGATTTGGAAAAAGCTGAAAAAAAGGTTGAAACCAGTTCCTGAAATTATTCCCCTAC  
 TTGACTAATAAGTATATAAAGACGGTAGGTATTGATTGTAATTCTGTAAATCTATTCTTAAA  
 CTTCTTAAATTTACTTTTATAGTTAGTCTTTTTTTTAGTTTTTAAACACCAAGAACTTAG  
 TTTGGAATAAACACACATAAACAAAC

cctaggacttctaagcgccgc

TTTGTTCCTTCTGACTTTTAGCTCGTACAAAATACAATATACTTTTCATTTCT

CCGTAAACAACATGTTTTCCCATGTAATATCCTTTTCTATTTTTTCGTTCCGTTACCAACT

TTACACATACTTTATATAGCTATTCACTTCTATACACTAAAAAACTAAGACAATTTTAAT

TTTGCTGCCTGCCATATTTCAATTTGTTATAAATTCCTATAATTTATCCTATTAGTAGCT

AAAAAAAGATGAATGTGAATCGAATCCTAAGA

**FIG. 3A**

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GCTAGCGCTATATGCGTTGATGCAATTTCTATGCGCACCCGTTCTCGGAGCACTGTCC  
GACCGCTTTGGCCGCGCCAGTCCTGCTCGCTTCGCTACTTGGAGCCACTATCGACTACGCGATCATGGCGACC  
ACACCCGCTCTGTGGATCAGATCCAATTTCTCTTAGGATTCGATTACATTTCATCTTTTTTTAGCTACTAATAGGA  
TAAATTATAGGAATTTATAACAAATTGAAATATGGCAGGCAGCAAAATTAATTTGTCTTAGTTTTTTAGTGTAT  
AGAAGTGAATAGCTATATAAAGTATGTGTAAAGTTGGTAACGGAACGAAAAATAGAAAAGGATATTACATGGGAA  
AACATGTTGTTTACGGAGAAATGAAAAGTATATTGTATTTTGTACGAGCTAAAAGTACAGTGGGAACAAA

GTCCGACTTTCACAGGCAACGCGTGTGTTGTTTATGTGTGTTTATTCGAAACTAAGTTCTTG  
GTGTTTTAAACTAAAAAAGACTAAGTATAAAGTAGAATTTAAGAAGTTTAAGAAAT  
AGATTTACAGAATTACAATCAATACCTACCGTCTTTATATACTTATTAGTCAAGTAGGGG  
AATAATTTTCAGGGAAGTGGTTTCAACCTTTTTTTTTCAGCTTTTTCCAAATCAGAGAGAGC  
AGAAGGTAATAGAAGGTGTAAGAAAATGAGATAGATACATGCGTGGGTCAATTGCCTTGT  
GTCATCATTTACTCCAGGCAGGTTGCATCACTCCATTGAGGTTGTGCCCGTTTTTTGCOCT  
GTTTGTGCCCGTGTCTCTGTAGTTGCGCTAAGAGAATGGACCTATGAACTAAGGGTTTTT  
TTTTTTTCATTCTCTCAATCTGAAATTTCTCTTATTTCTCCAACCTATAAGTTGGAGATGC  
CCGGTGTTCGCGCAGAGGAGATCAGTCTCGTGAAGTGGATGGTTTTCCCGCCTGCGGGCAA  
AACGTACATAACATTTTTTATGAGCGAAGCCGTTAATGAAGACAAAAATCCCTTAATTAATAA  
CATTAGAAATGGTGATTAGAAAAGGCAGGATTAATCAGTTACACAGGCTGTAACCGGAGAGA  
CGGATCATAAGGCAATTTTTAGATAAGACTGGTTAGAGTTCTTGGCATCAGAAAAATTTGA  
GAAACGATTTTTCCGTTTGTGTTGCCCTACGTTTTGCCCTTTGATCAAACTATCAGTTA  
AGATATTAATTTTTTTGAGAAAACGATTCTTTGATTAGTCTCTTCAACCAAACAATGAGC  
TCTGAAGACGAATTGGGAAGTATCGGTACTGTGTTTCCCGGAAGTCCCATAGATAAGAGC  
ATTGGGAGTATTCTCCACAATTTGATGAAGAAGTGGAGACTTTGCTGGAAGATAGCTTC  
ACGTGGAACATTCTGACTGGAACGAGTTAACAACCCGAAATACAATTCGCCAGGTTT  
AGAATTTGGTGATTTTGAATGGGACATTCTATTATTCCTCAGGGAACCATATAAAGGT  
GTTGCGGTATATCTGGAACCTCATCCGGAAGAAAAATTAGATGAGACTACGGGAGAGATG  
GTGCCAGTTGATCCGGACTGGTATTGTTGTGCTCAGTTTGGCATTGGTATATCTAGACCT  
GGTAATGGTGACACCATCAATTTAATTAACAAATCGCATCACCGATTCAACGCTCTAGAT  
ACAGACTGGGGATTTGCAAAATTTGATAGATTTGAACAACCTGAAACATCCCTCAAAAGGA  
AGACCGCTTTCGTTCTTAAACGAAGGGACCTTGAACATAACAGCGTATGTGCCCATATTG  
AA

GGATCCTCGACCGAT  
GCCCTTGAGAGCCTTCAACCCAGTCAGCTCCTTCCGGTGGGCGCGGGGCATGACTATCGTCCGCGCACTTATGAC  
TGCTCTCTTTATCATGCAACTCGTAGGACAGGTGCCGGCAGCGCTCTGGGTCATTTTTCGGCGAGGACCGCTTTTCG  
CTGGAGCGCGACATGATCGGCCTGTGCGTTGCGGTATTGCGAATCTTGACACGCCCTCGCTCAAGCCTTCGTAC  
TGGTCCCGCCACCAACGTTTTCGGCGAGAAGCAGGCCATTATCGCCGGCATGGCGGCCGACGCGCTGGGCTACGT  
CTTGCTGGCGTTTCGCGACGCGAGGCTGGATGGCCTTCCCCATTATGATTCTTCTCGCTTCCGGCGGCATCGGGAT  
GCCCGCGTTGCAGGCCATGCTGTCCAGGCAGGTAGATGACGACCATCAGGGACAGCTTCAAGGATCGCTCGCGGC  
TCTTACCAGCCTAATTCGATCATTGGACCGCTGATCGTCACGGCGATTATGCCCGCTCGGCGAGCACATGGAA  
CGGGTTGGCATGGATTGTAGGCGCCGCCCTATACCTTGTCTGCCCTCCCCGCGTTGCGTCCGGTGCATGGAGCCG  
GGCCACCTCGACCTGAATGGAAGCCGCGCGCACCTCGCTAACGGATTCAACACTCCAAGAATTGGAGCCAATCAA  
TTCTTGGGAGAACTGTGAATGCGCAAAACCAACCTTGGCGAACAATATCCATCGCGTCCGCCATCTCCAGCAGC  
CGACGCGCGCGCATCTCGGGCAGCGTTGGGTCTTGGCCACGGGTGCGCATGATCGTGCTCCTGTGCTTGAGGACC  
CGGCTAGGCTGGCGGGGTTGCCCTTACTGGTTAGCAGAATGAATCACCGATACGCGAGCGAACGTGAAGCGACTGC  
TGCTGCAAAACGCTTGCGACCTGAGCAACAACATGAATGGTCTTCGGTTTTCCGTGTTTCGTAAAGTCTGGAAACG  
CGGAAGTCAGCGCCCTGCACCATTATGTTCCGGATCTGCATCGCAGGATGCTGCTGGCTACCCCTGTGGAACACCT  
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TTCATCGGTATCATTACCCCCATGAACAGAAATCCCCCTTACACGGAGGCATCAGTGACCAAAACAGGAAAAAAC  
GCCCTTAACATGGCCCCGCTTTATCAGAAGCCAGACATTAAACGCTTCTGGAGAAACTCAACGAGCTGGACCGGAT  
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GATGACGGTGAAAACCTCTGACACATGCAGCTCCCGGAGACGGTCACAGCTTGTCTGTAAGCGGATGCCGGGAGC  
AGACAAGCCCGTCAGGGCGCGTCAGCGGGTGTGGCGGGTGTCCGGGCGCAGCCATGACCCAGTCACGTAGCGAT  
AGCGGAGTGATACTGGCTTAACTATGCGGCATCAGAGCAGATTGTACTGAGAGTGACCATATGCGGTGTGAAA  
TACCGCACAGATGCGTAAGGAGAAAAATACCGCATCAGGCGCTCTTCCGCTTCTCTCGCTCACTGACTCGCTGCGCT  
CGGTCGTTCCGGTGCGGCGAGCGGTATCAGCTCACTCAAAGGCGGTAAATACGGTTATCCACAGAATCAGGGGATA  
ACGCAGGAAAGAACATGTGAGCAAAAGGCCAGCAAAAGGCCAGGAACCGTAAAAAGGCCCGCTTGTGGCGTTTTT  
TCCATAGGCTCCGCCCCCTGACGAGCATCACAAAAATCGACGCTCAAGTCAGAGGTGGCGAAACCCGACAGGAC  
TATAAGATACACAGGCGTTTTCCCCCTGGAAGCTCCCTCGTGCGCTCTCTGTTCCGACCCCTGCCGCTTACCGGAT  
ACCTGTCCGCTTTCTCCCTTCCGGAAGCGTGGCGCTTTCTCATAGCTCACGCTGTAGGTATCTCAGTTCCGGTGT  
AGGTGCTTCCGCTCAAGCTGGGCTGTGTGCACGAACCCCCGTTCCAGCCGACCGCTGCGCCTTATCCGGTAACCT  
ATCGTCTTGAGTCCAACCCGGTAAGACACGACTTATCGCCACTGGCAGCAGCCACTGGTAACAGGATTAGCAGAG

**FIG. 3B**

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CGAGGTATGTAGGCGGTGCTACAGAGTTCTTGAAGTGGTGGCCTAACTACGGCTACACTAGAAGGACAGTATTTG  
GTATCTGCGCTCTGCTGAAGCCAGTTACCTTCGGAAAAAGAGTTGGTAGCTCTTGATCCGGCAAAACAAACCCCG  
CTGGTAGCGGTGGTTTTTTTTGTTTTGCAAGCAGCAGATTACGCGCAGAAAAAAGGATCTCAAGAAGATCCTTTGA  
TCTTTTCTACGGGCTCTGACGCTCAGTGGAAACGAAAACTCACGTTAAGGGATTTTGGTCATGAGATTATCAAAAA  
GGATCTTCACCTAGATCCTTTTAAATTAAAAATGAAGTTTTAAATCAATCTAAAGTATATATGAGTAAACTTGGT  
CTGACAGTTACCAATGCTTAATCAGTGAGGCACCTATCTCAGCGATCTGTCTATTTTCGTTTCATCCATAGTTGCCCT  
GACTCCCCGTCGTGTAGATAACTACGATACGGGAGGGCTTACCATCTGGCCCCAGTGCTGCAATGATACCGCGAG  
ACCCACGCTCACCAGCTCCAGATTTATCAGCAATAAACAGCCAGCCGGAAGGGCCGAGCGCAGAACTGGTCCTG  
CACTTTTATCCGCTCCATCCAGTCTATTAATTGTTGCCGGGAAGCTAGAGTAAGTACTTCGCCAGTTAATAGTT  
TGCGCAACGTTTGTGTCATTGCTGCAGGCATCGTGGTGTACGCTCGTCTGTTGGTATGGCTTCATTCAGCTCCG  
GTTCCCAACGATCAAGGCGAGTTACATGATCCCCATGTTGTGCAAAAAAGCGGTTAGCTCCTTCGGTCTCCGA  
TCGTTGTCAGAAAGTAAGTTGGCCGAGTGTATCACTCATGTTATGGCAGCACTGCATAATTCTCTTACTGTCA  
TGCCATCCGTAAGATGCTTTTCTGTGACTGGTGAGTACTCAACCAAGTCATTCGTAGAATAGTGTATGCGGCGAC  
CGAGTTGCTCTTGCCCGGCGTCAACACGGGATAATACCGCGCCACATAGCAGAACCTTAAAGTGCTCATCATTG  
GAAAACGTTCTTCGGGGCGAAACTCTCAAGGATCTTACCGCTGTTGAGATCCAGTTCCGATGTAACCCACTCGTG  
CACCCAACGATCTTCAGCATCTTTTACTTTTACCAGCGTTTCTGGGTGAGCAAAAAACAGGAAGGCAAAATGCCG  
CAAAAAAGGGAATAAGGGCGACACGGAAATGTTGAATACTCATACTCTTCTTTTCAATATTATTGAAGCATT  
ATCAGGGTTATTGTCTCATGAGCGGATACATATTTGAATGTATTTAGAAAAATAACAAATAGGGGTTCCGCGCA  
CATTTCCCGAAAAAGTGCCACCTGACGTCTAAGAAACCATTTATTATCATGACATTAACCTATAAAAAATAGGCGTA  
TCACGAGGCCCCCTTCGTCTTCAAGAATTCTCATGTTTGACAGCTTATCATCGATCCACTTGTATATTTGGATGAA  
TTTTTGAGGAATTCTGAACAGTCTTAAACGAGTAAATAGGACCGGCAATCTTCAAGCAATAAACAGGAATAC  
CAATTATTAAAGATAACTTAGTCAGATCGTACAATAAGCTTTGAAGAAAAATGCGCCTTATTCAATCTTTGCT  
ATAAAAAATGGCCCAAAATCTCACATTGGAAGACATTTGATGACCTCATTTCTTCAATGAAGGCGCTAACGGAG  
TTGACTAATGTTGTGGGAAATTGGAGCGATAAGCGTGCTTCTGCGGTGGCCAGGCAACGTATACCTCATCAGATA  
ACAGCAATACCTGATCACTACTTCGCACTAGTTTCTCGGTACTATGCATATGATCCAATATCAAAGGAAATGATA  
GCATTGAAGGATGAGACTAATCCAATTGAGGAGTGGCAGCATATAGAACAGCTAAAGGGTAGTGCTGAAGGAAGC  
ATACGATACCCCGCATGGAATGGGATAATATCACAGGAGGTACTAGACTACCTTTTCTCTACATAAAATAGACGC  
ATATAAGTACGCATTTAAGCATAAACACGCACTATGCCGTTCTTCTCATGTATATATATACAGGCAACACGCA  
GATATAGGTGCGACGTGAACAGTGAGCTGTATGTGCGCAGCTCGCGTTGCATTTTCGGAAGCGCTCGTTTTCCGA  
AACGCTTTGAAGTCTCTATTCCGAAGTCTCTATTCTCTAGAAAGTATAGGAACTTCAGAGCGCTTTTGAAACCA  
AAAGCGCTCTGAAGACGCACTTTCAAAAAACCAAAAAACGCACCGGACTGTAACGAGCTACTAAAAATATTGCGAAT  
ACCGCTTCCACAAACATTGCTCAAAAGTATCTCTTGTCTATATATCTCTGTGCTATATCCCTATATAACCTACCC  
ATCCACCTTTTCGCTCCTTGAACCTGCATCTAACTCGACCTCATACATTTTATGTTTATCTCTAGTATTACTCT  
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TTTCTGTTTCAAAAGTATGCGCAATCCACATCGGTATAGAATATAATCGGGGATGCCTTTATCTTGAAAAAATGC  
ACCCGCAGCTTCGCTAGTAATCAGTAAACGCGGGAAGTGGAGTCAAGCTTTTATGGAAGAGAAAAATAGACAC  
CAAAGTAGCCTTCTTCTAACCTTAACGGACCTACAGTGCAAAAAAGTTATCAAGAGACTGCATTATAGAGCGCACA  
AAGGAGAAAAAAGTAATCTAAGATGCTTTGTTAGAAAAATAGCGCTCTCGGGATGCAATTTTGTAGAACAAAA  
AGAAGTATAGATTCTTTGTTGGTAAAAATAGCGCTC  
TCGCGTTGCATTTCTGTTCTGTAAAAATGCAGCTCAGATTCTTTGTTTGAAAAATTAGCGCTCTCGCGTT  
GCATTTTGTGTTTTACAAAAATGAAGCACAGATTCTTCGTTGGTAAAAATAGCGCTT  
CTCGGTTGCATTTCTGTTCTGTAAAAATGCAGCTCAGATT  
CTTTGTTTGAAAAATTAGCGCTCTCGCGTTGCATTTTGTGTTTCTACAAAAATGAAGCACAGATGCTTCGTTAACAAA  
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TACGATCGTAAATCTGTAAAAACAGTTTGTGCGATATTAGGCTGTATCTCCTCAAAGCGTATTCCAATATCATTGA  
GAAGCTGCATTTTTTTTTTTTTTTTTTTTTTTTTTTTTTATATATATTTCAAGGATATACCATTTGTAATGTCTG  
CCCCTAAGAAGATCGTCGTTTTGCCAGGTGACCACGTTGGTCAAGAAATCACAGCCGAAGCCATTAAAGGTTCTTA  
AAGCTATTTCTGATGTTTCGTTCCAATGTCAAGTTCGATTTCGAAAAATCATTTAATTGGTGGTGCTGCTATCGATG  
CTACAGGTGTTCCACTTCCAGATGAGGCGCTGGAAGCCTCCAAAGAGGCTGATGCCGTTTTGTTAGGTGCTGTGG  
GTGGTCTTAATGGGGTACCGGTAGTGTTAGACCTGAACAAGGTTTACTAAAAATCCGTAAAGAACTTCAATTGT  
ACGCCAACTTAAGACCATGTAACCTTGCATCCGACTCTCTTTAGACTTATCTCCAATCAAGCCACAATTTGCT  
AAAGGTACTGACTTCGTTGTTGTTAGAGAATTAGTGGGAGGTATTTACTTTGGTAAGAGAAAGGAAGACGATGGT  
GATGGTGTCTGCTTGGGATAGTGAACAATACACCGTTCCAGAAGTGCAAGAAATCACAAGAAATGGCCGCTTTCATG  
GCCCTACAACATGAGCCACCATTGCCTATTTGGTCTTGGATAAAGCTAATGTTTTGGCCTCTTCAAGATTATGG  
AGAAAACTGTGGAGGAAACCATCAAGAACGAATTCCTACATTGAAAGTTCAACATCAATTGATTGATTCTGCC  
GCCATGATCCTAGTTAAGAACCCAAACCCACCTAAATGGTATTATAATCACCAGCAACATGTTTGGTGATATCATC  
TCCGATGAAGCCTCCGTTATCCAGGCTCCTTGGGTTTGTGTCATCTGCGTCTTGGCCTCTTGGCCAGCAAG  
AACACCGCATTTGGTTTGTACGAACCATGGTTCGGCTCCAGATTGCCAAAGAAATAGGTCAACCCATC  
GCCATCTTTGCTGCTGCAATGATGTTGAAATGTTCATTGAACTTGCTGAAGAAGGTAAAGCCATTGAAGAT  
GCAGTTAAAAAGGTTTTGGATGCAGGTATCAGAACTGGTGATTAGGTGGTTCCAACAGTACCACCGAAGTCGGT

**FIG. 3C**

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GATGCTGTGCGCCGAAGAAGTTAAGAAAATCCTTGCTTAAAAAGATTCTCTTTTTTATGATATTTGTACCAAAAA  
 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAATGCAGCGTCACATCGGATAATAATGAT  
 GGCAGCCATTGTAGAAGTGCCTTTTGCACTTCTAGTCTCTTTCTCGGTCTAGCTAGTTTTACTACATCGCGAAGA  
 TAGAATCTTAGATCACACTGCCTTTGCTGAGCTGGATCAATAGAGTAACAAAAGAGTGGTAAGGCCTCGTTAAAG  
 GACAAGGACCTGAGCGGAAGTGTATCGTACAGTAGACGGAGTATACTAGTATAGTCTATAGTCCGTGGAATTCTA  
 AGTGCCAGCTTTATAATGTCACTTCTCCTTACTACAGACCCGCTGAAAGTAGACACATCATCATCAGTAAGCTTT  
 GACAAAAGCATTGAGTAGCTAACTCTTCTATGCAATCTATAGCTGTTTTATAAGGCATTCAATGGACAGATTGA  
 GGTTTTTGAAACATACTAGTGAAATTAGCCTTAATCCCTTCTCGAAGTTAATCATGCATTATGGTGTAAAAATG  
 CAACTCGCGTTGCTCTACTTTTTCCCGAATTTCCAAATACGCAGCTGGGGTGATTGCTCGATTTCTGAACGAAAG  
 TTTTGTATATAAAAAACCGCGAAAACCTTCTGTAACAGATAGATTTTACAGCGCTGATATACAATGACATCAGCT  
 GTAATGGAAAATAACTGAAATATGAATGGCGAGAGACTGCTTGCTTGTATTAGCAATGTATTATGCAGCACTTC  
 CAACCTATGGTGTACGATGAAAGTAGGTGTGTAATCGAGACGACAAGGGGGACTTTTCCAGTTCCTGACAATTAT  
 AAGAAATACAAAACGTTAGCATTTCGATTGTTTGGACATGTACTGAATACAGACGACACACCGGTAAATTGAAAAA  
 GAAGTGGATTGGCCTGATCCTGCCTAGTGTACAATAACAATGTCTGATCGAATCATAAATCAGCCGAAATTATCA  
 CAGTTTATATCGGTTGCATTTATTAGTCAGTTAAAGGCCACCATCGGAGAGGGTTTAGATATTAATGTAAGGCG  
 ACGCTAAACCGCAGGGGAAAGGGTATCAGAAGGCCTAAAGGCGTATTTTTTAGATACATGGAATCTCCATTTGTC  
 AATACAAAGGTCAGTGCATTTCTTCTTATCTTCGAGATTATAATAAAATTCGCTCAGAATATCACAATAATACT  
 AAATTCATTCTCAGTTTTTCATGTCAAGCATATTGGGCATCTGGCCCAAACCTTCTCCGCTTGAAGAATGTTATT  
 AGGTGCTCCATAATTTCATGAATACATTTCTAAGTTTGTGGAAAGAGAACAGGATAAAGGTCATATAGGAGATC  
 AGGAGCTACCGCCTGAAGAGGACCCCTTCTCGTGAACATAACAATGTACAACATGAAGTCAATAGTTTAAACGGAAC  
 AAGATGCGGAGGCGGATGAAGGATTGTGGGGTGAAATAGATTCAATTATGTGAAAATGGCAGTCTGAAGCGGAAG  
 ATCAAACCTGAGGCGGAGATAATAGCCGACAGGATAATTGGAAATAGCCAGAGGATGGCGAACCTCAAATTCGTC  
 GTACAAAGTTCAAAGTGTCTTGTATCATATACTAAAGGAACATAATCAATCTCAGGGAACCGTAAAGGTTTATC  
 GCGGTAGTAGTTTTTACACGATTGATAAAGATAAGCTTACATTATGAAGAGCAGCATATTACAGCCGTATGGG  
 TCTACTTGACAGTAAATTTGAAGAGCATTGGAAGCCTGTTGATGTAGAGGTGAGTTTAGATGCAAGTTCAAGG  
 AGCGAAAGGTGGATGGGTAGGTTATATAGGGATATAGCACAGAGATATATAGCAAAGAGATACTTTTGAGCAATG  
 TTTGTGGAAGCGTATTTCGCAATATTTTAGTAGCTCGTTACAGTCCGGTGGCTTTTTTGGTTTTTTGAAAGTGGCT  
 CTTCAGAGCGCTTTTGGTTTTCAAAGCGCTCTGAAGTTCTTATACTTTCTAGAGAATAGGAACCTTCGGAATAGG  
 AACTTCAAAGCGTTTCCGAAAACGAGCGCTTCCGAAAATGCAACGCGAGCTGCGCACATACAGCTCACTGTTTAC  
 GTCGACCTATATCTGCGTGTGGCTGTATATATATATACATGAGAAGAACGGCATAGTGCGTGTATTATGCTTAA  
 ATGCGTACTTATATGCGTCTATTTATGTAGGATGAAAGGTAGTCTAGTACCTCCTGTGATATTATCCCATTCAT  
 CCGGGTATCGTATGCTTCTTCCAGCACTACCTTTAGCTGTTCTATATGCTGCCACTCCTCAATTGGATTAGTC  
 TCATCCTTCAATGCTATCATTTCCTTTGATATTGGATCATACCTAGAAGTATTACGTGATTTTCTGCCCCCTTAC  
 CCTCGTTGCTACTCTCCTTTTTCGTGGGAACCGCTTTAGGGCCCTCAGTGATGGTGTGTTTCTGAATTTATATG  
 TCTCTTGCTATTTGTGCTCTACTTCTTGTTCGCTGGAGGGAACCTTCTTCAATTGTATTAGCATGGTTCACTTC  
 AGTCTTCTCTTCCAACCTCACTCTTTTTTTGCTGTAAACGATTCTCTGCCGCCAGTTCTATTGAACTATTGAATAT  
 ATCCTTTAGAGATTCCGGGATGAATAAATCACCTATTAAAGCAGCTTGACGATCTGGTGGAACTAAAGTAAGCAA  
 TTGGGTAAACGACGCTTACGAGCTTCATAACATCTTCTTCCGTTGGAGCTGGTGGGACTAATAACTGTGTACAATC  
 CATTTTTCTCATGAGCATTTCGGTAGCTCTCTTCTTGTCTTTCTCGGGCAATCTTCTTATTATTATAGCAATAGA  
 TTTGTATAGTTGCTTTCTATTGTCTAACAGCTTGTTATTCTGTAGCATCAAATCTATGGCAGCCTGACTTGCTTC  
 TTGTGAAGAGAGCATACCATTTCCAATCGAATCAAACCTTTCTTAAACCATCTTCGCAGCAGGCAAAATTACCTC  
 AGCACTGGAGTCAGAAGATACGCTGGAATCTTCTGCGCTAGAATCAAGACCATAACGCCCTACCGGTTGTGAGAGA  
 TTCCATGGGCTTATGACATATCCTGGAAAGAGTAGCTCATCAGACTTACGTTTACTCTCTATATCAATATCTAC  
 ATCAGGAGCAATCATTTCAATAAACAGCCGACATACATCCAGACGCTATAAGCTGTACGTGCTTTTACCGTCAG  
 ATTCTTGGCTGTTTCAATGTCTGTCATTTTGGTTTTCTTTTACCAGTATTGTTTCTGTTGATAATGTATTCTTGCT  
 TATTACATTATAAAATCTGTGTCAGATCACATGTCAAACAACCTTTTATCACAAAGATAGTACCGCAAAACGAACC  
 TGCGGGCCGTCTAAAAATTAAGGAAAAGCAGCAAAGGTGCATTTTTTAAATATGAAATGAAGATACCGCAGTACC  
 AATTATTTTCGCACTACAAATAATGCGCGGCCGGTGCATTTTTTCGAAAGAACGCGAGACAAAACAGGACAATTAAA  
 GTTAGTTTTTCGAGTTAGCGTGTGTAATACTGCAAGATACAAGATAAATAGAGTAGTTGAAACTAGATATCAAT  
 TGCACACAAGATCGGCGCTAAGCATGCCACAATTTGATATATTATGTAAAACACCACCTAAGGTGCTTGTTCGTC  
 AGTTTGTGGAAGGTTTGAAGACCTTCAGGTGAGAAAATAGCATTATGTGCTGCTGAACTAACCTATTTATGTT  
 GGATGATTACACATAACGGAACAGCAATCAAGAGAGCCACATTTCATGAGCTATAATACTATCATAAGCAATTCCG  
 TGAGTTTCGATATTGTCAATAAATCACTCCAGTTTAAATACAAGACGCAAAAAGCAACAATCTGGAAGCCTCAT  
 TAAAGAAATTGATTCTGCTTGGGAATTTACAATTATCTTACTATGGACAAAACATCAATCTGATATCACTG  
 ATATTGTAAGTAGTTTGAATTACAGTTTGAATCATCGGAAGAAGCAGATAAGGGAAATAGCCACATGAAAAAA  
 TGCTTAAAGCACTTCTAAGTGAGGGTGAAGCATCTGGGAGACTGAGAGAAATACTAAATTCGTTTGAAGTATA  
 CTTTCGAGATTTACAAAACAAAACTTTATACCAATTCCTCTTCTAGCTACTTTTCATCAATTTGGAAGATTCA  
 GCGATATTAAGAACGTTGATCCGAAATCATTTAAATTAGTCCAAATAAGTATCTGGGAGTAATAATCCAGTGTT  
 TAGTGACAGAGACAAAGACAAGCGTTAGTAGGCACATATACTTCTTTAGCGCAAGGGGTAGGTAGC

**FIG. 3D**

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ParvoB19.Opti.VP1

acgcgtacaaacaaaATGTCTAAGAAAATCTGGTAAATCGTGGAAATCTGATGATAAAATTTGCTAAGGCTGTTTACC  
AACAAATTTGTTGAATTTTACGAAAAAGTTACTGCTACTGATTTGGAAATTTGATTCAAAATTTTGAAGGATCATTAACAAC  
ATTTCTTTGGATAATCCATTTGGAAAAATCCATCTTCAATTTGTTGATTTGGTTGCTAGAAATTAAGAACAACTTGAAGAA  
CTCTCCAGATTTGTATTCCTCATCATTTCCAAATCFCATGGTCAAAATTTGCTGATCATCCACATGCTTTATCTTCATCTTT  
CATCTCATGCTGAACCAAGAGGTGAAAAATGCTGTTTTATCTTCTGAAGATTTGCATAAACCCAGGTCAAGTTTCTGTT  
CAATTGCCAGGTACTAATTTACGTTTGTCACAGGTAAATGAATGCAAGCTGGTCCACCAACAATCTGCTGTTGATTTCTGCTG  
TGCTAGAAATTCATGATTTTCAGATACTCTCAATTGGCTAAGTTGGGTATTAATCCATATACCTCATTTGGACTGTTGCTG  
ATGAAGAAATTTGTTGAAGAACATTAAGAAATGAAACTGGTTTTCAAGCTCAAGTTGTTAAAGATTACTTCACTTTTGAAA  
GGTGCTGCTCCAGTTGCTCATTTTCAAGGTTCTTTGCCAGAAAGTTCCAGCTTATAACGCTTCTGAAAAATATCC  
ATCTATGACATCTGTTAATTTCTGCTGAAGCATCTACTGGTGCAGGTGGAGGTGTTCTAAATTTCTGTTAAATCTATGT  
GGTCTGAAGGTCTACTTTTTCTGCTGTAATTCAGTTACTTTCTCTAGACAAATTCCTTGATTTCCATATGATCCCA  
GAACATCATTAACAAGTTTTTTCACCAGCTGCTTCATCTTGTCAATAATGCTTCAGGTAAAGAACTAAGGTTTGTAC  
TAATTTCTCCAATTAATGGGTTATTTCTACTCTCTGGAGATACTTTGGAATTTTAATGCTTTGAACCTTGTTTTTCTCCAT  
TGAAATTTTCAACATTTTGATTGAAAACTACGGTTCTATTGCTCCAGATGCTTTGACTGTTACTATTTCTGAAATTTGCT  
GTTAAGGATGTTACTGATAAAACAGGTGCTGTTCAAGTTACTGATTTCTACTACTGTTAGATTTGTCATGTTGGT  
TGATCATGAATACAAATACCCATACGTTTGGGTCAGGTCAAGATACCTTTGGCTCCAGAAATTTGCCAATTTGGGTTT  
ATTTTCCACCACAATCTGCTTTTACGTTTGGAAACATTTCTTTTCAATTTGTTGGTACTGTTGGTACTGCTTCTAT  
GCTTCTGAAGAAATCTGCTTTTACGTTTGGAAACATTTCTTTTCAATTTGTTGGTACTGTTGGTACTGCTTCTAT  
GTCCTACAAATTTCCACCAGTTCCACCTGAAAATTTGGGAAGTTGTTCTCAACATTTTACGAAATGTACAAATCCAT  
TGTAATGGTTCTAGATTTGGGTGTTCCAGATACTTTGGGTGGTGATCCAAAATTTAGATCTTTGACTCATGAAGATCAT  
GCTATTCAACCCACAATAATTCATGCCAGGTCCATTTGGTTAAATTCGTTTCTACTAAAGAAAGTGATTTCTCTAATAC  
AGGTGCTGTTAAAGCATTGACTGGTTTGTCTACTGGTACTTCTCAAAACACTAGAAATTTCTTTAAGACCCAGGTCCAG  
TTTCACAACCATATCATCATTTGGGATACCTGATAAGTACGTTACTGGTATTAATGCTATTTCAATGTTCAAACTACT  
TATGGTAATGCTGAAGATAAAGAAATATCAACAAGGTGTTGGTAGATTTCCAAACGAAAAAGAACAAATTTGAAACAAT  
GCAAGGTTTGAATATGCATACTTACTTTCCAAACAAAGGTACTCAACAATACACTGATCAAAATTTGAAAGACCATTGA  
TGCTTGGTTCTGTTTGGAAATAGAAGAGCTTTGCTATTAATGAACTCAATTTGTTGGTCTAAGATTTCCAAATTTAGATGAT  
TCTTTCAAGACTCAATTTGCTGCTTTTGGGTGGTTGGATCACTCAACCTCCACCACAAATTTTCTTTGAAGATTTT  
GCCACAATCTGGTCCAAATTTGGTGGTATTAATCTATGGGTATTACTACTTTTGGTTCAATATGCTGTTGGTATTAATGA  
CTGTTACAATGACTTTTAAGTTGGGTCCAAGAAAAGCTACAGGTAGATGGAATCCACAACCCAGGTGTTTATCCACCA  
CATGCTGCTGGTCAATTTGCCCTTACGTTTGTATGATCCAACTGCTACTGATGCTAAACAACATCATAGACATGGTTA  
TGAAAAACCTGAAGAATTTGTGGACTGCTAAATCTAGAGTTTCATCCATTTGTAATGAGTcgac

**FIG. 4**

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**ParvoB19.Opti.VP2**

cctaggacaaaacaaaATGACATCTGTTAATTCTGCTGAAGCATCTACTGGTGCAGGTGGAGGTGGTTCTAATTCTG  
TTAAATCTATGTGGTCTGAAGGTGCTACTTTTTCTGCTAATTCAGTTACTTGTACTTTCTCTAGACAATTCTTGATT  
CCATATGATCCAGAACATCATTACAAAGTTTTTTCACCAGCTGCTTCATCTTGTCTATAATGCTTCAGGTAAAGAAGC  
TAAGGTTTGTACTATTTCTCCAATTATGGGTTATTCTACTCCTTGGAGATACTTGGATTTTAATGCTTTGAACCTGT  
TTTTTCTCCATTGGAATTTCAACATTTGATTGAAAACCTACGGTTCTATTGCTCCAGATGCTTTGACTGTTACTATT  
TCTGAAATTGCTGTTAAGGATGTTACTGATAAAACAGGTGGTGGTGTTCAGTTACTGATTCTACTACTGGTAGATT  
GTGCATGTTGGTTGATCATGAATACAAATACCCATACGTTTTGGGTCAAGGTCAAGATACTTTGGCTCCAGAATTGC  
CAATTTGGGTTTATTTCCACCACAATACGCTTATTTGACTGTTGGTGATGTTAATACTCAAGGTATTTCTGGTGAT  
TCTAAAAGTTGGCTTCTGAAGAATCTGCTTTTTACGTTTTGGAACATTCTTCTTTTCAATTGTTGGGTACTGGTGG  
TACTGCTTCTATGTCTTACAAATTTCCACCAGTTCCACCTGAAAATTTGGAAGGTTGTTCTCAACATTTTACGAAA  
TGTACAATCCATTGTATGGTTCTAGATTGGGTGTTCCAGATACTTTGGGTGGTGATCCAAAATTTAGATCTTTGACT  
CATGAAGATCATGCTATTCAACCACAAAATTTTCATGCCAGGTCCATTGGTTAATTCTGTTTCTACTAAAGAAGGTGA  
TTCTTCTAATACAGGTGCTGGTAAAGCATTGACTGGTTTGTCTACTGGTACTTCTCAAAACACTAGAATTTCTTTAA  
GACCAGGTCCAGTTTCCAAACCATATCATCATTGGGATACTGATAAGTACGTTACTGGTATTAATGCTATTTCCACAT  
GGTCAAACCTACTTATGGTAATGCTGAAGATAAAGAATATCAACAAGGTGTTGGTAGATTTCCAAACGAAAAAGAACA  
ATTGAAACAATTGCAAGGTTTGAATATGCATACTTACTTTCCAAACAAAGGTACTCAACAATACACTGATCAAATTG  
AAAGACCATTGATGGTTGGTTCTGTTTGGAAATAGAAGAGCTTTGCATTATGAATCTCAATTGTGGTCTAAGATTCCA  
AATTTAGATGATTCTTTCAAGACTCAATTTGCTGCTTTGGGTGGTTGGGGTTTGCATCAACCTCCACCACAAATTTT  
CTTGAAGATTTTGGCACAATCTGGTCCAATTGGTGGTATTAAATCTATGGGTATTACTACTTTGGTTCAATATGCTG  
TTGGTATTATGACTGTTACAATGACTTTTAAGTTGGGTCCAAGAAAAGCTACAGGTAGATGGAATCCACAACCAGGT  
GTTTATCCACCACATGCTGCTGGTCAATTTGCCTTACGTTTGTATGATCCAACCTGCTACTGATGCTAAACAACATCA  
TAGACATGGTTATGAAAAACCTGAAGAATTGTGGACTGCTAAATCTAGAGTTCATCCATTGTAATGAgcgccgc

**FIG. 5**

**ADH2 / GAPDH promoter (as BamHI / HindIII fragment)**  
**used in the expression of ParvoB19 VP2**

GGATCCTTCAATATGCGCACATACGCTGTTATGTTCAAGGTCCCTTCGTTTAAGAACGAAAGCGGTCTTCCTTTTGA  
GGGATGTTTCAAGTTGTTCAAATCTATCAAAATTTGCAAAATCCCCAGTCTGTATCTAGAGCGTTGAATCGGTGATGCC  
ATTTGTTAATTAAATTGATGGTGTCAACATTACCAGGTCTAGATATACCAATGGCAAACCTGAGCACAACAATACCAG  
TCCGGATCAACTGGCACCATCTCTCCCGTAGTCTCATCTAATTTTTCTTCCGGATGAGGTTCCAGATATACCGCAAC  
ACCTTTATTATGGTTTCCCTGAGGGAATAATAGAATGTCCCATTCGAAATCACCAATTCTAAACCTGGGCGAATTGT  
ATTTCCGGGTTTGTAACTCGTTCCAGTCAGGAATGTTCCACGTGAAGCTATCTCCAGCAAAGTCTCCACTTCTTCA  
TCAAATTTGTGGAGAATACTCCCAATGCTCTATCTATGGGACTTCCGGGAAACACAGTACCGATACTTCCCAATTC  
GTCTTCAGAGCTCATGTTTGTGTTGAAGAGACTAATCAAAGAATCGTTTTCTCAAAAAAATTAATATCTTAACTGAT  
AGTTTGATCAAAGGGGCAAAACGTAGGGGCAAAACCGGAAAAATCGTTTTCTCAAATTTTCTGATGCCAAGAACTC  
TAACCAGTCTTATCTAAAAATTCCTTATGATCCGTCTCTCCGGTTACAGCCTGTGTAACCTGATTAATCCTGCCTTT  
CTAATCACCATTTCTAATGTTTAAATTAAGGGATTTTGTCTTCATTAAACGGCTTTTCGCTCATAAAAAATGTTATGACGT  
TTTGGCCGAGGCGGGAACCATCCACTTCACGAGACTGATCTCCTCTGCCGGAACACCGGGCATCTCCAACCTATA  
AGTTGGAGAAATAAGAGAATTTTCAAGATTGAGAGAATGAAAAAACCCTTAGTTCATAGGTCCATTCTCTTAGC  
GCAACTACAGAGAACAGGGGCACAAACAGGCCAAAAACGGGCACAACCTCAATGGAGTGATGCAACCTGCCTGGAGT  
AAATGATGACACAAGGCAATTGACCCACGCATGTATCTATCTATTTTCTTACACCTTCTATTACCTTCTGCTCTCT  
CTGATTTGGAAAAAGCTGAAAAAAAGGTTGAAACCAGTTCCTGAAATTATTCCTTACTTGACTAATAAGTATAT  
AAAGACGGTAGGTATTGATTGTAATCTGTAAATCTATTTCTTAACTTCTTAAATCTACTTTTATAGTTAGTCTT  
TTTTTTAGTTTTTAAACACCAAGAAGCTTAGTTTCGAATAAACACACATAAACAAACAAGCTT

**FIG. 6**

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The “delta As” version of the ADH2 / GAPDH promoter in which 10 bases of GAPDH sequence has been replaced with 14 bases of parvovirus B19 native viral sequence (underlined) as it naturally occurs immediately upstream from the initiating Met of VP1 (highlighted in black)

GGATCCTTCAATATGCGCACATACGCTGTTATGTTCAAGGTCCTTCGTTTAAGAACGAA  
 AGCGGTCTTCCTTTTGAGGGATGTTTCAAGTTGTTCAAATCTATCAAATTTGCAAATCCC  
 CAGTCTGTATCTAGAGCGTTGAATCGGTGATGCGATTTGTTAATTAAATTGATGGTGTCA  
 CCATTACCAGGTCTAGATATACCAATGGCAAACCTGAGCACACAATACCAGTCCGGATCA  
 ACTGGCACCATCTCTCCCGTAGTCTCATCTAATTTTTCTTCCGGATGAGGTTCCAGATAT  
 ACCGCAACACCTTTATTATGGTTTCCCTGAGGGAATAATAGAATGTCCATTTCGAAATCA  
 CCAATTCTAAACCTGGGCGAATTGTATTTCTGGGTTTGTAACTCGTTCCAGTCAGGAATG  
 TTCCACGTGAAGCTATCTTCCAGCAAAGTCTCCACTTCTTCATCAAATTGTGGGAGAATA  
 CTCCCAATGCTCTTATCTATGGGACTTCCGGGAAACACAGTACCGATACTTCCCAATTCTG  
 TCTTCAGAGCTCATTGTTTGTGTTGAAGAGACTAATCAAAGAATCGTTTTCTCAAAAAAAT  
 TAATATCTTAACTGATAGTTTGATCAAAGGGGCAAACCTAGGGGCAAACAAACGGAAAA  
 ATCGTTTCTCAAATTTTCTGATGCCAAGAACTCTAACCAGTCTTATCTAAAAATTGCCTT  
 ATGATCCGTCTCTCCGGTTACAGCCTGTGTAACCTGATTAATCCTGCCTTTCTAATCACCA  
 TTCTAATGTTTTAATTAAGGGATTTTGTCTTCATTAACGGCTTTTCGCTCATAAAAAATGTT  
 ATGACGTTTTGCCCGCAGGCGGGAAACCATCCACTTCACGAGACTGATCTCCTCTGCCGG  
 AACACCGGGCATCTCCAACCTTATAAGTTGGAGAAATAAGAGAATTTAGATTGAGAGAAT  
 GAAAAAAAAAAACCCTTAGTTTCATAGGTCCATTCTCTTAGCGCAACTACAGAGAACAGGG  
 GCACAAACAGGCAAAAAACGGGCACAACCTCAATGGAGTGATGCAACCTGCCTGGAGTAA  
 ATGATGACACAAGGCAATTGACCCACGCATGTATCTATCTCATTTTCTTACACCTTCTAT  
 TACCTTCTGCTCTCTCTGATTTGGAAAAAGCTGAAAAAAAAGGTTGAAACCAGTTCCTTG  
 AAATTATTCCCTACTTGACTAATAAGTATATAAAGACGGTAGGTATTGATTGTAATTCT  
 GTAAATCTATTTCTTAACTTCTTAAATTCTACTTTTATAGTTAGTCTTTTTTTTAGTTT  
 TAAAACACCAAGAAGCTTAGTTTTCGAATAAACACACATAAACAAACAAGCTTTGTAGATT**ATG**

**FIG. 7**

ADH2 / GAPDH promoter (as BamHI / MluI fragment) with TATA deletion, used in expression of ParvoB19 VP1 and VP1mut

GGATCCTTCAATATGCGCACATACGCTGTTATGTTCAAGGTCCTTCGTTTAAGAACGAA  
 AGCGGTCTTCCTTTTGAGGGATGTTTCAAGTTGTTCAAATCTATCAAATTTGCAAATCCC  
 CAGTCTGTATCTAGAGCGTTGAATCGGTGATGCGATTTGTTAATTAAATTGATGGTGTCA  
 CCATTACCAGGTCTAGATATACCAATGGCAAACCTGAGCACACAATACCAGTCCGGATCA  
 ACTGGCACCATCTCTCCCGTAGTCTCATCTAATTTTTCTTCCGGATGAGGTTCCAGATAT  
 ACCGCAACACCTTTATTATGGTTTCCCTGAGGGAATAATAGAATGTCCATTTCGAAATCA  
 CCAATTCTAAACCTGGGCGAATTGTATTTCTGGGTTTGTAACTCGTTCCAGTCAGGAATG  
 TTCCACGTGAAGCTATCTTCCAGCAAAGTCTCCACTTCTTCATCAAATTGTGGGAGAATA  
 CTCCCAATGCTCTTATCTATGGGACTTCCGGGAAACACAGTACCGATACTTCCCAATTCTG  
 TCTTCAGAGCTCATTGTTTGTGTTGAAGAGACTAATCAAAGAATCGTTTTCTCAAAAAAAT  
 TAATATCTTAACTGATAGTTTGATCAAAGGGGCAAACCTAGGGGCAAACAAACGGAAAA  
 ATCGTTTCTCAAATTTTCTGATGCCAAGAACTCTAACCAGTCTTATCTAAAAATTGCCTT  
 ATGATCCGTCTCTCCGGTTACAGCCTGTGTAACCTGATTAATCCTGCCTTTCTAATCACCA  
 TTCTAATGTTTTAATTAAGGGATTTTGTCTTCATTAACGGCTTTTCGCTCATAAAAAATGTT  
 ATGACGTTTTTGCCCGCAGGCGGGAAACCATCCACTTCACGAGACTGATCTCCTCTGCCGG  
 AACACCGGGCATCTCCAACCTTATAAGTTGGAGAAATAAGAGAATTTAGATTGAGAGAAT  
 GAAAAAAAAAAACCCTTAGTTTCATAGGTCCATTCTCTTAGCGCAACTACAGAGAACAGGG  
 GCACAAACAGGCAAAAAACGGGCACAACCTCAATGGAGTGATGCAACCTGCCTGGAGTAA  
 ATGATGACACAAGGCAATTGACCCACGCATGTATCTATCTCATTTTCTTACACCTTCTAT  
 TACCTTCTGCTCTCTCTGATTTGGAAAAAGCTGAAAAAAAAGGTTGAAACCAGTTCCTTG  
 AAATTATTCCCTAGCATTGTAATTCTGTAAATCTATTTCTTAACTTCTTAAATTCTAC  
 TTTTATAGTTAGTCTTTTTTTTAGTTTAAACACCAAGAAGCTTAGTTTTCGAATAAACAC  
 ACATAAACAAAC**ACGCGT**

**FIG. 8**

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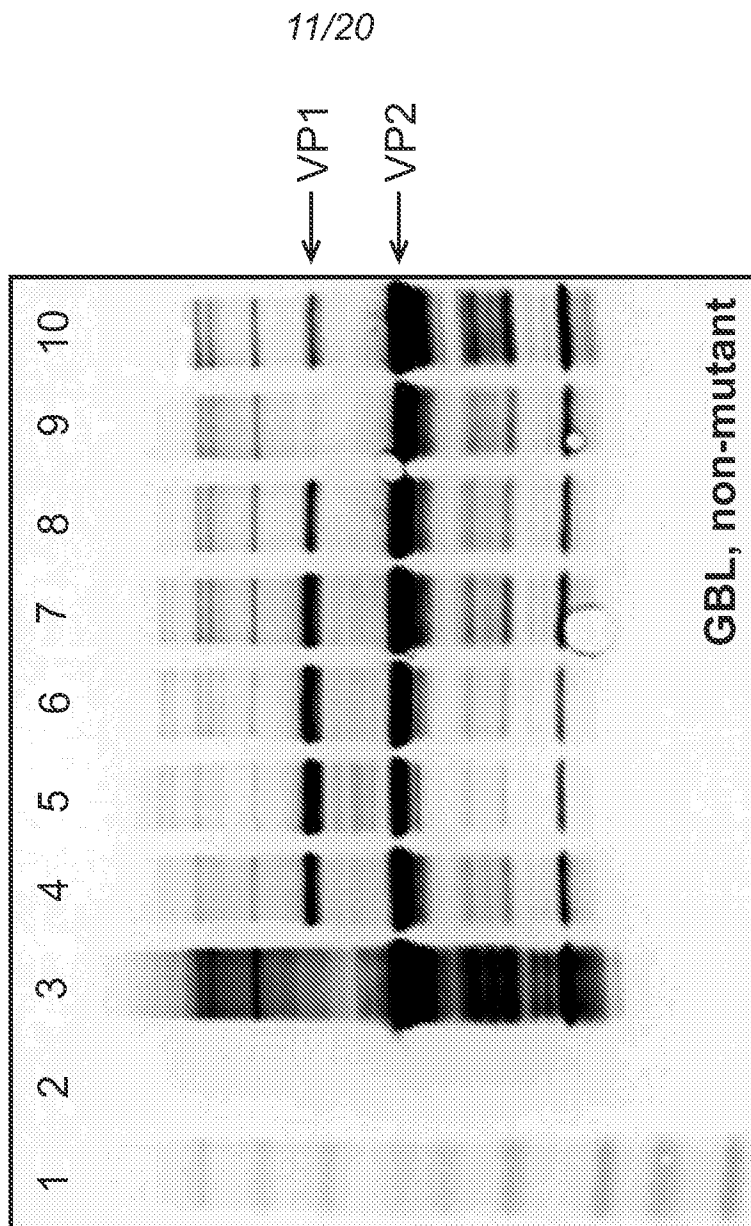
ParvoB19 VP1 with 1SS (synthetic start-stop sequence, underlined) upstream from the junction between the ADH2 / GAPDH promoter and the initiating Met of VP1 (highlighted in black)

ACGCGTATGATGCCTAGTTAATGAACAAAACAAA**ATG**TCTAAGAAATCCGGAAAATGGTG  
 GGAATCTGATGATAAATTTGCTAAGGCTGTTTACCAACAATTTGTTGAATTTTACGAAAA  
 GGTTACTGGTACTGATTTGGAATTGATTCAAATTTTGAAGGATCATTACAACATTTCTTT  
 GGATAATCCATTGGAAAATCCATCTTCATTGTTTGATTGGTTGCTAGAATTAAGAACAA  
 CTTGAAGAACTCTCCAGATTTGTATTCTCATCATTTCCAATCTCATGGTCAATTGTCTGA  
 TCATCCACATGCTTTATCTTCATCTTCATCTCATGCTGAACCAAGAGGTGAAAATGCTGT  
 TTTATCTTCTGAAGATTTGCATAAACAGGTCAAGTTTCTGTTCAATTGCCAGGTACTAA  
 TTACGTTGGTCCAGGTAATGAATTGCAAGCTGGTCCACCACAATCTGCTGTTGATTCTGC  
 TGCTAGAATTCATGATTTTCAGATACTCTCAATTGGCTAAGTTGGGTATTAATCCATATAC  
 TCATTGGACTGTTGCTGATGAAGAATTGTTGAAGAACATTAAGAATGAACTGGTTTTCA  
 AGCTCAAGTTGTTAAAGATTACTTCACTTTGAAAGGTGCTGCTGCTCCAGTTGCTCATTT  
 TCAAGTTTCTTTGCCAGAAGTTCCAGCTTATAACGCTTCTGAAAAATATCCATCTATGAC  
 ATCTGTAAATTCTGCTGAAGCATCTACTGGTGCAGGTGGAGGTGGTTCTAATTCTGTAA  
 ATCTATGTGGTCTGAAGGTGCTACTTTTTCTGCTAATTCAGTTACTTGTACTTTCTCTAG  
 ACAATTCTTGATTCCATATGATCCAGAACATCATTACAAAGTTTTTTCACCAGCTGCTTC  
 ATCTTGTCATAATGCTTCAGGTAAAGAAGCTAAGGTTTGTACTATTTCTCCAATTATGGG  
 TTATTCTACTCCTTGGAGATACTTGGATTTTAATGCTTTGAACTTGTTTTTTCTCCATT  
 GGAATTTCAACATTTGATTGAAAACACGGTCTATTGCTCCAGATGCTTTGACTGTTAC  
 TATTTCTGAAATTGCTGTTAAGGATGTTACTGATAAAACAGGTGGTGGTGTTCAGTTAC  
 TGATTCTACTACTGGTAGATTGTGCATGTTGGTTGATCATGAATACAAATACCCATACGT  
 TTTGGGTCAAGGTCAAGATACTTTGGCTCCAGAATTGCCAATTTGGGTTTATTTTCCACC  
 ACAATACGCTTATTTGACTGTTGGTGATGTTAATACTCAAGGTATTTCTGGTGATTCTAA  
 AAAGTTGGCTTCTGAAGAATCTGCTTTTTACGTTTTTGGAACATTCTTCTTTTCAATTGTT  
 GGGTACTGGTGGTACTGCTTCTATGTCTTACAAATTTCCACCAGTTCCACCTGAAAATTT  
 GGAAGGTTGTTCTCAACATTTTACGAAATGTACAATCCATTGTATGGTTCTAGATTGGG  
 TGTTCCAGATACTTTGGGTGGTGATCCAAAATTTAGATCTTTGACTCATGAAGATCATGC  
 TATTCAACCACAAAATTTTCATGCCAGGTCCATTGGTTAATTCTGTTTCTACTAAAGAAGG  
 TGATTCTTCTAATACAGGTGCTGGTAAAGCATTGACTGGTTTGTCTACTGGTACTTCTCA  
 AAACACTAGAATTTCTTTAAGACCAGGTCCAGTTTCACAACCATATCATCATTGGGATAC  
 TGATAAGTACGTTACTGGTATTAATGCTATTTTACATGGTCAAACACTATTATGGTAATGC  
 TGAAGATAAAGAATATCAACAAGGTGTTGGTAGATTTCCAAACGAAAAAGAACAATTGAA  
 ACAATTGCAAGGTTTGAATATGCATACTTACTTTCCAAACAAAGGTACTCAACAATACAC  
 TGATCAAATTGAAAGACCATTGATGGTTGGTTCTGTTTGAATAGAAGAGCTTTGCATTA  
 TGAATCTCAATTGTGGTCTAAGATTCCAAATTTAGATGATTCTTTCAAGACTCAATTTGC  
 TGCTTTGGGTGGTTGGGGTTTGCATCAACCTCCACCACAAATTTTCTTGAAGATTTTGCC  
 ACAATCTGGTCCAATTGGTGGTATTAAATCTATGGGTATTACTACTTTGGTTCAATATGC  
 TGTTGGTATTATGACTGTTACAATGACTTTTAAAGTTGGGTCCAAGAAAAGCTACAGGTAG  
 ATGGAATCCACAACCAGGTGTTTATCCACCACATGCTGCTGGTCATTTGCCTTACGTTTT  
 GTATGATCCAACTGCTACTGATGCTAAACAACATCATAGACATGGTTATGAAAAACCTGA  
 AGAATTGTGGACTGCTAAATCTAGAGTTCATCCATTGTAATGAGTCGAC

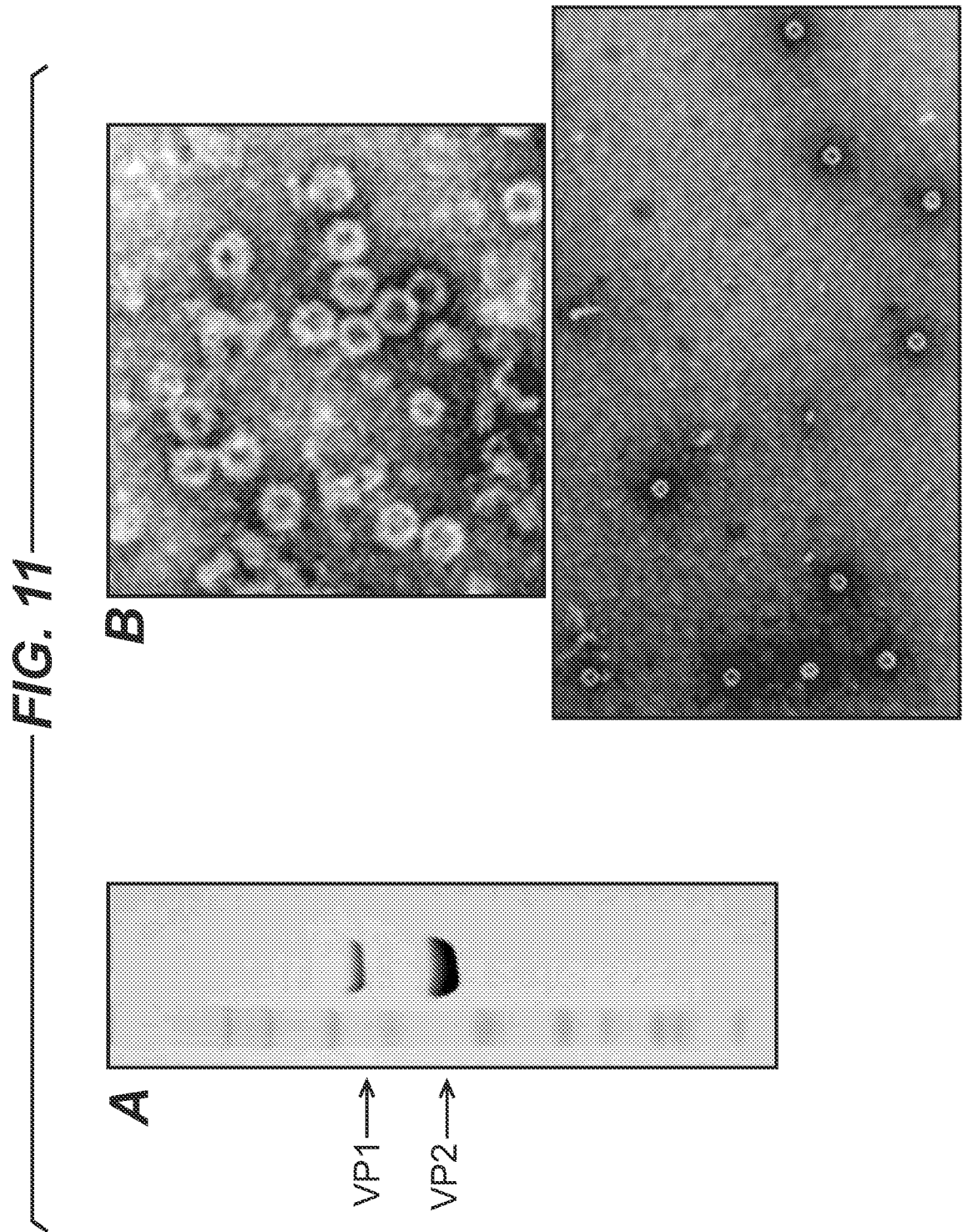
**FIG. 9**

**FIG. 10**

|    |                                      |
|----|--------------------------------------|
| 1  | Invitrogen Benchmark stds            |
| 2  | pCDC7-empty vector control           |
| 3  | VP2                                  |
| 4  | VP2+ $\Delta$ TATA.VP1               |
| 5  | $\Delta$ TATA.VP2+ $\Delta$ TATA.VP1 |
| 6  | VP2+dAs.VP1                          |
| 7  | VP2+1SS.VP1                          |
| 8  | VP2+2SS.VP1                          |
| 9  | VP2+1AUG.VP1                         |
| 10 | VP2+mc.VP1                           |



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3wp1 Pooled Sera Wild Type VLP

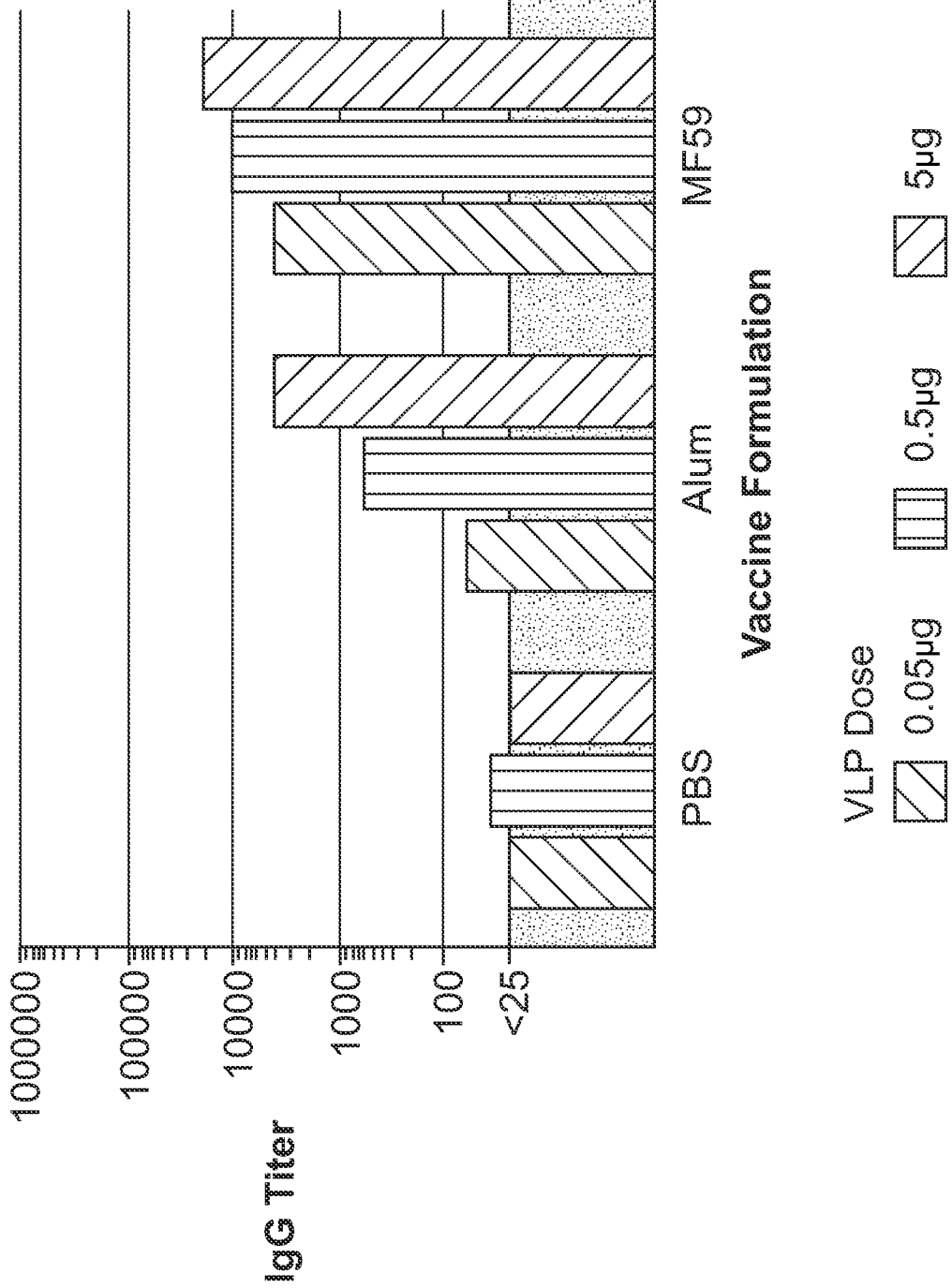


FIG. 12A

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3wp2 Pooled Sera Wild Type VLP

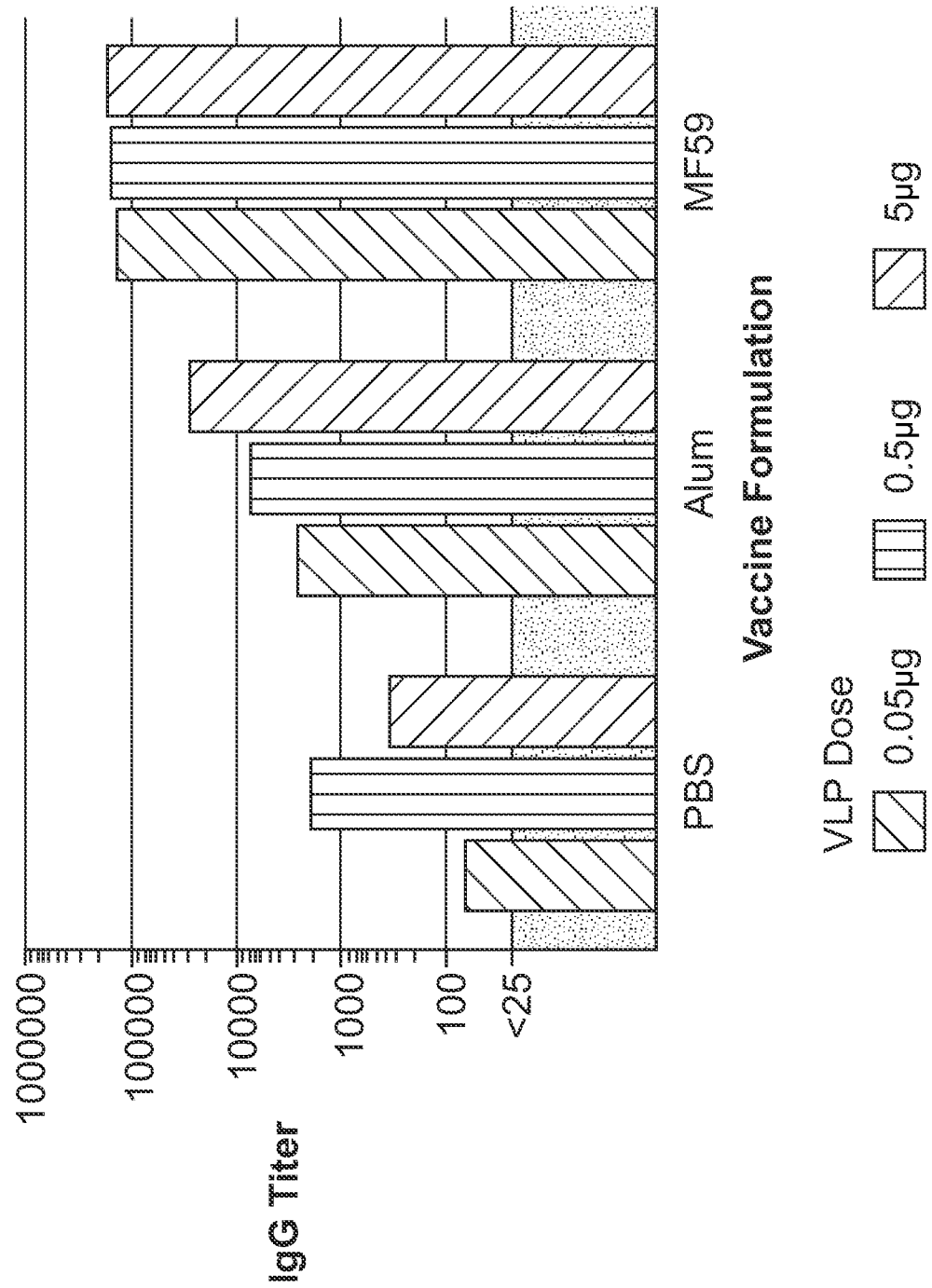


FIG. 12B

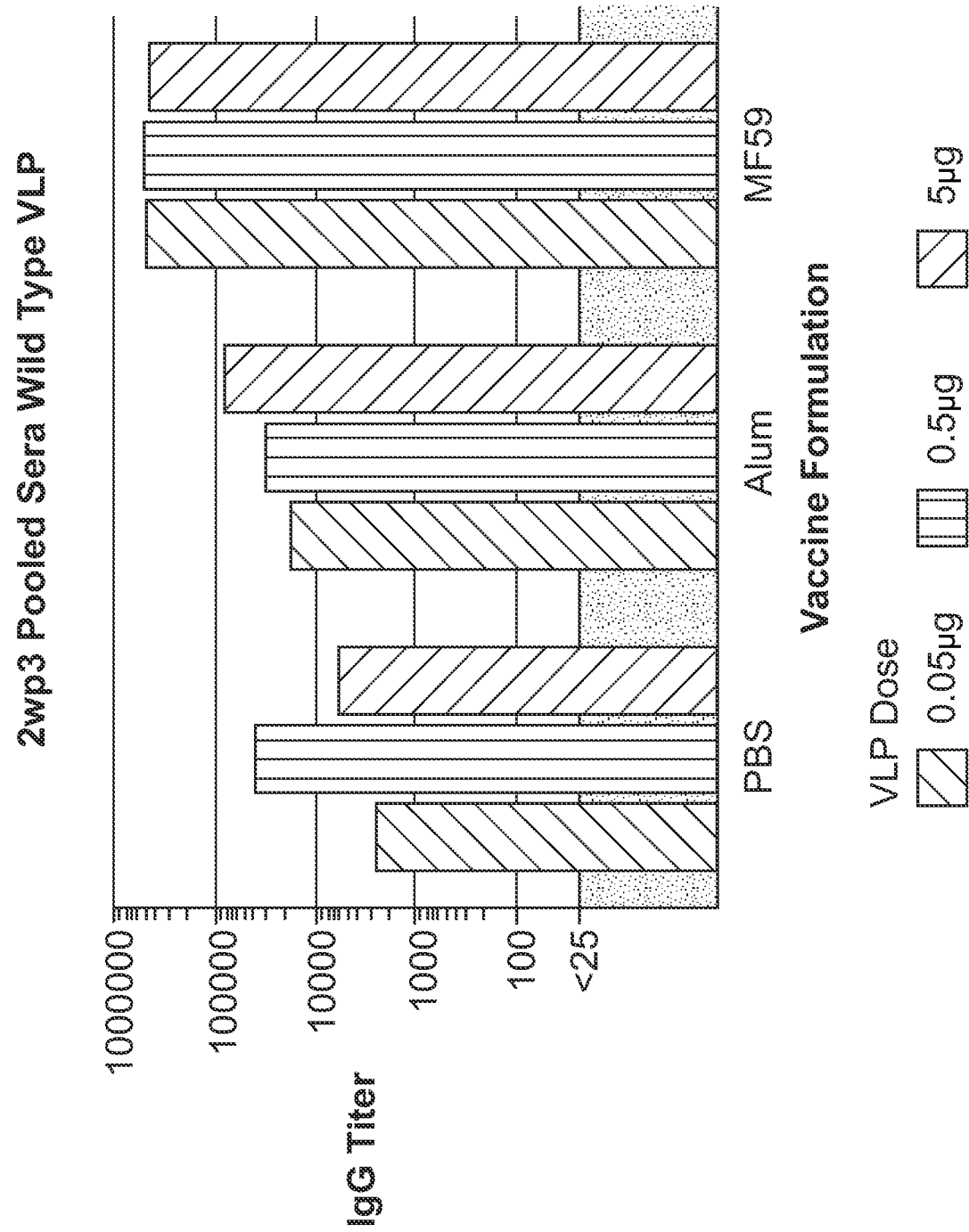


FIG. 12C

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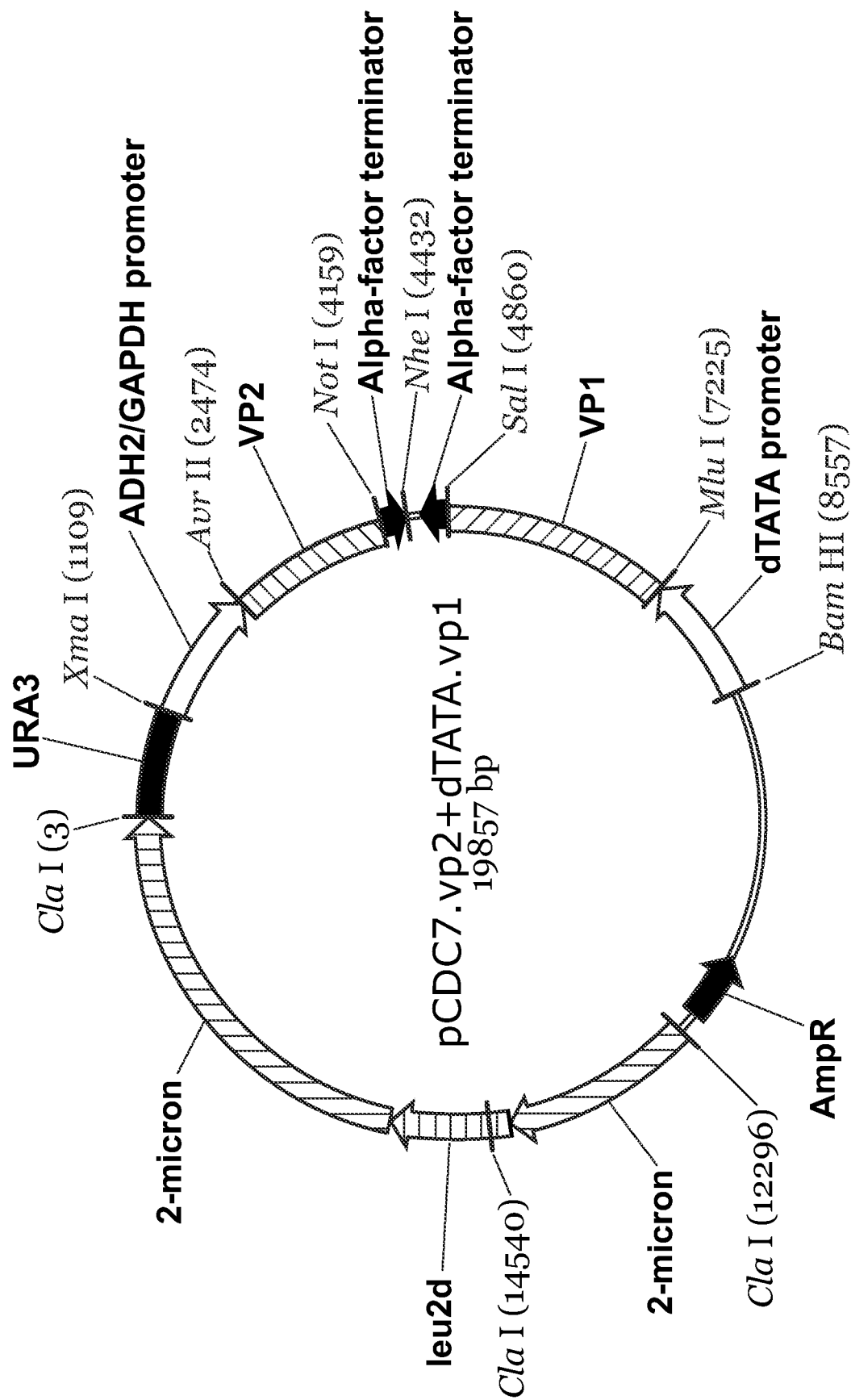


FIG. 13

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VP1 wild type amino acid sequence

Amino Acids :781

Molecular Weight : 84 kDa

MSKKSGKWWESDDKFAKAVYQQFVEFYEKVTGTDLELIQILKDHYNISLDNPLENPSSLF  
 DLVARIKNNLKNSPDLYSHHFQSHGQLSDHPHALSSSSSSHAEPGENAVLSSSEDLHKPGQ  
 VSVQLPGTNYVGPNGNELQAGPPQSAVDSAARIHDFRYSQLAKLGINPYTHWTVADEELLK  
 NIKNETGFQAQVVKDYFTLKGAAAPVAHFQGSLEVPAYNASEKYPSTSVNSAEASTGA  
 GGGGSNSVKSMWSEGATFSANSVTCTFSRQFLIPYDPEHHYKVFSPAASSCHNASGKEAK  
 VCTISPIMGYSTPWRYLDFNALNLFFSPLEFQHLENIYGSIAPDALTVTISEIAVKDVT  
 KTGGGVQVTDSTTGRLCMLVDHEYKYPYVLGQGQDTLAPELPIWVYFPPQYAYLTVGDVN  
 TQGISGDSKKLASEESAFYVLEHSSFQLLGTGGTASMSYKFPVPPEPPELEGCSQHFYEM  
 NPLYGSRLGVPDTLGGDPKFRSLTHEDHAIQPQNFMGPPLVNSVSTKEGDSSNTGAGKAL  
 TGLSTGTSQNTRISLRPGPVSQPYHHWDTDKYVTGINAISHGQTTYGNAEDKEYQQGVGR  
 FPNEKEQLKQLQGLNMHTYFPNKGTTQYTDQIERPLMVGSVWNRRALHYESQLWSKIPNL  
 DDSFKTQFAALGGWGLHQPPPQIFLKILPQSGPIGGIKSMGITTLLVQYAVGIMTVTMTFK  
 LGPRKATGRWNPQPGVYPPHAAGHLPYVLYDPTATDAKQHHRHGYEKPEELWTAKSRVHP  
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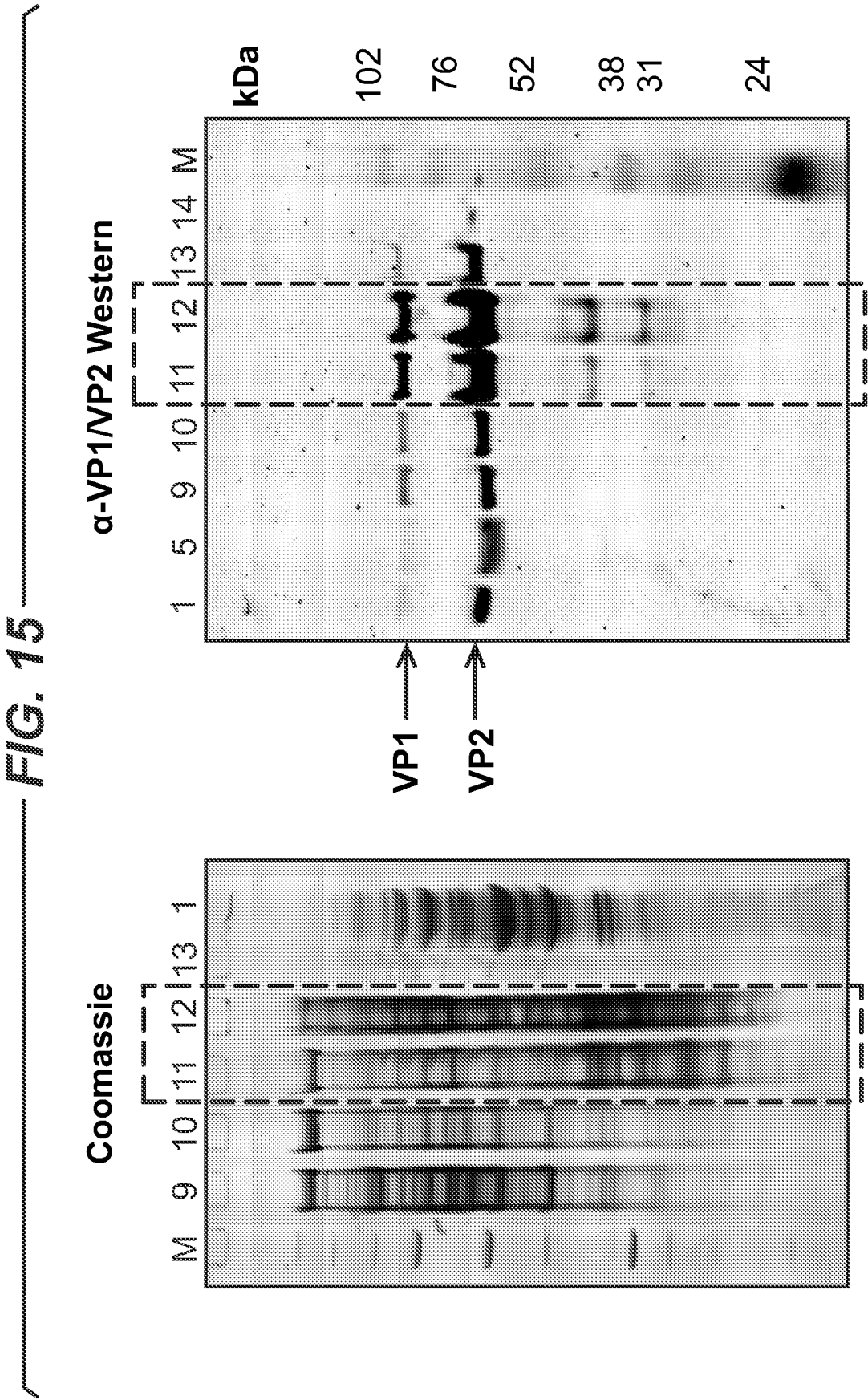
VP2 amino acid sequence

Amino Acids :554

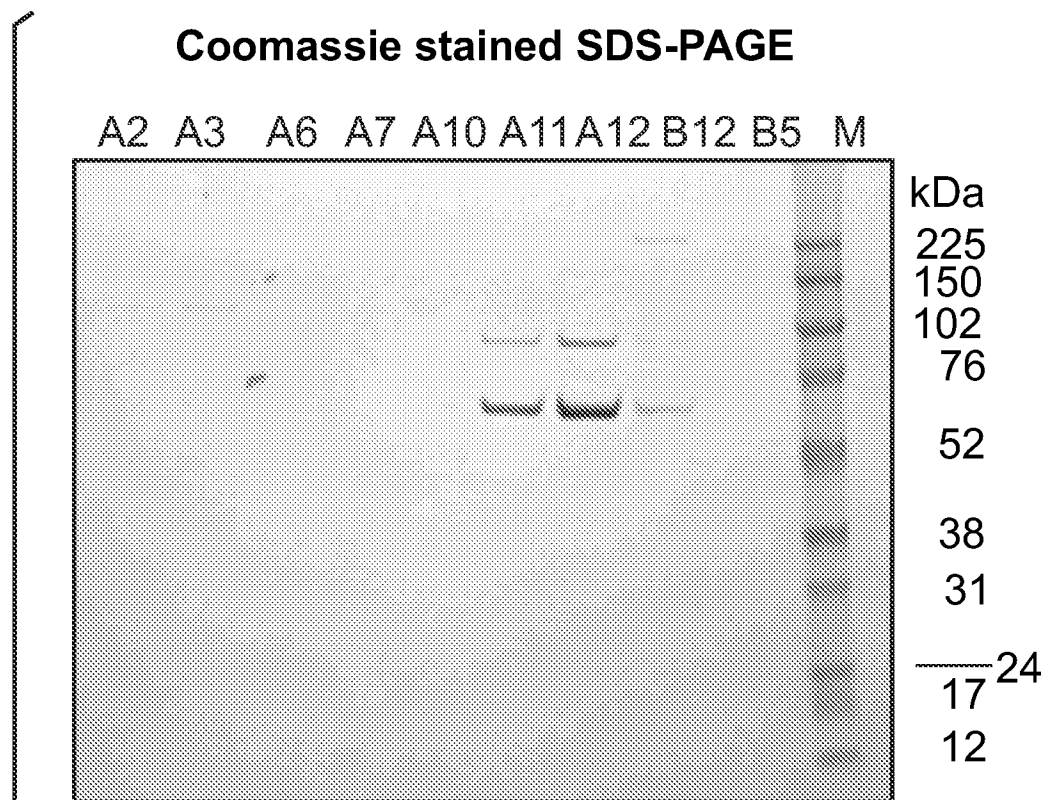
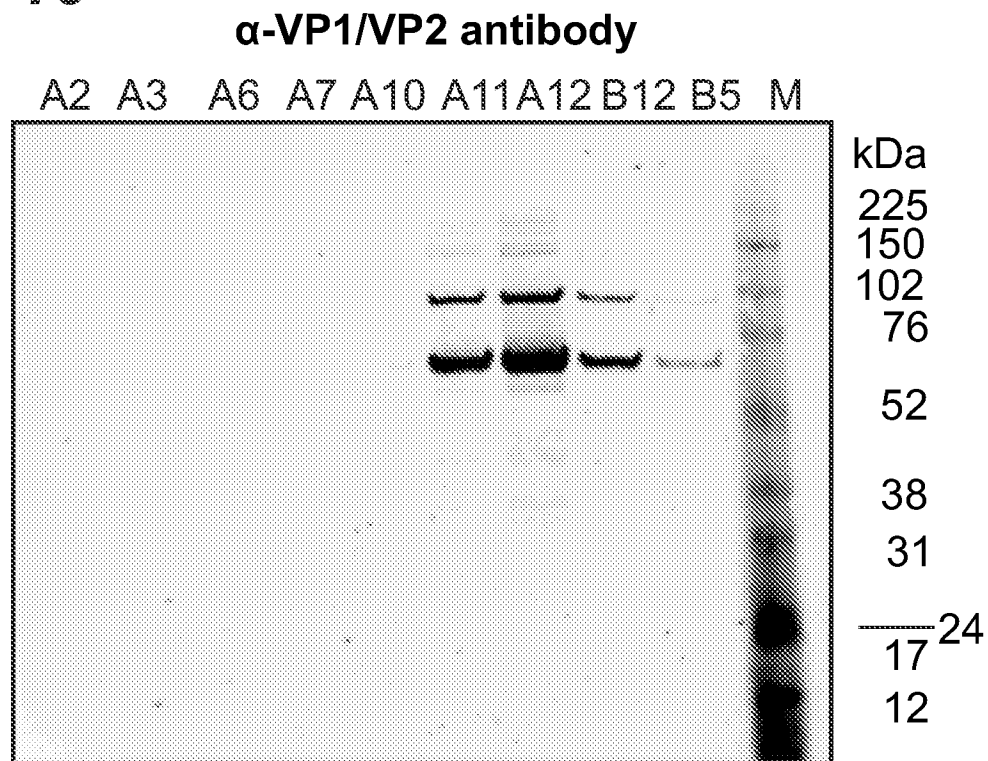
Molecular Weight : 58 kDa

MTSVNSAEASTGAGGGGSNSVKSMWSEGATFSANSVTCTFSRQFLIPYDPEHHYKVFSPA  
 ASSCHNASGKEAKVCTISPIMGYSTPWRYLDFNALNLFFSPLEFQHLENIYGSIAPDALT  
 VTISEIAVKDVTDKTGGGVQVTDSTTGRLCMLVDHEYKYPYVLGQGQDTLAPELPIWVYF  
 PPQYAYLTVGDVNTQGISGDSKKLASEESAFYVLEHSSFQLLGTGGTASMSYKFPVPPE  
 NLEGCSQHFYEMYNPLYGSRLGVPDTLGGDPKFRSLTHEDHAIQPQNFMGPPLVNSVSTK  
 EGDSSNTGAGKALTGLSTGTSQNTRISLRPGPVSQPYHHWDTDKYVTGINAISHGQTTYG  
 NAEDKEYQQGVGRFPNEKEQLKQLQGLNMHTYFPNKGTTQYTDQIERPLMVGSVWNRRAL  
 HYESQLWSKIPNLDDSFKTQFAALGGWGLHQPPPQIFLKILPQSGPIGGIKSMGITTLLVQ  
 YAVGIMTVTMTFKLGPRKATGRWNPQPGVYPPHAAGHLPYVLYDPTATDAKQHHRHGYEK  
 PEELWTAKSRVHPL

**FIG. 14**

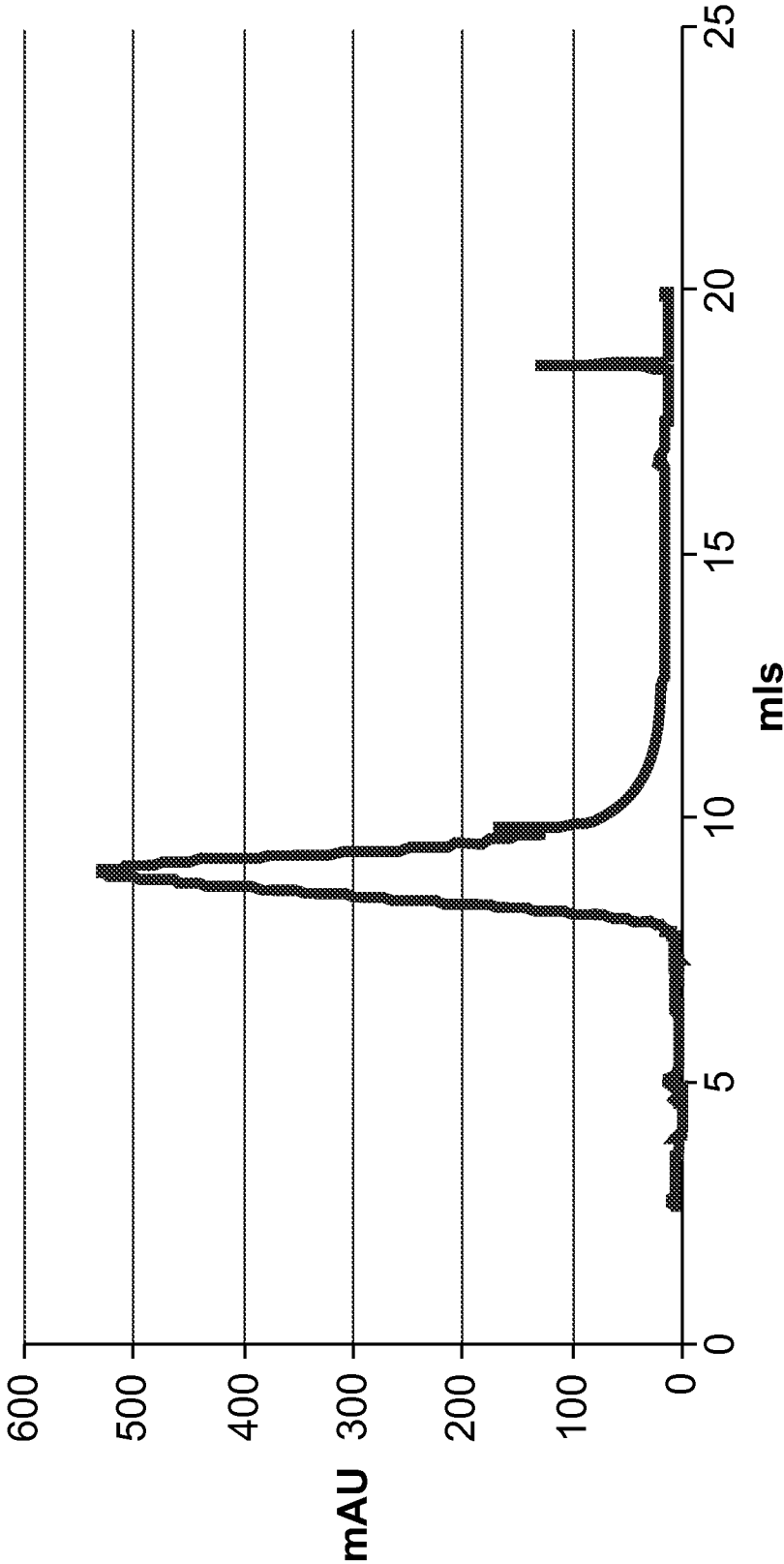


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**FIG. 16**

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FIG. 17



## INTERNATIONAL SEARCH REPORT

International application No

PCT/US2011/031630

A. CLASSIFICATION OF SUBJECT MATTER  
 INV. C12N7/04 C07K14/015  
 ADD.

According to International Patent Classification (IPC) or to both national classification and IPC

## B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

C12N C07K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

EPO-Internal, BIOSIS, CHEM ABS Data, WPI Data

## C. DOCUMENTS CONSIDERED TO BE RELEVANT

| Category* | Citation of document, with indication, where appropriate, of the relevant passages                                                                                                                                                                                                                                                                                                                                                                                                                                                               | Relevant to claim No.                                                       |
|-----------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----------------------------------------------------------------------------|
| X         | <p>Shelly D. and Van Cleave V.: "Parvovirus B19 VLP vaccine manufacturing", Genetic Engineering and Biotechnology News, vol. 29, no. 16<br/>           15 September 2009 (2009-09-15), pages 1-3, XP002647110,<br/>           Retrieved from the Internet:<br/>           URL: <a href="http://www.genengnews.com/keywordsandtools/print/1/12962/">http://www.genengnews.com/keywordsandtools/print/1/12962/</a><br/>           [retrieved on 2011-07-01]<br/>           the whole document</p> <p style="text-align: center;">-----<br/>-/-</p> | <p>1-3,<br/>           8-15,<br/>           27-29,<br/>           34,35</p> |



Further documents are listed in the continuation of Box C.



See patent family annex.

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"&" document member of the same patent family

Date of the actual completion of the international search

4 July 2011

Date of mailing of the international search report

22/07/2011

Name and mailing address of the ISA/

European Patent Office, P.B. 5818 Patentlaan 2  
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Authorized officer

Mandl, Birgit

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International application No

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| C(Continuation). DOCUMENTS CONSIDERED TO BE RELEVANT |                                                                                                                                                                                                                                                                                    |                                  |
|------------------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----------------------------------|
| Category*                                            | Citation of document, with indication, where appropriate, of the relevant passages                                                                                                                                                                                                 | Relevant to claim No.            |
| X                                                    | KAJIGAYA S ET AL: "SELF-ASSEMBLED B19 PARVOVIRUS CAPSIDS, PRODUCED IN A BACULOVIRUS SYSTEM, ARE ANTIGENICALLY AND IMMUNOGENICALLY SIMILAR TO NATIVE VIRIONS", MICROBIOLOGY, vol. 88, 1 June 1991 (1991-06-01), pages 4646-4650, XP001118322, ISSN: 1350-0872<br>the whole document | 1-3,<br>8-15,<br>27-29,<br>34,35 |
| X                                                    | US 2002/119527 A1 (BROWN CAROLINE SARAH [NL]) 29 August 2002 (2002-08-29)<br><br>paragraphs [0002], [0014]<br>paragraphs [0022] - [0024]<br>paragraph [0060]                                                                                                                       | 1-3,<br>8-15,<br>27-29,<br>34,35 |
| A                                                    | WO 2007/084773 A2 (ASKLEPIO BIOPHARMACEUTICAL INC [US]; SAMULSKI RICHARD J [US]; CHEN HAI)<br>26 July 2007 (2007-07-26)<br>paragraph [0017]                                                                                                                                        | 1-35                             |

# INTERNATIONAL SEARCH REPORT

Information on patent family members

International application No

PCT/US2011/031630

| Patent document<br>cited in search report | Publication<br>date | Patent family<br>member(s) | Publication<br>date      |
|-------------------------------------------|---------------------|----------------------------|--------------------------|
| US 2002119527                             | A1                  | 29-08-2002                 | NONE                     |
| -----                                     |                     |                            |                          |
| WO 2007084773                             | A2                  | 26-07-2007                 | EP 1981548 A2 22-10-2008 |
|                                           |                     | US 2009191597 A1           | 30-07-2009               |
| -----                                     |                     |                            |                          |