METHOD OF PREPARING PLANT-DERIVED VLPS

FIELD OF INVENTION

[0001] The present invention relates to methods of preparing plant-derived virus-like particles (VLPs).

BACKGROUND OF THE INVENTION

[0002] Current recombinant expression strategies in host cells such as *E. coli*, insect cell culture, and mammalian cell culture express and secrete proteins at very high level in the culture media. Using these systems high levels of expression, proper protein folding and post-translational modification of proteins, is achieved. Furthermore, purification of the expressed protein is simplified since intracellular proteins may be readily segregated from other components (DNA, vesicle, membranes, pigments, and so on). For plant or yeast expression systems, the cell wall prevents secretion of expressed protein into the culture media.

[0003] One of the primary methods to combat viral infections is by vaccination. Production of vaccines in response to an outbreak or epidemic, or to meet seasonal demands (e.g. the annual 'flu season' occurring in the fall, or the recent 'swine flu' outbreaks observed worldwide) requires the generation of sufficient quantity of vaccine given the short notice period. Current worldwide production of influenza vaccine may be insufficient in the face of a worldwide flu pandemic. Furthermore, dominant influenza strains change from year to year, thus stockpiling at low-need times in the year is not practical. Economical, large scale production of an effective influenza vaccine is of significant value.

[0004] Virus-like particles (VLPs) may be employed to prepare influenza vaccines. Suprastructures such as VLPs mimic the structure of the viral capsid, but lack a genome, and thus cannot replicate or provide a means for a secondary infection. VLPs offer an improved alternative to isolated (soluble) recombinant antigens for stimulating a strong immune response. VLPs are assembled upon expression of specific viral proteins and present an external surface resembling that of their cognate virus but, unlike true viral particle, do not incorporate genetic material. The presentation of antigens in a particulate and multivalent structure similar to that of the native virus achieves an enhanced stimulation of the immune response with balanced humoral and cellular components. Such improvement over the stimulation by isolated antigens is

believed to be particularly true for enveloped viruses as enveloped VLPs present the surface antigens in their natural membrane-bound state (Grgacic and Anderson, 2006, Methods 40, 60-65). Furthermore, Influenza VLPs, with their nanoparticle organization, have been shown to be better vaccine candidates compared to recombinant hemagglutinin (HA) (i.e. monomeric HA, or HA organized into rosettes; assembly of 3-8 trimers of HA), and they are able to activate both humoral and cellular immune response. (Bright, R.A., et. al.,. 2007, *Vaccine* **25**, 3871-3878).

[0005] The vast majority of the influenza vaccines currently on the market are composed of viral particle or virus antigens obtained from egg-grown virions. The production of egg-derived vaccines relies on the culture of live viruses in embryonated hen eggs. Split-influenza vaccines are obtained after chemical inactivation and disruption of purified virions with a detergent. Recombinant influenza antigens are an effective alternative to virus-derived antigens as pandemic vaccine products. Recombinant antigens can be produced from information on the genetic makeup of a new strain once this information is made available, and allows a rapid initiation of the production process. However, purified recombinant HA subunits appear less efficacious than inactivated split-influenza vaccines and higher antigen content is required to generate a potent immune response (Treanor et al., 2007, *J. Am. Med. Assoc.* **297**, 1577-1582).

[0006] Influenza VLPs have been obtained in cultured mammalian cells from the co-expression of all 10 influenza proteins (Mena et al., 1996, *J. Virol.* **70**, 5016-5024). Several viral proteins are dispensable for the production of VLPs, and influenza VLPs in vaccine development programs have been produced from the co-expression of the 2 major antigenic envelope proteins (HA and NA) with M1 or from the co-expression of HA and M1 only (Kang et al., 2009, *Virus Res.* **143**, 140-146). Chen et al. (2007, *J. Virol.* **81**, 7111-7123) have shown that HA alone is capable of driving VLP formation and budding and M1 co-expression could be omitted in their system. However, since HA was found to bind to sialylated glycoproteins on the surface of the mammalian cells producing the VLPs, a viral sialidase was co-expressed to allow the release of VLPs from the producing cell after budding.

[0007] A simpler VLP production system, for example, one that relies on the expression of only one or a few viral proteins without requiring expression of non-structural viral proteins is desirable to accelerate the development of vaccines. Production of viral antigens, including VLPs, in plant systems provides an advantage for production, in that they may be grown in a greenhouse or field, and don't require aseptic tissue culture methods and handling.

[0008] PCT Publication WO 2006/119516 (to Williamson and Rybicki) discloses expression of full length and truncated human-codon optimized H5 HA of Influenza A/Vietnam/1194/2004 in plants. The truncated construct lacks the membrane anchoring domain. The highest accumulation of HA protein was obtained with constructs that targeted to the ER. Constructs lacking a membrane targeting domain did not yield detectable HA. The production of VLPs was not reported.

[0009] The production of influenza HA VLPs that comprise a lipid envelope has been previously described by the inventors in WO 2009/009876 and WO 2009/076778 (to D'Aoust et al.; both of which are incorporated herein by reference). For enveloped viruses, it may be advantageous for a lipid layer or membrane to be retained by the virus. The composition of the lipid may vary with the system (e.g. a plant-produced enveloped virus would include plant lipids or phytosterols in the envelope), and may contribute to an improved immune response.

[0010] The assembly of enveloped VLPs in transgenic tobacco expressing the HBV surface antigen (HBsAg) was described by Mason et al.(1992, *Proc. Natl. Acad. Sci. USA* **89**, 11745-11749). Plant-produced HBV VLPs were shown to induce potent B- and T-cell immune responses in mice when administered parenterally (Huang et al., 2005, *Vaccine* **23**, 1851-1858) but oral immunization through feeding studies only induced a modest immune response (Smith et al., 2003, *Vaccine* **21**, 4011-4021). Greco (2007, *Vaccine* **25**, 8228-8240) showed that human immunodeficiency virus (HIV) epitopes in fusion with HBsAg accumulated as VLP when expressed in transgenic tobacco and *Arabidopsis*, creating a bivalent VLP vaccine.

[0011] Expression of the viral capsid protein (NVCP) in transgenic tobacco and potato plants resulted in the assembly of non-enveloped VLPs (Mason et al., 1996, *Proc. Natl. Acad. Sci. USA* **93**, 5335-5340). NVCP VLPs have been produced in agroinfiltrated *N. benthamiana* leaves (Huang et al. 2009, *Biotechnol. Bioeng.* **103**, 706-714) and their immunogenicity upon oral administration demonstrated in mice (Santi et al., 2008, *Vaccine* **26**, 1846-1854). Administration of 2 or 3 doses of raw potatoes containing 215-751 μg of NVCP in the form of VLPs to healthy adult volunteers resulted in development of an immune response in and 95% of the immunized volunteers (Tacket et al. 2000, *J. Infect. Dis.* **182**, 302-305). Non-enveloped VLPs have also been obtained from the expression of HBV core antigen (HBcAg; Huang et al., 2009, *Biotechnol. Bioeng.* **103**, 706-714), and the human papillomavirus (HPV) major capsid protein L1 (Varsani et al., 2003, Arch. *Virol.* **148**, 1771-1786).

[0012] It may be desirable to separate the VLPs from some, or all of the proteins, carbohydrates, etc. present in the plant or plant matter before the VLP is used in vaccine formulation. A method for extracting protein from the intercellular space of plants, comprising a vacuum and centrifugation process to provide an interstitial fluid extract comprising the protein of interest is described in PCT Publication WO 00/09725 (to Turpen et al.). This approach is suitable for small proteins (of 50 kDa or smaller) that pass through pores under vacuum and centrifugation, but is not suitable for larger superstructure proteins or protein complexes such as a VLP.

[0013] McCormick et al 1999 (Proc Natl Acad Sci USA 96:703-708) discloses use of a rice amylase signal peptide fused to a single-chain Fv (scFv) epitope to target the expressed protein to the extracellular compartment, followed by vacuum infiltration of leaf and stem tissue for recovery of the scFv polypeptides. Moehnke et al., 2008 (Biotechnol Lett 30:1259-1264) describes use of the vacuum infiltration method of McCormick to obtain a recombinant plant allergen from tobacco using an apoplastic extraction. PCT Publication WO 2003/025124 (to Zhang et al) discloses expression of scFv immunoglobulins in plants, targeting to the apoplastic space using murine signal sequences.

[0014] Given the complexity of VLPs and the plant tissue in which they may be produced, methods of preparing VLPs that are substantially free of, or easily separated from plant proteins, yet retain the structural and immunogenic characteristics of the enveloped virus are desired.

SUMMARY OF THE INVENTION

[0015] The present invention relates to methods of preparing plant-derived virus-like particles (VLPs). More specifically, the present invention is directed to methods of preparing VLPs comprising influenza antigens.

[0016] It is an object of the invention to provide an improved method of preparing plant-derived virus-like particles.

[0017] The present invention provides a method (A) of preparing plant derived VLPs comprising obtaining a plant or plant matter comprising plant-derived VLPs localized within the apoplast; producing a protoplast and an apoplast fraction, the apoplast fraction comprising plant-derived VLPs; and recovering the apoplast fraction. The method may further comprise a step of purifying the plant derived VLPs from the apoplast fraction. The plant-derived VLP may be a chimeric plant-derived VLP. The plant derived VLP may be selected from the group of viral

envelope proteins, viral structural proteins, viral capsid proteins, and viral coat proteins. The plant derived VLPs may comprise influenza hemagglutinin.

[0018] The apoplast and protoplast fractions may be produced by treatment of the plant or plant matter by an enzyme composition. The enzyme composition may comprise one or more than one pectinase, one or more than one cellulase, or one or more than one pectinase and one or more than one cellulase. Furthermore, if desired, the enzyme composition does not include a lipase or protease, or the composition does not include an added lipase or protease.

[0019] Plant or plant matter may be obtained by growing, harvesting or growing and harvesting the plant. The plant matter may comprise some or all of the plant, one ore more than one plant cell, leaves, stems, roots or cultured plant cells.

[0020] The present invention provides a method of preparing plant derived VLPs as described above (Method A), wherein a nucleic acid encoding the VLP selected from the group of viral envelope proteins, viral structural proteins, viral capsid proteins, and viral coat proteins is introduced into the plant in a transient manner. Alternatively, the nucleic acid is stably integrated within a genome of the plant.

[0021] The present invention provides a method of preparing plant derived VLPs as described above (Method A) further comprising a step of purifying the plant derived VLPs from the apoplast fraction. The step of purifying may comprise filtering the apoplast fraction using depth filtration to produce a clarified extract, followed by chromatography of the clarified extract using a cation exchange resin.

[0022] Without wishing to be bound by theory, proteins obtained from the apoplast are more homogenous, as the intermediate forms of post-translationally modified proteins, or proteins comprising other types of processing that occurs in various intracellular compartments are not co-extracted. A higher degree of homogeneity of a recombinant protein typically results in a higher quality of a preparation comprising the protein, and may result in a product with beneficial properties including higher potency, longer half-life, or better immunogenic capacity. For example, blood proteins containing high-mannose glycosylation are eliminated in blood circulation more rapidly than proteins comprising complex glycosylation. A glycosylated protein produce in the apoplastic fraction exhibits more complex-type glycosylation. Therefore, an

apoplast-derived protein prepared using the methods described herein, involving cell-wall digestion, exhibit, for example, a better half life in circulation.

[0023] The present invention also provides for a method (B) of preparing plant-derived VLPs comprising a plant-derived lipid envelope, the method comprising, obtaining a plant, or plant matter comprising VLPs localized within the apoplast; treating the plant or plant matter with an enzyme composition to produce a protoplast fraction, and one or more than one apoplastic protein composition; separating the one or more than one apoplastic protein complex from the protoplast fraction, wherein the one or more than one apoplastic protein complexes comprise the VLPs. The enzyme composition may comprise one or more than one pectinase, one or more than one cellulose, or one or more than one pectinase and one or more than one cellulase. Furthermore, if desired, the enzyme composition does not include a lipase or protease, or the composition does not include an added lipase or protease. The plant-derived VLP may be a chimeric plant-derived VLP. The plant derived VLP may be selected from the group of viral envelope proteins, viral structural proteins, viral capsid proteins, and viral coat proteins. The plant derived VLPs may comprise influenza hemagglutinin.

[0024] The present invention provides a method of preparing plant derived VLPs as described above (Method B), wherein a nucleic acid encoding the VLP selected from the group of viral envelope proteins, viral structural proteins, viral capsid proteins, and viral coat proteins is introduced into the plant in a transient manner. Alternatively, the nucleic acid is stably integrated within a genome of the plant.

[0025] The present invention provides a method of preparing plant derived VLPs as described above (Method B) further comprising a step of purifying the plant derived VLPs from the apoplast fraction. The step of purifying may comprise filtering the apoplast fraction using depth filtration to produced a clarified extract, followed by chromatography of the clarified extract using a cation exchange resin.

[0026] The plant derived VLPs may include VLPs comprising one or more influenza HA polypeptides. The influenza HA polypeptide may also be a chimeric HA polypeptide. The plant-derived VLPs may further comprise hemagglutinating activity. Plant or plant matter may be obtained by growing, harvesting or growing and harvesting the plant. The plant matter may comprise some or all of the plant, or one or more than one plant cell, leaves, stems, roots or cultured cells.

[0027] The present invention also provides a method (C) of preparing plant derived VLPs, comprising obtaining a plant or plant matter comprising plant-derived VLPs, digesting the plant matter using a cell wall degrading enzyme composition to produced a digested fraction, and filtering the digested fraction to produced a filtered fraction and recovering the plant-derived VLPs from the filtered fraction.

[0028] The enzyme composition may comprise one or more than one pectinase, one or more than one cellulose, or one or more than one pectinase and one or more than one cellulase. Furthermore, if desired, the enzyme composition does not include a lipase or protease, or the composition does not include an added lipase or protease. The plant-derived VLP may be a chimeric plant-derived VLP. The plant derived VLP may be selected from the group of viral envelope proteins, viral structural proteins, viral capsid proteins, and viral coat proteins. The plant derived VLPs may comprise influenza hemagglutinin.

[0029] The present invention provides a method of preparing plant derived VLPs as described above (Method C), wherein a nucleic acid encoding the VLP selected from the group of viral envelope proteins, viral structural proteins, viral capsid proteins, and viral coat proteins is introduced into the plant in a transient manner. Alternatively, the nucleic acid is stably integrated within a genome of the plant.

[0030] The present invention provides a method of preparing plant derived VLPs as described above (Method C) further comprising a step of separating the VLPs in the filtered fraction from the cellular debris and insoluble materials. The step of separating may be performed by centrifugation, by depth filtration, or bother centrifugation and depth filtration to produce a clarified fraction. The plant derived VLPs may be further purified by chromatography, for example, the clarified extract may be purified using a cation exchange resin.

[0031] The plant derived VLPs may include VLPs comprising one or more influenza HA polypeptides. The influenza HA polypeptide may also be a chimeric HA polypeptide. The plant-derived VLPs may further comprise hemagglutinating activity. Plant or plant matter may be obtained by growing, harvesting or growing and harvesting the plant. The plant matter may comprise some or all of the plant, or one or more than one plant cell, leaves, stems, roots or cultured cells.

[0032] Without wishing to be bound by theory, plant-made VLPs comprising plant derived lipids, may induce a stronger immune reaction than VLPs made in other manufacturing systems and that the immune reaction induced by these plant-made VLPs is stronger when compared to the immune reaction induced by live or attenuated whole virus vaccines.

[0033] The composition of a protein extract obtained from a host cell is complex and typically comprises intercellular and intracellular components along with a protein or suprastructure of interest that is to be isolated. Preparation of an apoplastic fraction, followed by a step to segregate the intracellular proteins and components is advantageous since the protein or suprastructure of interest can be enriched and increase efficiency within a manufacturing process. Having a simpler process, comprising fewer efficient steps, may result in significant yield increases, and cost reduction. It has also been found that the process of digesting the cell wall using cell wall degrading enzymes increases VLP protein yield even if protoplasts do not remain intact during the extraction procedure. Without wishing to be bound by theory, the step of cell wall digestion may loosen the polymeric components of the cells wall and assist in release of the VLPs otherwise associated within the cell wall. This protocol may also minimize contamination of the VLPs within intracellular components.

[0034] Methods to digest plant cell-wall are known, and enzyme cocktail mixtures that digest cell walls may vary. The present invention is not limited by the cell wall digestion method used.

[0035] The methods described herein result in less disruption, and contamination of a plant-derived VLP extract when compared to methods for preparing plant-derived VLPs involving homogenization, blending or grinding. The methods described herein provide an apoplast fraction of the plant tissue and that may maintain the integrity of protoplasts and their components. The method as described herein is effective in purifying VLPs even if the protoplasts, or a portion of the protoplasts, lose their integrity and are no longer intact.

[0036] These methods provide a higher yield of VLPs when compared to methods of VLP extraction involving standard tissue disruption techniques, for example, homogenization, blending or grinding. The greater yield may be due to, in part, a reduction of the shearing forces that disrupt the structural integrity of the VLPs and/or the lipid envelope. Preparation of VLPs from an apoplastic fraction may be advantageous, as apoplastic fractions are significantly reduced, or free of, cytoplasmic proteins. Therefore, VLP separation from other proteins and matter, including HA monomers, trimers or fragments of HA, in the apoplastic fraction is easily

carried out. However, increased yields of VLPs may also be obtained using the methods described herein, even if the protoplast preparation, or a portion of the protoplast preparation, is not intact.

[0037] The VLPs of the present invention are also characterized as exhibiting a greater hemagglutinating activity than those obtained using standard tissue disruption techniques. This improved hemagglutinating activity may result from a greater yield of intact VLPs (fewer HA monomers or trimers free in solution), a greater yield of intact VLPs with intact lipid envelopes, or a combination thereof.

[0038] Vaccines made using VLPs provide the advantage, when compared to vaccines made of whole viruses, that they are non-infectious. Therefore, biological containment is not an issue and it is not required for production. Plant-made VLPs provide a further advantage by allowing the expression system to be grown in a greenhouse or field, thus being significantly more economical and suitable for scale-up.

[0039] Additionally, plants do not comprise enzymes involved in synthesizing and adding sialic acid residues to proteins. VLPs may be produced in the absence of neuraminidase (NA), and there is no need to co-express NA, or to treat the producing cells or extract with sialidase (neuraminidase), to ensure VLP production in plants

[0040] This summary of the invention does not necessarily describe all features of the invention.

BRIEF DESCRIPTION OF THE DRAWINGS

[0041] These and other features of the invention will become more apparent from the following description in which reference is made to the appended drawings wherein:

[0042] **Figure 1** shows a schematic representation of CPMVHT-based expression cassette (construct 685) for the expression of H5 A/Indonesia/5/05 hemagglutinin.

[0043] **Figure 2** shows A) the nucleic acid sequence (SEQ ID NO. 1) of a portion of construct for expressing H5/Indo (construct number 685) from PacI (upstream of the 35S promoter) to AscI (immediately downstream of the NOS terminator). Coding sequence of H5 from A/Indonesia/5/2005 is underlined. **Figure 2B** shows the amino acid sequence (SEQ ID NO. 2) of H5 A/Indonesia/5/05 hemagglutinin encoded by construct number 685.

[0044] **Figure 3** shows characterization of hemagglutinin (HA)-containing structures by size exclusion chromatography (SEC). Following centrifugation of the digested plant extract, the pellet was resuspended and fractionated by SEC. **Figure 3A** shows the total soluble protein content per fraction (solid triangles; % of maximum, left-side Y-axis; determined using the Bradford method). The hemagglutinating activity of the collected fractions (solid bars; right-side Y axis) is also shown. **Figure 3B** shows SDS-PAGE analysis of SEC eluted fractions. Fractions were precipitated by acetone and re-suspended in 1/40 volume of reducing sample loading buffer prior to analysis. Gel was stained with 0.1% Coomassie R-250 solution. Purified VLPs were run as a control. The band corresponding to the HA0 monomer is indicated by an arrow. MW - Molecular weight standards (kDa); C - Purified VLPs (control); lanes 7 through 10 and 14 through 16 correspond to fractions number eluted from SEC analysis, shown in Figure 3A.

[0045] **Figure 4** shows a comparison of protein profiles obtained after enzymatic digestion and by mechanical homogenization using a Comitrol™ homogenizer. Samples were treated in denaturing sample loading buffer and proteins were separated by SDS-PAGE analysis of elution fractions. Gels were stained with 0.1% Coomassie R-250 solution. MW - Molecular weight standards (kDa); lane 1 – 25 µl enzyme mixture; lane 2 – 25 µl enzymatic digestion of plant tissue and lane 3 – 5 µl extract obtained with the Comitrol homogenizer.

[0046] **Figure 5** shows the nucleic acid sequence (SEQ ID NO: 9) of an HA expression cassette comprising alfalfa plastocyanin promoter and 5' UTR, hemagglutinin coding sequence of H5 from A/Indonesia/5/2005 (Construct # 660), alfalfa plastocyanin 3' UTR and terminator sequences.

[0047] **Figure 6** shows the capture of HA-VLP on cationic exchange resin directly form separation of HA-VLP in the apoplastic fraction. Samples were treated in non-reducing, denaturing sample loading buffer and proteins were separated by SDS-PAGE. Gels were stained with 0.1% Coomassie R-250 solution. Lane 1: Apoplastic fraction after centrifugation, Lane 2-3: Apoplastic fraction after successive microfiltration; Lane 4: Load of the cationic exchange; Lane 5: Flow through fraction of the cationic exchange. Lane 6; elution from cationic exchange, concentrated 10X; Lane 7: Molecular weight standards (kDa).

[0048] **Figure 7** shows the Nanoparticle Tracking analysis (NTA) profile of H5/Indo VLP (A) and H1/Cal VLP (B) after clarification without addition of NaCl to digestion buffer and of H1/Cal VLP (C) with this addition. NTA experiments were carried out with NanoSight LM20

(NanoSight, Amesbury, UK). The instrument is equipped with a blue laser (405 nm), a sample chamber and a Viton fluoroelastomer o-ring. Videos were recorded at room temperature and analysed using the NTA 2.0 software. The samples were recorded for 60 sec. The shutter and gain were manually chosen so that optimal particle resolution was obtained.

[0049] **Figure 8** shows a Western blot of extract of H3/Brisbane VLP generated by enzymatic digestion using different buffers. Lane 1) Pure recombinant HA standard (5 µg, from Immune Technology Corp. IT-003-0042p) Lane 2 to 5 contain 7 µl of centrifuged enzymatic extract performed in the following buffers: Lane 2) 600mM Mannitol + 125mM citrate+ 75mM $NaPO_4$ + 25mM EDTA + 0.04% bisulfite pH6.2, Lane 3) 600mM Mannitol + 125mM citrate+ 75mM $NaPO_4$ + 50mM EDTA + 0.04% bisulfite pH6.2, Lane 4) 200mM Mannitol + 125mM citrate+ 75mM $NaPO_4$ + 25mM EDTA + 0.03% bisulfite pH6.2, Lane 5) 200mM Mannitol + 125mM citrate+ 75mM $NaPO_4$ + 50mM EDTA + 0.03% bisulfite pH6.2. The arrow represents the immunodetection signal of HA0.

DETAILED DESCRIPTION

[0050] The present invention relates to methods of preparing plant-derived virus-like particles (VLPs). More specifically, the present invention is directed to methods of preparing VLPs comprising influenza hemagglutinin (HA).

[0051] The following description is of a preferred embodiment.

[0052] The present invention provides a method for obtaining a protein, or protein suprastructure of interest. The protein of interest may be present in the apoplast or extracellular compartment, corresponding to the plant cell portion excluding the protoplast/spheroplast compartment. The method involves removing, digesting or both digesting and removing the cellulosic plant cell wall that surrounds plant cells. By digesting the cell wall the polymeric components of the cell wall are loosened, and the protein or protein suprastructure of interest may be more readily released. By using this method, the protein or protein suprastructure of interest is enriched since the protoplast/spheroplast compartment that contains a majorly host-cell proteins and components is segregated from the apoplast. As noted below, the method as provided herein is still effective in obtaining a protein or protein suprastructure of interest if, during the process, the integrity of the protoplast/spheroplast compartment is lost, if the protoplast/spheroplast compartment is not intact, and if a portion of host cell proteins and components from the protoplast/spheroplast compartment are present in the apoplast fraction.

[0053] Examples of protein suprastructures are structures comprised of two or more polypeptides; the polypeptides may be the same, or different; if different, they may be present in a ratio of about 1:1 to about 10:1 or greater. The protein suprastructure may further comprise one or more lipids, phospholipids, nucleic acids, membranes or the like. The two or more polypeptides may be connected by a covalent bond, a disulfide bridge, charge interaction, hydrophobic attraction, van der waals forces, hydrogen bonds or the like. An example of a protein suprastructure is a virus like particle (VLP), which may be enveloped, or non-enveloped, for example, a viral envelope protein, a viral structural protein, a viral capsid protein, or a viral coat protein.

[0054] The present invention also provides a method of preparing plant-derived virus like particles (VLPs). The method involves obtaining a plant or plant matter comprising plant-derived VLPs localized within the apoplast; producing a protoplast/spheroplast fraction, and an apoplast fraction from the plant matter, the apoplast fraction comprising plant-derived VLPs, and recovering the apoplast fraction. If desired, the plant derived VLPs may be purified from the apoplast fraction.

[0055] The present invention also provides a method of preparing VLPs comprising a plant-derived lipid envelope. The method includes obtaining a plant, or plant matter comprising VLPs, treating the plant or plant matter with an enzyme composition to produce one or more than one apoplastic protein complex and a protoplast/spheroplast fraction, and separating the one or more than one apoplastic protein complex from the protoplast fraction. The one or more than one apoplastic protein complex comprises the VLPs comprising a plant derived lipid envelope.

[0056] The present invention also provides a method of preparing plant derived VLPs, comprising obtaining a plant or plant matter that comprise the plant-derived VLPs, digesting the plant matter using a cell wall degrading enzyme composition to produced a digested fraction, and filtering the digested fraction to produced a filtered fraction and recovering the plant-derived VLPs from the filtered fraction. In this method, integrity of the protoplasts may not be required.

[0057] A protoplast is a plant cell that has had its cell wall completely or partially removed. A spheroplast may have partial removal of the cell wall. A protoplast, a spheroplast, or both a protoplast and spheroplast (protoplast/spheroplast) may be used as described herein, and the terms as used herein are interchangeable. The cell wall may be disrupted and removed mechanically (e.g. via homogenization, blending), the cell wall may be fully or partially digested

enzymatically, or the cell wall may be removed using a combination of mechanical and enzymatic methods, for example homogenization followed by treatment with enzymes for digestion of the cell wall. Protoplasts may also be obtained from cultured plant cells, for example liquid cultured plant cells, or solid cultured plant cells.

[0058] Standard reference works setting forth the general principles of plant tissue culture, cultured plant cells, and production of protoplasts, spheroplasts and the like include: *Introduction to Plant Tissue Culture*, by MK Razdan 2nd Ed. (Science Publishers, 2003; which is incorporated herein by reference), or see for example, the following URL: molecular-plant-biotechnology.info/plant-tissue-culture/protoplast-isolation.htm. Methods and techniques relating to protoplast (or spheroplast) production and manipulation are reviewed in, for example, Davey MR et al., 2005 (Biotechnology Advances 23:131-171; which is incorporated herein by reference). Standard reference works setting forth the general methods and principles of protein biochemistry, molecular biology and the like include, for example Ausubel et al, Current Protocols In Molecular Biology, John Wiley & Sons, New York (1998 and Supplements to 2001; which is incorporated herein by reference); Sambrook et al, Molecular Cloning: A Laboratory Manual, 2d Ed., Cold Spring Harbor Laboratory Press, Plainview, New York, 1989 (which is incorporated herein by reference); Kaufman et al , Eds., Handbook Of Molecular And Cellular Methods In Biology And Medicine, CRC Press, Boca Raton ,1995 (which is incorporated herein by reference); McPherson, Ed., Directed Mutagenesis: A Practical Approach, IRL Press, Oxford, 1991 (which is incorporated herein by reference).

[0059] Enzymes useful for digesting or degrading plant cell walls for release or protoplasts or spheroplasts are known to one of skill in the art and may include cellulase (EC 3.2.1.4), pectinase (EC 3.2.1.15), xylanase (EC 3.2.1.8), chitinases (EC 3.2.1.14), hemicellulase, or a combination thereof. Non- limiting examples of suitable enzymes includes a multi-component enzyme mixture comprising cellulase, hemicellulase, and pectinase, for example MACEROZYME™ (containing approximately: Cellulase: 0.1U/mg, Hemicellulase: 0.25U/mg, and Pectinase: 0.5U/mg). Other examples of commercial enzymes, enzyme mixtures and suppliers are listed in Table 1 (see: *Introduction to Plant Tissue Culture*, by MK Razdan 2nd Ed.., Science Publishers, 2003).

[0060] Alternate names, and types of cellulases include endo-1,4-β-D-glucanase; β-1,4-glucanase; β-1,4-endoglucan hydrolase; cellulase A; cellulosin AP; endoglucanase D; alkali

cellulase; cellulase A 3; celludextrinase; 9.5 cellulase; avicelase; pancellase SS and 1,4-(1,3;1,4)-β-D-glucan 4-glucanohydrolase. Alternate names, and types of pectinases (polygalacturonases) include pectin depolymerase; pectinase; endopolygalacturonase; pectolase; pectin hydrolase; pectin polygalacturonase; endo-polygalacturonase; poly-α-1,4-galacturonide glycanohydrolase; endogalacturonase; endo-D-galacturonase and poly(1,4-α-D-galacturonide) glycanohydrolase. Alternate names, and types of xylanases include hemicellulase, endo-(1→4)-β-xylan 4-xylanohydrolase; endo-1,4-xylanase; xylanase; β-1,4-xylanase; endo-1,4-xylanase; endo-β-1,4-xylanase; endo-1,4-β-D-xylanase; 1,4-β-xylan xylanohydrolase; β-xylanase; β-1,4-xylan xylanohydrolase; endo-1,4-β-xylanase; β-D-xylanase. Alternate names, and types of chitinases include chitodextrinase; 1,4-β-poly-*N*-acetylglucosaminidase; poly-β-glucosaminidase; β-1,4-poly-*N*-acetyl glucosamidinase; poly[1,4-(*N*-acetyl-β-D-glucosaminide)] glycanohydrolase.

Table 1: Non-limiting examples of commercially available enzymes for protoplast isolation

Enzyme	**Source**	**Supplier**
Cellulases		
Cellulase ONOZUKA R-10	*Trichoderma viride*	Kinki Yakult Mfg. Col. Ltd. 8-12, Shinglkancho Nishinomiya, Japan
Cellulase ONOZUKA RS	*T. viride*	Yakult Honsha Co., Tokyo, Japan
Cellulase YC	*T. viride*	Seishin Pharma Co. Ltd. 9-500-1, Nagareyama Nagareyama-shi, Chiba-kan, Japan
Cellulase CEL	*T. viride*	Cooper Biomedical Inc. Malvern, PA, USA
Cellulysin	*T. viride*	Calbiochem, San Diego, CA, USA
Driselase	*Irpex locteus*	Kyowa Hakko Kogyo Co. Ltd., Tokyo, Japan
Melcelase P-1	*T. viride*	Meiji Seiki Kaisha Ltd. No.8, 2-Chome Kyobashi, Chou-Ku, Japan
Multifect CX GC	*T. viride*	Genencor
Multifect CX B	*T. viride*	Genencor
Hemicellulases		
Hellcase	*Helix pomatia*	Industrie Biologique Francaise, Gennevilliers, France
Hemicellulase	*Aspergillus niger*	Sigma Chemical Co., St. Louis, MO, USA
Hemicellulase H-2125	*Rhizopus* sp.	Sigma, Munchen
Rhozyme HP 150	*Aspergillus niger*	Genencor Inc., South San Francisco, CA, USA
Pectinases		
MACERASE	*Rhizopus arrhizus*	Calbiochem, San Diego, CA, USA

MACEROZYME R-10	*R. arrhizus*	Yakult Honsha Co., Tokyo, Japan
Multifect Pectinase FE	*A. niger*	Genencor
PATE	*Baccilus polymyza*	Farbwerke-Hoechst AG, Frankfurt, FRG
Pectinol	*Aspergillus* sp.	Rohm and Haas Co. Independence Hall West, Philadelphia, PA 19105, USA
Pectolyase Y-23	*Aspergillus joponicus*	Seishin Pharma Co. Ltd., Japan
Zymolyase	*Arthrobacter luteus*	Sigma Chemical Co., USA

[0061] Choice of a particular enzyme or combination of enzymes, and concentration and reaction conditions may depend on the type of plant tissue used from which the protoplast and apoplast fraction comprising the VLPs is obtained. A mixture of cellulase, hemicellulase and pectinase, for example, a pectinase MACEROZYME™ or Multifect, may be used in a concentration ranging from 0.01% to2.5% (v/v), for example 0.01, 0.02, 0.04, 0.06, 0.08, 0.1, 0.2, 0.3, 0.4, 0.5, 0.6, 0.7, 0.8, 0.9, 1.0, 1.1,.1.2, 1.3, 1.4, 1.5, 1.6, 1.7, 1.8, 1.9, 2.0, 2.1, 2.2, 2.3, 2.4, or 2.5% (v/v), or any amount therebetween. MACEROZYME™ or Multifect may be used alone, or in combination with other enzymes, e.g cellulase, pectinase, hemicellulase, or a combination thereof. Cellulase may be used in a concentration ranging from 0.1% to 5%, for example 0.1, 0.25, 0.5, 0.75, 1.0, 1.25, 1.5, 1.75, 2.0, 2.25, 2.5, 2.75. 3.0. 3.25, 3.5, 3.75, 4.0, 4.25, 4.5, 4.75, 5.0% (w/v) or any amount therebetween.

[0062] The enzyme solution (alternately referred to as a cell wall degrading composition, digesting solution) will generally comprise a buffer or buffer system, an osmoticum, and one or more than one salts, divalent cations or other additives. The buffer or buffer system is selected to maintain a pH in the range suitable for enzyme activity and the stability of the protein(s), or VLP, to purify, for example, within the range of about pH 5.0 to about 8.0, or any value therebetween. The selected pH used may vary depending upon the VLP to be recovered, for example the pH may be 5.0, 5.2, 5.4, 5.6, 5.8, 6.0, 6.2, 6.4, 6.6, 6.8, 7.0, 7.2, 7.4, 7.6, 7.8, 8.0, or any pH therebetween. Examples of buffers or buffer systems include, but are not limited to, MES, phosphate, citrate and the like. One or more buffers or buffer systems may be combined in an enzyme solution (digesting solution); the one or more buffers may be present at a concentration from 0 mM to about 200 mM, or any amount therebetween, for example 10, 20, 30, 40, 50, 60, 70, 80, 90, 100, 110, 120, 130, 140, 150, 160, 170, 180 or 190 mM or any amount therebetween.

Depending on the suitability, an osmoticum component can be added if desired. The osmoticum and its concentration are selected to raise the osmotic strength of the enzyme solution. Examples of osmoticum include mannitol, sorbitol or other sugar alcohols, polyethylene glycol (PEG) of varying polymer lengths, and the like. Concentration ranges of osmoticum may vary depending on the plant species, the type of osmoticum used, and the type of plant tissue selected (species or organ of origin e.g. leaf or stem) – generally the range is from 0M to about 0.8 M, for example 0.05, 0.1, 0.15, 0.2, 0.25, 0.3. 0.35, 0.4, 0.5, 0.6, 0.7, or 0.75 M, or any amount therebetween, for example, 0, 50, 100, 150, 200, 250, 300, 350, 400, 450, 500, 550, 600 nM mannitol, or any amount therebetween. The concentration of osmoticum may also be expressed as a percentage (w/v). For some plant or tissue types, it may be beneficial to employ a slightly hypertonic preparation, which may facilitate separation of plant cell plasma membrane from the cell wall. The osmoticum can also be omitted during digestion.

[0063] Another parameter to set for the plant digestion is the temperature. Temperature may be controlled if desired during the digestion process. Useful temperature range should be between 4°C and 40 °C or any temperature therebetween, for example from about 4°C to about 15°C, or any amount therebetween, or from about 4°C to about 22°C, or any temperature therebetween. Depending to the temperature chosen, the other digestion experimental parameters may be adjusted to maintain optimal extraction conditions.

[0064] Cations, salts or both may be added to improve plasma membrane stability, for example divalent cations, such as Ca^{2+}, or Mg^{2+}, at 0.5-50mM, or any amount therebetween, salts, for example $CaCl_2$, NaCl, $CuSO_4$, KNO_3, and the like, from about 0 to about 750 mM, or any amount therebetween, for example 10, 20, 30, 40, 50, 100, 200, 300, 400, 500, 600, 700 or 750 mM. Other additives may also be added including a chelator for example, but not limited to, EDTA, EGTA, from about 0 to about 200 mM, or any amount therebetween, for example 5, 10, 15, 20, 25, 50, 75, 100, 125, 150, 175, 200 mM, or any amount therebetween, a reducing agent to prevent oxidation such as, but not limited to, sodium bisulfite or ascorbic acid, at 0.005-0.4% or any amount therebetween, for example 0.01, 0.02, 0.03, 0.04, 0.05, 0.06, 0.07, 0.08, 0.09, 0.1, 0.15, 0.2, 0,25, 0.3, 0.35, 0.4%, or any amount therebetween, specific enzyme inhibitors (see below), and if desired, an inhibitor of foliar senescence, for example, cycloheximide, kinetin, or one or more polyamines.

[0065] The digestion solution may also comprise one or more of mannitol from about 0 to about 600 mM, NaCl from about 0 to about 500 mM, EDTA from about 0 to about 50 mM, cellulose from about 1% to about 2% v/v, pectinase from about 0 to about 1% v/v, sodium metabisulfite from about 0.03 to about 0.04%, citrate from about 0 to about 125 mM or $NaPO_4$ from about 0 to 75 mM.

[0066] The plant matter may be treated to enhance access of the enzymes or enzyme composition to the plant cell wall. For example, the epidermis of the leaf may be removed or 'peeled' before treatment with an enzyme composition. The plant matter may be cut into small pieces (manually, or with a shredding or cutting device such as an Urschel slicer); the cut up plant matter may be further infiltrated with an enzyme composition under a partial vacuum (Nishimura and Beevers 1978, Plant Physiol 62:40-43; Newell et al., 1998, J. Exp Botany 49:817-827). Mechanical perturbation of the plant matter may also be applied to the plant tissues (Giridhar et al., 1989. Protoplasma 151:151-157) before or during treatment with an enzyme composition. Furthermore, cultured plant cells, either liquid or solid cultures, may be used to prepare protoplasts or spheroplasts.

[0067] It may be desired to use an enzyme composition that lacks, or that has inactivated lipases or proteases. In some embodiments, one or more protease, or lipase inhibitors may be included in the enzyme composition. Examples of lipase inhibitors include RHC80267 (SigmaAldrich); examples of protease inhibitors include E-64, Na_2EDTA, Pepstatin, aprotinin, PMSF, Pefabloc, Leupeptin, bestatin and the like.

[0068] Any suitable method of mixing or agitating the plant matter in the enzyme composition may be used. For example, the plant matter may be gently swirled or shaken in a tray or pan or via a rotary shaker, tumbled in a rotating or oscillating drum. Precaution should be taken in order to minimize the protoplast (and/or spheroplast) damage until they are removed form the digestion soup. The digestion vessel should be selected accordingly.

[0069] As a non-limiting example, an enzyme composition comprising 1.5% cellulase (Onozuka R-10) and 0.375% MACEROZYME^{TM} in 500 mM mannitol, 10 m $CaCl_2$ and 5 mM MES (pH 5.6) may be used for protoplast (or spheroplast) production from some *Nicotiana* tissues. As described herein, the concentration of mannitol may also be varied from about 0 to about 500mM, or any amount therebetween. One of skill in the art, provided with the information

disclosed herein, will be able to determine a suitable enzyme composition for the age and strain of the *Nicotiana sp*, or for another species used for production of VLPs.

[0070] Upon disruption of the cell wall, or partial digestion of the cell wall, a protoplast fraction (comprising protoplasts and/or spheroplasts), and an "apoplast fraction" are obtained. Alternatively, a "digested fraction" may be obtained. As noted below, integrity of the protoplast fraction may not be required to produce high yields of protein as described herein, therefore, an apoplast fraction or a digested fraction may be used for the extraction of proteins, for example, but not limited to, VLPs, viral envelope proteins, viral structural proteins, viral capsid proteins, viral coat proteins.

[0071] By "apoplast fraction" it is meant a fraction that is obtained following enzymatic digestion, or partial enzymatic digestion, using cell wall degrading enzymes of the plant matter in the presence of an osmoticum and/or other ingredients that may be used to assist in maintaining integrity of the protoplast. The apoplast fraction may comprise some components arising from disrupted protoplasts (or spheroplasts). For example, the apoplast fraction may comprise from about 0 to about 50% (v/v) or any amount therebetween, of the components from the protoplast fraction, or 0, 0.1, 0.5, 1, 5, 10, 15, 20, 25, 30, 35, 40, 45, or 50% (v/v) or any amount therebetween of the components from the protoplast fraction.

[0072] By a "digested fraction" it is meant the fraction that remains following enzymatic digestion, or partial enzymatic digestion, using cell wall degrading enzymes of the plant matter, however, integrity of the protoplast is not required, and the digested fraction may comprise intact, disrupted, or both intact and disrupted protoplasts. The composition comprising the cell wall degrading enzymes used to produce the digested fraction may comprise an osmoticum, or the osmoticum may be present at a reduced amount when compared to the amount present in standard procedures used to obtain protoplasts, or the osmoticum may be absent from the composition. The digested fraction comprises the apoplast fraction and the protoplast/spheroplast fraction, however, the protoplast/spheroplast fraction may or may not be intact. The digested fraction contains intracellular components and extracellular components. Intracellular components may be found in the form of protoplasts/spheroplasts if an osmoticum is used to maintain the protoplast/spheroplast intact. If no osmoticum is used in the digestion solution, then the protoplasts/spheroplasts may be disrupted and the intracellular and extracellular components may be combined in the digested fraction. As described herein , the

VLPs, may be separated from components of the digested fraction using any suitable technique. Without wishing to be bound by theory, the step of cell wall digestion may loosen the polymeric components of the cells wall and assist in release VLPs, otherwise trapped within the cell wall. This protocol also minimizes contamination of theVLPs, with the intracellular components. The VLPs may be separated from cellular debris following enzymatic digestion using low speed centrifugation followed by filtration, depth filtration, sedimentation, precipitation for example , but not limited to ammonium sulfate precipitation, or a combination thereof to obtain a separated fraction comprising the proteins or suprastructure proteins of interest.

[0073] If an osmoticum is used, the protoplast/spheroplast fraction, or fraction comprising protoplasts, may be separated from the apoplast fraction using any suitable technique, for example but not limited to, centrifugation, filtration, depth filtration, sedimentation, precipitation, or a combination thereof to obtain a separated fraction comprising the VLPs and/or comprising protoplasts/spheroplasts that comprise the VLPs.

[0074] The protoplast (and spheroplast) fraction, or fraction comprising protoplasts, may be separated from the apoplast fraction using any suitable technique, for example but not limited to, centrifugation, filtration, depth filtration, sedimentation, precipitation, or a combination thereof to obtain a separated fraction.

[0075] The separated fraction may be for example a supernatant (if centrifuged, sedimented, or precipitated), or a filtrate (if filtered), and is enriched for VLPs. The separated fraction may be further processed to isolate, purify, concentrate or a combination thereof, the VLPs by, for example, additional centrifugation steps, precipitation, chromatographic steps (e.g. size exclusion, ion exchange chromatography), tangential flow filtration, or a combination thereof. The presence of purified VLPs may be confirmed by, for example, native or SDS-PAGE, Western analysis using an appropriate detection antibody, capillary electrophoresis, or any other method as would be evident to one of skill in the art.

[0076] The apoplast is the portion of the plant cell outside the plasma membrane, and includes the cell wall and intercellular spaces of the plant. While it is preferred that the integrity of the protoplasts (and/or spheroplasts) be maintained during digestion and further processing, it is not required that the protoplasts remain intact in order to enrich for VLPs.

[0077] During synthesis, VLPs are excreted outside of the plasma membrane. VLPs are of an average size of about 20 nm to 1 μm, or any amount therebetween, for example 60, 65, 70, 75, 80, 85, 90, 95, 100, 105, 110, 120, 130, 140, 150 160, 170, 180, 190, or 200 nm, or any amount therebetween, for example 100 nm, and may include a lipid membrane. Due to their size, once synthesized, VLPs may remain trapped between the plasma membrane and cell wall and may be inaccessible for isolation or further purification using standard mechanical methods used to obtain plant proteins. In order to maximize yields, minimize contamination of the VLP fraction with cellular proteins, maintain the integrity of the VLPs and, in some embodiments, the associated lipid envelope or membrane, methods of disrupting the cell wall to release the VLPs that minimize mechanical damage to the protoplast (and/or spheroplasts) may be useful, such as the enzymatic methods described herein. However, it is not required that the integrity of all of the protoplasts be retained during the procedure.

[0078] A VLP produced in a plant according to some aspects of the invention may be complexed with plant-derived lipids. The VLP may comprise an HA0 precursor form, or the HA1 or HA2 domains retained together by disulphide bridges form. The plant-derived lipids may be in the form of a lipid bilayer, and may further comprise an envelope surrounding the VLP. The plant derived lipids may comprise lipid components of the plasma membrane of the plant where the VLP is produced, including, but not limited to, phosphatidylcholine (PC), phosphatidylethanolamine (PE), glycosphingolipids, phytosterols or a combination thereof. A plant-derived lipid may alternately be referred to as a 'plant lipid'. Examples of phytosterols are known in the art, and include, for example, stigmasterol, sitosterol, 24-methylcholesterol and cholesterol (Mongrand et al., 2004, J. Biol Chem 279:36277-86).

[0079] Correct folding of viral structural proteins such as HA, and formation of trimers of HA is desired for assembly of VLPs. VLPs, and in particular VLPs comprising a plant derived lipid envelope, may provide for a superior immune response when administered to a subject, relative to administration of the structural protein monomer.

[0080] In some embodiments, polypeptide expression may be targeted to any intracellular or extracellular space, organelle or tissue of a plant. In order to localize the expressed polypeptide to a particular location, the nucleic acid encoding the polypeptide may be linked to a nucleic acid sequence encoding a signal peptide. A signal peptide may alternately be referred to as a transit peptide or signal sequence. Signal peptides or peptide sequences for directing localization of an

expressed polypeptide to the apoplast include, but are not limited to, a rice amylase signal peptide (McCormick 1999, Proc Natl Acad Sci USA 96:703-708), protein disulfide isomerase signal peptide (PDI) having the amino acid sequence:

MAKNVAIFGLLFSLLLLVPSQIFAEE; SEQ ID NO. 10,

plant pathogenesis related protein (PRP; Szyperski et al. PNAS 95:2262-2262), for example, Tobacco plant pathogenesis related protein 2 (PRP), human monoclonal antibody signal peptide (SP), or any native hemagglutinin signal peptide.

[0081] In some examples, an expressed polypeptide may accumulate in specific intercellular or extracellular space (such as the apoplast), organelle or tissue, for example when the polypeptide is expressed and secreted in the absence of a signal peptide or transit peptide.

[0082] The term "virus like particle" (VLP), or "virus-like particles" or "VLPs" refers to structures that self-assemble and comprise viral surface proteins, for example an influenza HA protein, or a chimeric influenza HA protein. VLPs and chimeric VLPs are generally morphologically and antigenically similar to virions produced in an infection, but lack genetic information sufficient to replicate and thus are non-infectious. VLPs and chimeric VLPs may be produced in suitable host cells including plant host cells, and if desired further purified.

[0083] While influenza VLPs and chimeric influenza VLPs are exemplified herein, the methods described herein may be used for any plant-derived VLPs that localize in, or are secreted to, the apoplast.

[0084] By "chimeric protein" or "chimeric polypeptide", it is meant a protein or polypeptide that comprises amino acid sequences from two or more than two sources, for example but not limited to, two or more influenza types or subtypes, that are fused as a single polypeptide. The chimeric protein or polypeptide may include a signal peptide that is the same (i.e. native) as, or heterologous with, the remainder of the polypeptide or protein. The chimeric protein or chimeric polypeptide may be produced as a transcript from a chimeric nucleotide sequence, and remain intact, or if required, the chimeric protein or chimeric polypeptide may be cleaved following synthesis. The intact chimeric protein, or cleaved portions of the chimeric protein, may associate to form a multimeric protein. A chimeric protein or a chimeric polypeptide may also include a protein or polypeptide comprising subunits that are associated via disulphide bridges (i.e. a multimeric protein). For example, a chimeric polypeptide comprising amino acid sequences

from two or more than two sources may be processed into subunits, and the subunits associated via disulphide bridges to produce a chimeric protein or chimeric polypeptide.

[0085] The polypeptide may be influenza hemagglutinin (HA), and each of the two or more than two amino acid sequences that make up the polypeptide may be obtained from different HA's to produce a chimeric HA, or chimeric influenza HA. A chimeric HA may also include a amino acid sequence comprising heterologous signal peptide (a chimeric HA pre-protein) that is cleaved after synthesis. Examples of HA proteins that may be used in the invention described herein may be found in WO 2009/009876; WO 2009/076778; WO 2010/003225 (which are incorporated herein by reference). A nucleic acid encoding a chimeric polypeptide may be described as a "chimeric nucleic acid", or a "chimeric nucleotide sequence". A virus-like particle comprised of chimeric HA may be described as a "chimeric VLP". Chimeric VLPs are further described in PCT Application No. PCT/CA2010/000983 filed June 25, 2010, and U.S. Provisional Application No. 61/220,161 (filed June 24, 2009; which is incorporated herein by reference). VLPs can be obtained from expression of native or chimeric HA.

[0086] The HA of the VLPs prepared according to a method provided by the present invention, include known sequences and variant HA sequences that may be developed or identified. Furthermore, VLPs produced as described herein do not comprise neuraminidase (NA) or other components for example M1 (M protein), M2, NS and the like. However, NA and M1 may be co-expressed with HA should VLPs comprising HA and NA be desired.

[0087] Generally, the term "lipid" refers to a fat-soluble (lipophilic), naturally-occurring molecules. A chimeric VLP produced in a plant according to some aspects of the invention may be complexed with plant-derived lipids. The plant-derived lipids may be in the form of a lipid bilayer, and may further comprise an envelope surrounding the VLP. The plant derived lipids may comprise lipid components of the plasma membrane of the plant where the VLP is produced, including phospholipids, tri-, di- and monoglycerides, as well as fat-soluble sterol or metabolites comprising sterols. Examples include phosphatidylcholine (PC), phosphatidylethanolamine (PE), phosphatidylinositol, phosphatidylserine, glycosphingolipids, phytosterols or a combination thereof. A plant-derived lipid may alternately be referred to as a 'plant lipid'. Examples of phytosterols include campesterol, stigmasterol, ergosterol, brassicasterol, delta-7-stigmasterol, delta-7-avenasterol, daunosterol, sitosterol, 24-methylcholesterol, cholesterol or beta-sitosterol (Mongrand et al., 2004, J. Biol Chem

279:36277-86). As one of skill in the art will readily understand, the lipid composition of the plasma membrane of a cell may vary with the culture or growth conditions of the cell or organism, or species, from which the cell is obtained.

[0088] Cell membranes generally comprise lipid bilayers, as well as proteins for various functions. Localized concentrations of particular lipids may be found in the lipid bilayer, referred to as 'lipid rafts'. These lipid raft microdomains may be enriched in sphingolipids and sterols. Without wishing to be bound by theory, lipid rafts may have significant roles in endo and exocytosis, entry or egress of viruses or other infectious agents, inter-cell signal transduction, interaction with other structural components of the cell or organism, such as intracellular and extracellular matrices.

[0089] VLPs comprising a lipid envelope has been previously described in WO 2009/009876; WO 2009/076778, and WO 2010/003225 (which are incorporated herein by reference). With reference to influenza virus, the term "hemagglutinin" or "HA" as used herein refers to a structural glycoprotein of influenza viral particles. The HA of the present invention may be obtained from any subtype. For example, the HA may be of subtype H1, H2, H3, H4, H5, H6, H7, H8, H9, H10, H11, H12, H13, H14, H15, or H16, or of influenza types B or C. The recombinant HA of the present invention may also comprise an amino acid sequence based on the sequence of any hemagglutinin. The structure of influenza hemagglutinin is well-studied and demonstrates a high degree of conservation in secondary, tertiary and quaternary structure. This structural conservation is observed even though the amino acid sequence may vary (see, for example, Skehel and Wiley, 2000 Ann Rev Biochem 69:531-69; Vaccaro et al 2005; which is incorporated herein by reference). Nucleotide sequences encoding HA are well known, and are available for example, from the BioDefense and Public Health Database (now Influenza Research Database; Squires et al., 2008 Nucleic Acids Research 36:D497-D503) for example at URL: biohealthbase.org/GSearch/home.do?decorator=Influenza) or the databases maintained by the National Center for Biotechnology Information (NCBI; for example at URL: ncbi.nlm.nih.gov/sites/entrez?db=nuccore&cmd=search&term=influenza), both of which are incorporated herein by reference.

[0090] The present invention also pertains to methods of preparing, isolating, or both preparing and isolating VLPs, including influenza VLPs of viruses which infect humans, or host animals, for example primates, horses, pigs, birds, sheep, avian water fowl, migratory birds, quail, duck,

geese, poultry, chicken, camel, canine, dogs, feline, cats, tiger, leopard, civet, mink, stone marten, ferrets, house pets, livestock, mice, rats, seal, whale and the like. Some influenza viruses may infect more than one host animal.

[0091] Amino acid variation is tolerated in hemagglutinins of influenza viruses. This variation provides for new strains that are continually being identified. Infectivity between the new strains may vary. However, formation of hemagglutinin trimers, which subsequently form VLPs is maintained. The present invention also includes methods of preparing any plant-derived VLPs, regardless of the HA subtype or sequence, or chimeric HA comprising the VLP, or species of origin.

[0092] Correct folding of the hemagglutinins may be important for stability of the protein, formation of multimers, formation of VLPs and function of the HA (ability to hemagglutinate), among other characteristics of influenza hemagglutinins. Folding of a protein may be influenced by one or more factors, including, but not limited to, the sequence of the protein, the relative abundance of the protein, the degree of intracellular crowding, the availability of cofactors that may bind or be transiently associated with the folded, partially folded or unfolded protein, the presence of one or more chaperone proteins, or the like.

[0093] Heat shock proteins (Hsp) or stress proteins are examples of chaperone proteins, which may participate in various cellular processes including protein synthesis, intracellular trafficking, prevention of misfolding, prevention of protein aggregation, assembly and disassembly of protein complexes, protein folding, and protein disaggregation. Examples of such chaperone proteins include, but are not limited to, Hsp60, Hsp65, Hsp 70, Hsp90, Hsp100, Hsp20-30, Hsp10, Hsp100-200, Hsp100, Hsp90, Lon, TF55, FKBPs, cyclophilins, ClpP, GrpE, ubiquitin, calnexin, and protein disulfide isomerases (see, for example, Macario, A.J.L., *Cold Spring Harbor Laboratory Res.* 25:59-70. 1995; Parsell, D.A. & Lindquist, S. *Ann. Rev. Genet.* 27:437-496 (1993); U.S. Patent No. 5,232,833). Chaperone proteins, for example but not limited to Hsp40 and Hsp70 may be used to ensure folding of a chimeric HA (PCT Application No. PCT/CA2010/000983 filed June 25, 2010, and U.S. Provisional Application No. 61/220,161, filed June 24, 2009; WO 2009/009876 and WO 2009/076778, all of which are incorporated herein by reference). Protein disulfide isomerase (PDI; Accession No. Z11499) may also be used.

[0094] Once recovered,VLPs may be assessed for structure, size potency or activity by, for example, hemagglutination assay, electron microscopy, light scattering, size exclusion chromatography, HPLC, Western blot analysis, or electrophoresis. These and other methods for assessing size, concentration, activity and composition of VLPs are known in the art.

[0095] For preparative size exclusion chromatography, a preparation comprising VLPs may be obtained by the methods described herein, and insoluble material removed by centrifugation. Precipitation with PEG may also be of benefit. The recovered protein may be quantified using conventional methods (for example, Bradford Assay, BCA), and the extract passed through a size exclusion column, using for example SEPHACRYL™, SEPHADEX™, or similar medium, and the fractions collected. Blue Dextran 2000 or a suitable protein, may be used as a calibration standard. The extract may also be passed through a cation exchange column and active fractions collected. Following chromatography, fractions may be further analyzed by protein electrophoresis, immunoblot, or both, to confirm the presence of VLPs and the protein complement of the fraction.

[0096] A hemagglutination assay may be used to assess the hemagglutinating activity of the VLP-containing fractions, using methods well-known in the art. Without wishing to be bound by theory, the capacity of HA to bind to RBC from different animals is driven by the affinity of HA for sialic acids α2,3 or α2,3 and the presence of these sialic acids on the surface of RBC. Equine and avian HA from influenza viruses agglutinate erythrocytes from all several species, including turkeys, chickens, ducks, guinea pigs, humans, sheep, horses and cows; whereas human HAs will bind to erythrocytes of turkey, chickens, ducks, guinea pigs, humans and sheep (Ito T. et al, 1997, Virology, 227:493-499; Medeiros R et al, 2001. Virology 289:74-85).

[0097] A hemagglutination inhibition (HI, or HAI) assay may also be used to demonstrate the efficacy of antibodies induced by a vaccine, or vaccine composition comprising chimeric HA or chimeric VLP can inhibit the agglutination of red blood cells (RBC) by recombinant HA. Hemagglutination inhibitory antibody titers of serum samples may be evaluated by microtiter HAI (Aymard et al 1973). Erythrocytes from any of several species may be used – e.g. horse, turkey, chicken or the like. This assay gives indirect information on assembly of the HA trimer on the surface of VLP, confirming the proper presentation of antigenic sites on HAs.

[0098] Cross-reactivity HAI titres may also be used to demonstrate the efficacy of an immune response to other strains of virus related to the vaccine subtype. For example, serum from a

subject immunized with a vaccine composition comprising a chimeric hemagglutinin comprising an HDC of a first influenza type or subtype may be used in an HAI assay with a second strain of whole virus or virus particles , and the HAI titer determined.

[0099] Methods for transformation, and regeneration of transgenic plants, plant cells, plant matter or seeds comprising VLPs are established in the art and known to one of skill in the art. The method of obtaining transformed and regenerated plants is not critical to the present invention.

[00100] By "transformation" it is meant the interspecific transfer of genetic information (nucleotide sequence) that is manifested genotypically, phenotypically or both. The interspecific transfer of genetic information from a chimeric construct to a host may be heritable (i.e. integrated within the genome of the host) and the transfer of genetic information considered stable, or the transfer may be transient and the transfer of genetic information is not inheritable.

[00101] By the term "plant matter", it is meant any material derived from a plant. Plant matter may comprise an entire plant, tissue, cells, or any fraction thereof. Further, plant matter may comprise intracellular plant components, extracellular plant components, liquid or solid extracts of plants, or a combination thereof. Further, plant matter may comprise plants, plant cells, tissue, a liquid extract, or a combination thereof, from plant leaves, stems, fruit, roots or a combination thereof. Plant matter may comprise a plant or portion thereof which has not been subjected to any processing steps. A portion of a plant may comprise plant matter. Plants or plant matter may be harvested or obtained by any method, for example, the whole plant may be used, or the leaves or other tissues specifically removed for use in the described methods. Transgenic plants expressing and secreting VLPs may also be used as a starting material for processing as described herein.

[00102] The constructs of the present invention can be introduced into plant cells using Ti plasmids, Ri plasmids, plant virus vectors, direct DNA transformation, micro-injection, electroporation, infiltration, and the like. For reviews of such techniques see for example Weissbach and Weissbach, *Methods for Plant Molecular Biology*, Academy Press, New York VIII, pp. 421-463 (1988); Geierson and Corey, *Plant Molecular Biology*, 2d Ed. (1988); and Miki and Iyer, Fundamentals of Gene Transfer in Plants. In *Plant Metabolism*, 2d Ed. DT. Dennis, DH Turpin, DD Lefebrve, DB Layzell (eds), Addison-Wesley, Langmans Ltd. London, pp. 561-579 (1997). Other methods include direct DNA uptake, the use of liposomes, electroporation, for

example using protoplasts, micro-injection, microprojectiles or whiskers, and vacuum infiltration. See, for example, Bilang, et al. (Gene 100: 247-250 (1991), Scheid et al. (Mol. Gen. Genet. 228: 104-112, 1991), Guerche et al. (Plant Science 52: 111-116, 1987), Neuhause et al. (Theor. Appl Genet. 75: 30-36, 1987), Klein et al., Nature 327: 70-73 (1987); Howell et al. (Science 208: 1265, 1980), Horsch et al. (Science 227: 1229-1231, 1985), DeBlock et al., Plant Physiology 91: 694-701, 1989), Methods for Plant Molecular Biology (Weissbach and Weissbach, eds., Academic Press Inc., 1988), Methods in Plant Molecular Biology (Schuler and Zielinski, eds., Academic Press Inc., 1989), Liu and Lomonossoff (J. Virol Meth, 105:343-348, 2002), U.S. Pat. Nos. 4,945,050; 5,036,006; 5,100,792; 6,403,865; 5,625,136, (all of which are hereby incorporated by reference).

[00103] Transient expression methods may be used to express the constructs of the present invention (see Liu and Lomonossoff, 2002, Journal of Virological Methods, 105:343-348; which is incorporated herein by reference). Alternatively, a vacuum-based transient expression method, as described in PCT Publications WO 00/063400, WO 00/037663 (incorporated herein by reference) may be used. These methods may include, for example, but are not limited to, a method of Agro-inoculation or Agro-infiltration, however, other transient methods may also be used as noted above. With either Agro-inoculation or Agro-infiltration, a mixture of *Agrobacteria* comprising the desired nucleic acid enter the intercellular spaces of a tissue, for example the leaves, aerial portion of the plant (including stem, leaves and flower), other portion of the plant (stem, root, flower), or the whole plant. After crossing the epidermis the *Agrobacterium* infect and transfer t-DNA copies into the cells. The t-DNA is episomally transcribed and the mRNA translated, leading to the production of the protein of interest in infected cells, however, the passage of t-DNA inside the nucleus is transient.

[00104] The influenza VLPs prepared by methods of the present invention may be used in conjunction with an existing influenza vaccine, to supplement the vaccine, render it more efficacious, or to reduce the administration dosages necessary. As would be known to a person of skill in the art, the vaccine may be directed against one or more than one influenza virus. Examples of suitable vaccines include, but are not limited to, those commercially available from Sanofi-Pasteur, ID Biomedical, Merial, Sinovac, Chiron, Roche, MedImmune, GlaxoSmithKline, Novartis, Sanofi-Aventis, Serono, Shire Pharmaceuticals and the like.

[00105] If desired, the VLPs of the present invention may be admixed with a suitable adjuvant as would be known to one of skill in the art. Furthermore, the VLP may be used in a vaccine composition comprising an effective dose of the VLP for the treatment of a target organism, as defined above. Furthermore, the VLP produced according to the present invention may be co-expressed with other protein components or reconstituted with other VLPs or influenza protein components, for example, neuraminidase (NA), M1, and M2, . It can also be co-expressed or reconstituted with other VLP made of vaccinal proteins such as malaria antigens, HIV antigens, respiratory syncytial virus (RSV) antigens, and the like.

[00106] The sequences described herein are summarized below.

SEQ ID NO:	**Description**	**Figure**
1	Nucleic acid sequence (construct 685)	2A
2	Amino acid sequence encoded by SEQ ID NO: 1	2B
3	**pBinPlus.2613c**: AGGAAGGGAAGAAAGCGAAAGGAG	
4	**Mut-ATG115.r**: GTGCCGAAGCACGATCTGACAACGT TGAAGATCGCTCACGCAAGAAAGACAAGAGA	
5	**Mut-ATG161.c**: GTTGTCAGATCGTGCTTCGGCACCAGTACAA CGTTTTCTTTCACTGAAGCGA	
6	**LC-C5-1.110r**: TCTCCTGGAGTCACAGACAGGGTGG	
7	**ApaI-H5 (A-Indo).1c**: TGTCGGGCCCATGGAGAAAATAGTGC TTCTTCTTGCAAT	
8	**H5 (A-Indo)-StuI.1707r**: AAATAGGCCTTTAAATGCAAATTC TGCATTGTAACGA	
9	nucleic acid sequence (construct 660)	5
10	PDI signal peptide: MAKNVAIFGLLFSLLLLVPSQIFAEE	
11	Plasto-443c	
12	supP19-plasto.r	
13	supP19-1c	
14	SupP19-SacI.r	

[00107] The present invention will be further illustrated in the following examples. However it is to be understood that these examples are for illustrative purposes only, and should not be used to limit the scope of the present invention in any manner.

Assembly of expression cassettes

[00108] Constructs that may be used for the production of VLPs are described U.S. Provisional Application No. 61/220,161 (filed June 24, 2009), WO 2009/009876, WO 2009/076778 and WO2010/003225 all of which are incorporated herein by reference. Constructs may also include those listed in Table 2. Assembly of these constructs is described in WO 2009/009876, WO 2009/076778, WO2010/003225 and US 61/220,161. However other constructs comprising known HA's, including but not limited to, those provided in Table 2, and combined with similar or different regulatory elements and promoters, may also be used for the production of VLPs as described herein.

Table 2: Non-limiting examples of constructs that can be used for hemagglutinin production.

Cassette number	Corresponding HA	HA abbreviation
540	SpPDI-H1 from strain A/New Caledonia/20/99 (H1N1)	H1/NC
560	SpPDI-H1 A/California/4/2009 in 2X35S/CPMV-*HT* expression cassette	H1/Cal WT
580	SpPDI-H1 A/New Caledonia/20/99 in 2x35S/CPMV-*HT* expression cassette	H1/NC
660	H5 from strain A/Indonesia/5/2005 (H5N1)	H1/Indo
663	H5 A/Indonesia/5/2005	H1/Indo
685	H5 A/Indonesia/5/2005 in CPMV-*HT* expression cassette	H1/Indo
686	SpPDI-H5 A/Indonesia/5/2005 in CPMV-*HT* expression cassette	H1/Indo
690	H1 A/Brisbane/59/07 receptor-binding (RB) domain in H5 A/Indonesia/5/05 backbone	H1/Bris
691	H1 A/Brisbane/59/07 esterase and receptor-binding domains (E1-RB-E2) in H5 A/Indonesia/5/05 backbone	H1/Bris
696	H5 A/Indonesia/5/05 receptor-binding (RB) domain in H1 A/New Caledonia/20/99 backbone	H1/Indo
732	H1 A/Brisbane/59/2007 in CPMV-*HT* expression cassette	H1/Bris
733	SpPDI-H1 A/Brisbane/59/2007 in CPMV-*HT* expression cassette	H1/Bris
734	H1 A/Brisbane/59/07 receptor-binding (RB) domain in H5 A/Indonesia/5/05 backbone in CPMV-HT expression cassette	H1/Bris
735	H3 A/Brisbane/10/2007 in CPMV-*HT* expression cassette	H3/Bris

736	SpPDI-H3 A/Brisbane/10/2007 in CPMV-*HT* expression cassette	H3/Bris
737	Assembly of chimeric SpPDI-H3 A/Brisbane/10/2007 (ectodomain) + H5 A/Indonesia/5/2005 (TmD + Cyto tail) in CPMV-*HT* expression cassette	H3/Bris-H5/Indo chimera
738	HA B/Florida/4/2006 in CPMV-*HT* expression cassette	B/Flo
739	SpPDI-HA B/Florida/4/2006 in CPMV-*HT* expression cassette	B/Flo
745	SpPDI-HA B/Florida/4/2006 (ectodomain) + H5 A/Indonesia/5/2005 (TmD + Cyto tail) in CPMV-*HT* expression cassette	B/Flo
747	SpPDI-HA B/Florida/4/2006+ H5 A/Indonesia/5/2005 (TmD + Cyto tail) in 2X35S-CPMV-*HT* expression cassette	B/Flo
774	HA of A/Brisbane/59/2007 (H1N1)	H1/Bris
775	HA of A/Solomon Islands 3/2006 (H1N1)	H1/Solomon
776	HA of A/Brisbane 10/2007 (H3N2)	H3/Bris
777	HA of A/Wisconsin/67/2005 (H3N2)	H3/Wisc
778	HA of B/Malaysia/2506/2004	B/Malaysia
779	HA of B/Florida/4/2006	B/Flo
780	HA of A/Singapore/1/57 (H2N2)	H2/Sing
781	HA of A/Anhui/1/2005 (H5N1)	H5/Anhui
782	HA of A/Vietnam/1194/2004 (H5N1)	H5/Vietnam
783	HA of A/Teal/HongKong/W312/97 (H6N1)	H6/HongKong
784	HA of A/Equine/Prague/56 (H7N7)	H7/Prague
785	HA of A/HongKong/1073/99 (H9N2)	H9/HongKong
787	H1 A/Brisbane/59/2007	H1/Bris
790	H3 A/Brisbane/10/2007	H3/Bris
798	HA B/Florida/4/2006	B/Flo

[00109] CPMV-*HT* expression cassettes included the 35S promoter to control the expression of an mRNA comprising a coding sequence of interest flanked, in 5' by nucleotides 1–512 from the Cowpea mosaic virus (CPMV) RNA2 with mutated ATG at positions 115 and 161 and in 3', by nucleotides 3330–3481 from the CPMV RNA2 (corresponding to the 3' UTR) followed by the NOS terminator. Plasmid pBD-C5-1LC, (Sainsbury et al. 2008; Plant Biotechnology Journal 6: 82-92 and PCT Publication WO 2007/135480), was used for the assembly of CPMV-*HT*-based hemagglutinin expression cassettes. The mutation of ATGs at position 115 and 161 of the CPMV RNA2 was done using a PCR-based ligation method presented in Darveau et al. (Methods in Neuroscience 26: 77-85 (1995)). Two separate PCRs were performed using pBD-C5-1LC as template. The primers for the first amplification were

pBinPlus.2613c (SEQ ID NO: 3) and Mut-ATG115.r (SEQ ID NO: 4). The primers for the second amplification were Mut-ATG161.c (SEQ ID NO: 5) and LC-C5-1.110r (SEQ ID NO: 6). The two fragments were then mixed and used as template for a third amplification using pBinPlus.2613c (SEQ ID NO: 3) and LC-C5-1.110r (SEQ ID NO: 6) as primers. The resulting fragment was digested with PacI and ApaI and cloned into pBD-C5-1LC digested with the same enzyme. The expression cassette generated was named 828.

Assembly of H5 A/Indonesia/5/2005 in CPMV-*HT* expression cassette (construct number 685).

[00110] The assembly of this cassette is described in WO 2009/009876, WO 2009/076778 and WO2010/003325, which are incorporated herein by reference.

[00111] Briefly, the coding sequence of H5 from A/Indonesia/5/2005 was cloned into CPMV-*HT* as follows: restriction sites ApaI (immediately upstream of the initial ATG) and StuI (immediately downstream of a stop codon) were added to the hemagglutinin coding sequence by performing a PCR amplification with primers ApaI-H5 (A-Indo).1c (SEQ ID NO: 7) and H5 (A-Indo)-StuI.1707r (SEQ ID NO: 8) using construct number 660 (D'Aoust et al., Plant Biotechnology Journal 6:930-940 (2008)) as template. Construct 660 comprises an alfalfa plastocyanin promoter and 5' UTR, hemagglutinin coding sequence of H5 from A/Indonesia/5/2005 (Construct # 660), alfalfa plastocyanin 3' UTR and terminator sequences (SEQ ID NO: 9; Fig. 5). The resulting fragment was digested with ApaI and StuI restriction enzymes and cloned into construct number 828, previously digested with the same enzymes. The resulting cassette was named construct number 685 (Fig. 1, 2).

Suppressors of silencing.

[001] Post-transcriptional gene silencing (PTGS) may be involved in limiting expression of transgenes in plants, and co-expression of a suppressor of silencing from the potato virus Y (HcPro) may be used to counteract the specific degradation of transgene mRNAs (Brigneti et al., 1998). Alternate suppressors of silencing are well known in the art and may be used as described herein (Chiba et al., 2006, Virology 346:7-14; which is incorporated herein by reference), for example but not limited to, TEV-p1/HC-Pro (Tobacco etch virus-p1/HC-Pro), BYV -p21, p19 of Tomato bushy stunt virus (TBSV p19), capsid protein of Tomato crinkle virus (TCV -CP), 2b of Cucumber mosaic virus; CMV-2b), p25 of Potato virus X (PVX-p25), p11 of Potato virus M

(PVM-p11), p11 of Potato virus S (PVS-p11), p16 of Blueberry scorch virus, (BScV –p16), p23 of Citrus tristeza virus (CTV-p23), p24 of Grapevine leafroll-associated virus-2, (GLRaV-2 p24), p10 of Grapevine virus A, (GVA-p10), p14 of Grapevine virus B (GVB-p14), p10 of Heracleum latent virus (HLV-p10), or p16 of Garlic common latent virus (GCLV-p16). Therefore, a suppressor of silencing, for example, but not limited to, HcPro, TEV -p1/HC-Pro, BYV-p21, TBSV p19, TCV-CP, CMV-2b, PVX-p25, PVM-p11, PVS-p11, BScV-p16, CTV-p23, GLRaV-2 p24, GBV-p14, HLV-p10, GCLV-p16 or GVA-p10, may be co-expressed along with the nucleic acid sequence encoding the protein of interest to further ensure high levels of protein production within a plant.

[002] The construction of p19 is described in described in WO 2010/0003225 (which is incorporated herein by reference). Briefly, the coding sequence of p19 protein of tomato bushy stunt virus (TBSV) was linked to the alfalfa plastocyanin expression cassette by the PCR-based ligation method presented in Darveau et al. (Methods in Neuroscience 26: 77-85(1995)). In a first round of PCR, a segment of the plastocyanin promoter was amplified using primers Plasto-443c:

GTATTAGTAATTAGAATTTGGTGTC (SEQ ID NO:11)

and supP19-plasto.r

CCTTGTATAGCTCGTTCCATTTTCTCTCAAGATG (SEQ ID NO:12)

with construct 660 (described in WO 2010/0003225, which is incorporated herein by reference) as template. In parallel, another fragment containing the coding sequence of p19 was amplified with primers supP19-1c

ATGGAACGAGCTATACAAGG (SEQ ID NO:13)

and SupP19-SacI.r

AGTCGAGCTCTTACTCGCTTTCTTTTTCGAAG (SEQ ID NO:14)

[00112] using construct 35S:p19 as described in Voinnet et al. (The Plant Journal 33: 949-956 (2003)) as template. Amplification products were then mixed and used as template for a second round of amplification (assembling reaction) with primers Plasto-443c and SupP19-SacI.r. The resulting fragment was digested with BamHI (in the plastocyanin promoter) and SacI (at the end of the p19 coding sequence) and cloned into construct number 660, previously

digested with the same restriction enzymes to give construct number R472. The plasmids were used to transform *Agrobacteium tumefaciens* (AGL1; ATCC, Manassas, VA 20108, USA) by electroporation (Mattanovich et al., 1989). The integrity of all *A. tumefaciens* strains were confirmed by restriction mapping. The *A. tumefaciens* strain comprising R472 is termed "AGL1/R472 ".

[00113] HcPro construct (35HcPro) was prepared as described in Hamilton et al. (2002). All clones were sequenced to confirm the integrity of the constructs. The plasmids were used to transform *Agrobacteium tumefaciens* (AGL1; ATCC, Manassas, VA 20108, USA) by electroporation (Mattanovich et al., 1989). The integrity of all *A. tumefaciens* strains were confirmed by restriction mapping.

Preparation of plant biomass, inoculum, agroinfiltration, and harvesting

[00114] *Nicotiana benthamiana* plants were grown from seeds in flats filled with a commercial peat moss substrate. The plants were allowed to grow in the greenhouse under a 16/8 photoperiod and a temperature regime of 25°C day/20°C night. Three weeks after seeding, individual plantlets were picked out, transplanted in pots and left to grow in the greenhouse for three additional weeks under the same environmental conditions. After six weeks, plants have an average weight of 80 g and 30 cm in height.

[00115] *Agrobacterium* strain AGL1 was transfected (electroporation) with constructs as identified below, using the methods described by D'Aoust et al 2008 (Plant Biotechnology Journal 6:930-940). Transfected *Agrobacterium* were grown in YEB medium supplemented with 10 mM 2-(N-morpholino)ethanesulfonic acid (MES), 20 µM acetosyringone, 50 µg/ml kanamycin and 25 µg/ml of carbenicillin pH5.6 to an OD_{600} between 0.6 and 1.6. *Agrobacterium* suspensions were centrifuged before use and resuspended in infiltration medium (10 mM $MgCl_2$ and 10 mM MES pH 5.6).

[00116] Plants were agroinfiltrated as described in D'Aoust et al (*supra*). Briefly, for vacuum-infiltration, *A. tumefaciens* suspensions were centrifuged, resuspended in the infiltration medium and stored overnight at 4°C. On the day of infiltration, culture batches were diluted in 2.5 culture volumes and allowed to warm before use. Whole plants of *N. benthamiana* were placed upside down in the bacterial suspension in an air-tight stainless steel tank under a vacuum of 20-40 Torr for 2-min. Unless otherwise specified, all infiltrations were performed as co-

infiltration with a bacterial transformed with R472 (strain AGL1/R472) at a 1:1 ratio. Following vacuum infiltration, plants were returned to the greenhouse for a 4-6 day incubation period until harvest.

Leaf sampling and total protein extraction (mechanical homogenization)

[00117] Following incubation of 4, 5, 6, 7 and 8 days, the aerial part of plants was harvested and used immediately. Total soluble proteins were extracted by homogenizing plant tissue in 3 volumes of cold 50 mM Tris pH 8.0, 0.15 M NaCl containing 1% Trition X-100 and 0.004% sodium metabisulfite. Plant tissue were mechanically homogenized using a POLYTRON™, grinding with mortar and pestle, or with a COMITROL™ in 1 volume of cold 50 mM Tris pH 8, 0.15 M NaCl. The buffer used with the COMITROL™ also contained 0.04% sodium metabisulfite. Following homogenization, the slurry of ground plant material was centrifuged at 5,000 g for 5min at 4°C and the crude extracts (supernatant) kept for analysis. The total protein content of clarified crude extracts was determined by the Bradford assay (Bio-Rad, Hercules, CA) using bovine serum albumin as the reference standard.

VLP extraction by cell wall digestion

[00118] Leaf tissue was collected from the *Nicotiana benthamiana* plants and cut into ~1 cm^2 pieces. The leaf pieces were soaked in 500 mM mannitol for 30 minutes at room temperature (RT). The mannitol solution was then removed and changed with the enzyme mix (mixture of cellulases from *Trichoderma viride* (Onozuka R-10; 3% v/v) and a mixture of pectinases from *Rhizopus* sp. (MACEROZYME™; 0.75% v/v; both from Yakult Pharmaceuticals) in protoplasting solution (500 mM mannitol, 10mM $CaCl_2$ and 5 mM MES/KOH (pH 5.6)). The ratio used was 20 g of leaf pieces per 100 mL solution. This preparation was spread evenly into a shallow vessel (~11x18 cm) and incubated for 16 hours on a rotary shaker at 40 rpm and 26°C.

[00119] Alternately, VLP extraction may be performed as follows: plants were agroinfiltrated with AGL1/#685 as described in example 1. Leaf tissue was collected from the *N. benthamiana* plants at day 6 post-infiltration and cut into ~1 cm^2 pieces. Multifect Pectinase FE, Multifect CX CG and Multifect CX B (Genencor) were added to 1.0% each (v/v) in a 600 mM Mannitol, 75 mM Citrate, 0.04% sodium bisulfite pH 6.0 buffer using a ratio of 1:2.5 (w/v) fresh biomass; digestion buffer. The biomass was digested for 15h at room temperature in a orbital shaker.

[00120] Following incubation, leaf debris was removed by filtration (nylon filter of 250 or 400 μm mesh). Protoplasts in suspension were collected by centrifugation at 200xg (15 min), followed by centrifugation of the supernatant at 5000xg (15 min) to further clarify the supernatant. Alternately, a single centrifugation step at 5000 xg for 15 minutes may be employed. Seventy mL of the supernatant was then centrifuged at 70,000xg for 30 minutes. The resulting pellet was resuspended in 1.7mL of PBS and analyzed immediately or frozen.

Protein Analysis

[00121] A hemagglutination assay for H5 was based on a method described by Nayak and Reichl (2004). Briefly, serial double dilutions of the test samples (100 μL) were made in V-bottomed 96-well microtiter plates containing 100 μL PBS, leaving 100 μL of diluted sample per well. One hundred microliters of a 0.25% turkey red blood cells suspension (Bio Link Inc., Syracuse, NY) were added to each well, and plates were incubated for 2h at room temperature. The reciprocal of the highest dilution showing complete hemagglutination was recorded as hemagglutination activity. In parallel, a recombinant HA5 standard (A/Vietnam/1203/2004 H5N1) (Protein Science Corporation, Meriden, CT) was diluted in PBS and run as a control on each plate.

ELISA

[00122] HA5 standard was prepared with purified virus-like particles which were disrupted by treatment with 1% Triton X-100 followed by mechanical agitation in a Tissue Lyser™ (Qiagen) for 1 min. U-bottom 96-well microtiter plates were coated with 10 μg/mL of capture antibody (Immune Technology Corporation, #IT-003-005I) in 50 mM carbonate-bicarbonate coating buffer (pH 9.6) for 16-18 hours at 4°C. All washes were performed with 0.01 M PBS (phosphate-buffered saline), pH 7.4 containing 0.1% Tween-20. After incubation, plates were washed three times and blocked with 1% casein in PBS for 1 hour at 37°C. After the blocking step, plates were washed three times. The HA5 standard was diluted in a mock extract (prepared from leaf tissue infiltrated with AGL1/R472 alone) to generate a standard curve from 500 to 50 ng/mL. Samples to quantify were treated in 1% Triton X-100 prior to loading the microplate. Plates were further incubated for 1 hour at 37°C. After washing, sheep polyclonal antibody raised against HA5 (CBER/FDA) diluted 1:1000 was added and the plates were incubated for 1 hour at 37°C. After washing, horseradish peroxidase-conjugated rabbit anti-sheep antibody diluted 1:1000 was added and the plates were incubated for 1 hour at 37°C. After

the final washes, the plates were incubated with SureBlue TMB peroxidase substrate (KPL) for 20 minutes at room temperature. Reaction was stopped by the addition of 1N HCl and A_{450} values were measured using a Multiskan Ascent plate reader (Thermo Scientific).

Example 1: Enzymatic extraction of plant tissue releases high quantities of HA having an elevated relative activity.

[00123] The quantity and relative activity of HA obtained from the present enzymatic extraction method were compared with that of HA obtained from common mechanical extraction methods. *N. benthamiana* plants were infiltrated with AGL1/685 and the leaves were harvested after a five to six-day incubation period. Leaf homogenates were prepared as follows : Two grams of leaves were homogenized with a Polytron homogenizer; 4g of leaves were ground with a mortar and a pestle; and 25kg of leaves were homogenized with a COMITROL™ processor (Urschel Laboratories) in an extraction buffer (50 mM Tris, 150 mM NaCl pH 8.0, ratio of 1:1 w/v). Enzymatic extraction was carried as follow: Twenty grams of harvested leaves were subjected to digestion with Macerozyme pectinases and Onozuka R-10 cellulases as described above. Following digestion, leaf debris were removed by filtration (nylon filter, 250 µm mesh). Protoplasts in suspension were removed by centrifugation at 200xg (15 min), and the supernatant further clarified by centrifugation at 5000xg (15 min).

[00124] The quantity and relative activity of HA in each of these plant extracts is shown in Table 3. The quantity of HA obtained from the enzymatic extraction method is significantly greater than that obtained from the mechanical methods.

Table 3: Comparative quantities and relative activities of HA obtained from mechanical and enzymatic extraction of plant leaves. All data have been adjusted to take into account the differences in volume of liquid added for each extraction method. The Comitrol™ extraction method was chosen as the standard value for the purpose of the present comparative analysis.

Extraction method	Relative activity	Quantity*
Comitrol™ extract	*100%*	*100%*
Polytron extract	50%	150%
Mortar extract	100%	220%
Digestion extract	440%	570%

*Quantity was evaluated by ELISA analysis

Example 2: Enzymatic digestion of plant tissue releases HA organized into VLPs.

[00125] A combination of differential centrifugation and size exclusion chromatography (SEC) was used to demonstrate that the HA obtained by the enzymatic extraction method described herein were organized as VLPs. *N. benthamiana* plants were agroinfiltrated with AGL1/685 as described in Example 1. Leaves were collected from the plants 6 days post-infiltration, cut into ~1 cm^2 pieces and then digested, coarse-filtered and centrifuged as described in Example 1.

[00126] The clarified samples were then centrifuged at 70,000xg to allow for segregation of VLPs. The centrifugation pellet, containing the VLPs, was gently resuspended in 1/50 volume of Phosphate buffered saline (PBS; 0.1M sodium phosphate, 0.15M NaCl pH 7.2) before being loaded on a SEC column.

[00127] SEC columns of 32 ml SEPHACRYL™ S-500 high resolution beads (S-500 HR : GE Healthcare, Uppsala, Sweden, Cat. No. 17-0613-10) were prepared with equilibration/elution buffer (50 mM Tris, 150 mM NaCl, pH8). SEC chromatography was performed with the loading of a 1.5 mL VLP sample onto the equilibrated column, and its elution with 45 mL of equilibration/elution buffer. The eluate was collected in fractions of 1.7 mL, and the protein content of each fraction was evaluated by mixing 10 µL of the eluate fraction with 200 µL of diluted Bio-Rad protein dye reagent (Bio-Rad, Hercules, CA). Each separation was preceded by a calibration with Blue Dextran 2000 (GE Healthcare Bio-Science Corp., Piscataway, NJ, USA). Comparison of the elution profiles of both Blue Dextran 2000 and host proteins was performed for each separation to ensure uniformity of the separations.

Protein Analysis of the SEC eluted fractions

[00128] Total protein content of clarified crude extracts was determined by the Bradford assay (Bio-Rad, Hercules, CA) using bovine serum albumin as the reference standard. Proteins present in SEC eluate fractions were precipitated with acetone (Bollag et al., 1996), resuspended in either 0.25 volume or 0.05 volume of denaturing sample loading buffer (0.1M Tris pH 6.8, 0.05% bromophenol blue, 12.5% glycerol, 4% SDS and 5% beta-mercaptoethanol) for SDS-PAGE analysis or immunoblot analysis, respectively. Separation by SDS-PAGE was performed under reducing conditions, and Coomassie Brillant Blue R-250 was used for protein staining.

[00129] Hemagglutination assay for H5 was performed based on a method described by Nayak and Reichl (2004). Briefly, successive double dilutions of the test samples (100 µL) were made in V-bottomed 96-well microtiter plates containing 100 µL PBS, leaving 100 µL of diluted sample per well. One hundred microliters of a 0.25% turkey red blood cells suspension (Bio Link Inc., Syracuse, NY) were added to each well, and plates were incubated for 2h at room temperature. The reciprocal of the highest dilution showing complete hemagglutination was recorded as hemagglutination activity. In parallel, a recombinant H5 standard (A/Vietnam/1203/2004 H5N1) (Protein Science Corporation, Meriden, CT) was diluted in PBS and run as a control on each plate.

[00130] Figure 3A shows that the hemagglutination activity is concentrated in the fractions corresponding to the void volume of the column, confirming that the hemagglutination activity originates from a high molecular weight structural organization. SDS-PAGE analysis (Fig. 3B) revealed that those same void volume fractions (fractions 7-10) also present the highest HA content, with a band corresponding to the HA0 monomer being detectable at approximately 75 kDa.

Example 3: Enzymatic digestion of plant tissue releases HA-VLPs with fewer contaminants

[00131] *N. benthamiana* plants were agroinfiltrated with AGL1/685 as described in Example 1. Leaves were collected on day 6 post-infiltration, cut into ~1 cm^2 pieces, digested, coarse-filtered and centrifuged as described in Example 1.

[00132] The controlled enzymatic digestion of the leaves removed the cell walls, at least partially, thus allowing for the release of proteins and components presents in the space between the cell wall and the plasma membrane into the extraction medium. Since most intracellular proteins and components were still undamaged and contained within the mostly intact protoplasts, an initial centrifugation step allowed for their removal, thus providing a resulting solution comprising cell wall degrading enzymes, in addition of the extracellular plant proteins and components (apoplastic fraction), as shown in Figure 4.

[00133] Figure 4 shows a SDS-PAGE analysis of the resulting solution obtained following the controlled enzymatic digestion of leaves tissue as described previously, with lane 1 showing the enzyme mixture used and lane 2 showing the resulting solution following the enzymatic digestion. The protein content of a crude extract from Comitrol™ is provided on lane 3 for

comparison. The biomass:buffer ratio for the extract presented in lane 2 was 1:5 (w/v) while it was 1:1 (w/v) for that in lane 3. Each of lanes 2 and 3 therefore contain proteins derived from an equivalent quantity of starting material. For approximately the same buffer:plant ratio, a mechanical plant extract contained a protein concentration of approximately 3.5-4 mg/ml, while the enzymatic plant extract obtained according to the present method presented a protein concentration of approximately 1 mg/ml.

[00134] The major contaminant present in lane 3 was found to be RubisCo (Ribulose-1,5-bisphosphate carboxylase oxygenase), which is made of two types of protein subunits: a large-chain (L, about 55 kDa) and a small-chain (S, about 13 kDa). A total of eight large-chain dimers and eight small-chains usually assemble with each other into a RubisCo 540 kDa larger complex. While this plant protein contaminant is found in large amount in plant extracts originating from mechanical extraction method (see arrow in Figure 4), it is virtually absent in plant extracts obtained by the enzymatic digestion method described herein. Therefore, the present method allows for the elimination of this major plant protein contaminant, amongst others, at an early stage of the process.

Example 4: Enzymatic digestion of plant tissue releases HA-VLP in conditions where it can be directly captured on a cation exchange resin.

[00135] *N. benthamiana* plants were agroinfiltrated with AGL1/685 as described in Example 1. Leaves were collected on day 6 post-infiltration, cut into ~1 cm^2 pieces and digested for 15h at room temperature in an orbital shaker. The digestion buffer contained 1.0% (v/v) Multifect Pectinase FE, 1.0% (v/v) Multifect CX CG orand 1.0% (v/v) Multifect CX B (all from Genencor), each in a solution of 600 mM Mannitol, 75 mM Citrate, 0.04% sodium bisulfite pH 6.0 buffer using a biomass : digestion buffer ratio of 1:2.5 (w/v).

[00136] Following digestion, the apoplastic fraction was filtered through a 400 µm nylon filter to remove coarse undigested vegetal tissue (<5% of starting biomass). The filtered extract was then centrifuged at room temperature for 15 min at 5000xg to remove protoplasts and intracellular contaminants (proteins, DNA, membranes, vesicles, pigments, etc). Next, the supernatant was depth-filtered (for clarification) using a 0.65µm glass fiber filter (Sartopore2/Sartorius Stedim) and a 0.45/0.2µm filter, before being subjected to chromatography.

[00137] The clarified apoplastic fraction was loaded over a cation exchange column (Poros HS Applied Biosystems) equilibrated with an equilibration/elution buffer (50 mM $NaPO_4$, 100 mM NaCl, 0.005% Tween 80 pH 6.0). Once the UV was back to zero, the extract was step-eluted with the equilibration/elution buffer containing increasing concentrations of NaCl (500 mM). Where necessary, the chromatographic fractions were concentrated 10 times using Amicon™ devices equipped with 10 kDa MWCO. Protein analysis was performed as described in previous examples.

[00138] Under the above-mentioned conditions, most enzymes and plant proteins did not bind to the cation exchange resin whereas the HA-VLP did bind, thus providing a considerable enrichment in HA-VLPs in the eluted fraction (Figure 6). In addition, as shown in Figure 6, lane 4 and 5, the cellulases and pectinases did not bind to the cation exchange column at pH under 7. Consequently, recovery of HA-VLP, based on HA hemagglutination activity, was of 92% prior to loading on the cation exchange column, and of 66% in the eluted fraction. A purification factor of 194 was measured on the eluted fraction from the cation exchange resin.

Example 5: Addition of NaCl to the digestion buffer

[00139] *N. benthamiana* plants were agroinfiltrated with *Agrobacterium* AGL1 strains carrying a construct expressing a hemagglutinin of interest (H1/Cal WT, B/Flo, H5/Indo or H1/Cal X179A) as described in Example 1. Leaves were collected on day 6 post-infiltration, cut into ~1 cm^2 pieces and digested according to Example 4, except where noted below. Filtration, centrifugation and clarification were performed as described in Example 4.

[00140] NaCl was added to digestion buffer to evaluate its potential effect on the HA-VLP recovery rate. The suspected advantages were the potential prevention of non-specific association of HA with plant cells or with particle in suspension that are removed during clarification and potential effect on achievement and/or maintenance and/or improvement of colloidal stability of the HA-VLP.

[00141] Addition of 500 mM NaCl to the digestion buffer resulted in an increase of HA-VLP recovery yield per gram of biomass after removal of protoplasts and cellular debris by centrifugation. However, this increase was only noted with the for the H1/Cal WT and B/Flo strains, while the recovery yield for H5 was not significantly increased by this approach (Table 4).

Table 4 : Effect of the addition of NaCl to the digestion step on the HA-VLP recovery yield (as measured by hemagglutination activity unit, dil : reciprocal of dilution)

HA strain	Digestion conditions	Concentration in HA (dil /ml)	Yields (dil/g)	Yield increased (X-fold)[1]
H5 Indo/05 (#972)	Ø NaCl	4608	12,430	1.2
	500 mM NaCl	4608	14,921	
H1 CA/07 WT (#604)	Ø NaCl	384	1,206	2.1
	500 mM NaCl	768	2,481	
H1 CA/07 X-179A (#605)	Ø NaCl	96	299	8.1
	500 mM NaCl	768	2,419	
B Flo/4 (475)	Ø NaCl	16	52	7.5
	500 mM NaCl	128	392	

[1] Yield (dil/g) with NaCl divided by Yield (dil/g) without NaCl

[00142] Addition of 500 mM NaCl during the digestion further resulted in an increase of the release of HA-VLP during digestion, which in turn resulted into increased recovery rate after clarification for both H1/Cal WT and H1/Cal X-179A strains (Table 5), but not for the H5/Indo strain.

Table 5 : Effect of the addition of NaCl to the digestion step on the HA-VLP recovery yield (as measured by hemagglutination activity unit) after the clarification step.

HA strain	Digestion conditions	Recovery after depth filtration[1]	Increase in recovery (X-fold)
H5/Indo (#972)	Ø NaCl	100%	1.0
	500 mM NaCl	100%	
H1/Cal WT (#604)	Ø NaCl	25%	3.0
	500 mM NaCl	75%	
H1/Cal X-179A (#605)	Ø NaCl	50%	2.0
	500 mM NaCl	100%	

[1] Recovery is expressed in percentage of hemagglutination activity obtained after depth filtration compared to the activity found in the centrifuged digested extract.

[00143] The association state of the HA-VLP, with and without the addition of NaCl during enzymatic digestion, was studied using Nanoparticle Tracking Analysis (NTA) for H5/Indo and H1/Cal WT (Figure 7A and 7B respectively). A monodisperse preparation of particles was observed for H5 when digestion was performed in absence of NaCl, while the H1/Cal preparation showed much larger array of particle species. The addition of NaCl to the digestion buffer reduced HA-VLP self-association for H1/Cal, as shown by the fairly monodisperse particle distribution found in Figure 7C. The number of particles at 150 nm for H1/Cal WT-VLPs was enhanced (ca 5-fold) by the addition of 500 mM NaCl to the digestion buffer.

Example 6: Controlling the release of pigments

[00144] *N. benthamiana* plants were agroinfiltrated with *Agrobacterium* AGL1 strains carrying a construct expressing a hemagglutinin of interest (H5/Indo) as described in Example 1. Leaves were collected on day 6 post-infiltration, cut into ~1 cm^2 pieces, and digested as described in Example 4, with addition of either 500 mM NaCl or 500 mM NaCl and 25 mM EDTA to the digestion buffer. Filtration, centrifugation and clarification were performed as described in Example 4.

[00145] Release of components having a green color during the enzymatic digestion step led to purified preparation of VLP having a greenish coloration. The composition of the cell wall digestion solution was therefore investigated and adjusted to obtain a VLP purified preparation having a reduced green coloration, and thus an increased purity. Without wishing to be bound by theory, since Ca^{2+} plays a critical role in the retention of constituents of the cell wall's middle lamellae together, and given the fact that there is usually a high concentration of Ca^{2+} in plant cell wall, the addition of Ca^{2+}-chelator EDTA could facilitate the enzymatic depolymerisation of the cell wall, thereby preserving intact intracellular organelles, such as chloroplasts, and preventing the release green-pigments components.

[00146] As shown in Table 6, the addition of 25mM EDTA to the digestion buffer allowed for the reduction of the green coloration of the purified H5-VLP preparation, as evaluated by measuring the difference in absorption of the preparation (OD_{672nm} - OD_{650nm}). When the green constituents were released in high quantity, or not suitably removed, VLP preparation exhibited a $\Delta OD > 0.040$.

Table 6 : Effect of the addition of 25 mM EDTA to the digestion buffer on green coloration of H5-VLP preparations.

	OD_{672nm} - OD_{650nm}
0 mM NaCl, 0 mM EDTA	**0.071 ± 0.061**
500 mM NaCl	**0.087 ± 0.060**
500 mM NaCl + 25 mM EDTA	**0.010 ± 0.002**

Example 7: Alternative digestion buffer compositions

[00147] *N. benthamiana* plants were agroinfiltrated with *Agrobacterium* AGL1 strains carrying a construct expressing a hemagglutinin of interest (H5/Indo) as described in Example 1. Leaves were collected on day 6 post-infiltration, cut into ~1 cm^2 pieces and digested according to Example 4, with modification of digestion buffer to include 0%, 0.25%, 0.5%, 0.75% or 1% v/v Multifect Pectinase FE, Multifect CX- CG cellulase and Multifect CX B cellulose as noted in Tables 7-9. Filtration, centrifugation and clarification were as described in Example 4.

[00148] As shown in following tables 7 and 8, pectinase has been demonstrated to be non-essential in the digestion buffer. Similar levels of H5/Indo or H1/Cal WT VLP can be extracted with the present method either in the presence or absence of pectinase. Furthermore, it has been found that reducing the concentration of cellulase when compared to previous examples had no significant impact on the quality of extraction (Table 9).

Table 7 : Release of H5/Indo VLP by digestion of *N. benthamiana* leaves. All conditions were tested in replicates. (Concentration in HA-VLP measured by hemagglutination activity, dil : reciprocal of dilution)

Pectinase (% v/v)	Cellulase* (% v/v)	Concentration in H5 VLP (dil/ml)
1	1	1152
0.5	1	6144
0	1	768
0	2	1536

*Multifect CX GC

Table 8: Release of H1/Cal WT VLP by digestion of *N. benthamiana* leaves. All conditions were tested in replicates. (Concentration in HA-VLP measured by hemagglutination activity, dil : reciprocal of dilution)

Pectinase (% v/v)	Cellulase* (% v/v)	Concentration in H1 VLP (dil/ml)
1	2	2304
0	2	3840

*1% each of Multifect CX GC and Multifect CX B

Table 9 : Release of H1/Cal WT VLP by digestion of *N. benthamiana* leaves. All conditions were tested in replicates. (Concentration in HA-VLP measured by hemagglutination activity, dil:, reciprocal of dilution)

Pectinase (% v/v)	Cellulase* (% v/v)	Concentration in H1 VLP (dil/ml)
1.0	1	384
0.75	1	480
0.50	1	480
0.25	1	480

*Multifect CX GC

Example 8: Enzymatic digestion in conditions near to neutral pH

[00149] Controlling the pH during the digestion can be critical for the extraction of some VLPs. Taking into account that the depolymerisation of the cell wall occurring during the digestion step can release acid sugars that could acidify the solution (i.e. from pH 6 to 5) in the presence of appropriate buffers, and that some VLPs (such as those comprising H3/Bris and B/Flo HA) have already demonstrated a strong sensitivity to mildly acidic conditions, impact of such a potential acidification on the yield of VLP produced was investigated.

[00150] *N. benthamiana* plants were agroinfiltrated with *Agrobacterium* AGL1 strains carrying a construct expressing a hemagglutinin of interest (B/Flo, H5/Indo, H3/Bris) as described in Example 1. Leaves were collected on day 6 post-infiltration, cut into ~1 cm^2 pieces and digested according to Example 4, with modification of digestion conditions to include 500 mM NaCl; 25 or 50 mM EDTA; 0.03 or 0.04 % sodium bisulfite; 0, 100, 200 or 600 mM mannitol, 75, 125 or 150 mM citrate; and/or 75 mM $NaPO_4$; with the pH of the digestion buffer

adjusted as set out in Tables 10-14. Filtration, centrifugation and clarification were as described in Example 4.

[00151] Various digestion buffer compositions were tested to achieve a pH of approximately 5.5 by the end of the enzymatic digestion, including increased concentration of citrate (buffer effect between pH 3.0 and 5.4) and addition of sodium phosphate (buffer effect at pH above 6.0). Table 10 shows that VLPs from the B strain were extracted more efficiently when post-digestion pH was close to pH 6.0.

Table 10: Effect of the digestion buffer composition on the extraction yield of B/Flo VLPs.

Buffer composition[1]	Concentration of B/Flo VLP (dil/ml)	Protein concentration (mg/ml)	pH post-digestion
75 mM Citrate + 500mM NaCl + 25 mM EDTA pH 6.0	1	0.92	5.0
75 mM Citrate pH 6.0	0	1.43	5.6
125 mM Citrate + 500mM NaCl + 25 mM EDTA pH 6.0	1.5	1.07	5.4
150 mM Citrate + 500mM NaCl + 25 mM EDTA pH 6.0	1.5	1.07	5.4
125 mM Citrate + 75mM $NaPO_4$ + 500mM NaCl + 25 mM EDTA pH 6.5	4	2.19	5.9

[1]All buffers contained 600 mM mannitol, sodium metabisulfite 0.04%

[00152] Next, the effect of initiating the digestion at a higher pH in order to reach final pH value close to pH 6.0 was tested. As shown in Table 11, the digestion of plant cell wall with such near-neutral conditions was possible, and did not impaired the extraction yield for H5/Indo VLPs.

Table11: Effect of the initial pH of the digestion buffer on the extraction yield of H5/Indo VLPs.

Initial pH of digestion solution[1]	Concentration of H5/Indo VLP (dil/ml)	Protein concentration (mg/ml)	pH post-digestion
6.5	2304	2.79	6.08
6.4	1536	2.31	5.93
6.3	2304	2.40	5.81
6.2	2304	2.09	5.73
6.1	2304	1.72	5.61

[1]All digestion buffers contained 600 mM mannitol, sodium metabisulfite 0.04%,125 mM Citrate + 75mM $NaPO_4$ + 500mM NaCl + 25 mM EDTA

[00153] Other components of the digestion solution were also shown to be modifiable without negatively affecting the extraction yield of VLPs. Table 12 illustrates modifications that can be applied to the digestion solution in order to enhance the extraction yield of B/Flo VLPs, while obtaining a post-digestion pH of 5.4 -5.7. Such modifications include increasing the concentration of citrate and adding a PO_4 buffer. It has been found that increasing the concentration of EDTA generally led to a more acidified extract and to lower VLP extraction yields.

Table 12: Effect of various digestion buffer components on the extraction yield of B/Flo VLPs.

Buffer composition[1]							
Mannitol (mM)	Citrate (mM)	PO_4 (mM)	EDTA (mM)	pH	Concentration of B VLP (dil/ml)	Protein concentration (mg/ml)	pH post-digestion
600	75	0	25	6.1	2	1.07	5.0
600	125	0	25	6.1	192	0.83	5.7
600	125	75	25	6.2	192	1.81	5.5
600	125	75	50	6.2	96	1.26	5.4
200	125	75	25	6.2	384	1.05	5.7
200	125	75	50	6.2	96	1.04	5.4
200	125	75	75	6.2	96	1.55	5.4

[1]All buffers contained 500 mM NaCl and sodium metabisulfite 0.04%.

[00154] Buffer composition was further modified to improve the extraction yield of H3/Brisbane VLPs (Table 13)

Table 13: Effect of the concentrations of mannitol and sodium bisulfite in the digestion solution on the extraction yield of H3/Bris VLPs.

Buffer composition					
Mannitol (mM)	Sodium bisulfite (%)	EDTA (mM)	pH	Protein concentration (mg/ml)	pH post-digestion
600	0.04	25	6.2	1.87	5.7
600	0.04	50	6.2	1.62	5.6
200	0.03	25	6.2	1.89	5.7
200	0.03	50	6.2	1.24	5.6

[1]All buffers containing 125 mM Citrate, 75 mM $NaPO_4$, 500 mM NaCl.

[00155] As shown in Tables 12 and 13, mannitol concentration could be reduced to 200 mM without significantly affecting VLPs extraction yield. Further reduction of mannitol concentrations to 100 mM, and even the total omission of mannitol from the digestion solution, did not significantly affect the level of HA-VLP obtained (Table 14).

Table 14: Released of H5/Indo VLP from digestion of biomass performed in buffers with different concentration of mannitol.

Mannitol concentration of the digestion solution [1]	Concentration of H5/Indo VLP (dil/ml)	Protein concentration (mg/ml)
[2]Trial 1: without mannitol	2304	1.62
[2]Trial 1: with 600 mM mannitol	3072	1.73
[2]Trial 2: with 100 mM mannitol	4608	1.77
[2]Trial 2: with 600 mM mannitol	4608	2.0

[1]All buffers containing 75 mM Citrate pH 6.0 + sodium metabisulfite 0.04%.
[2]Two trials were were performed to compare the extraction yields of VLPs without mannitol (Trial 1) and with 100mM mannitol (Trial 2) versus 600 mM mannitol.

Example 9: Suitability of enzymatic digestion to a broad variety of HA-VLPs

[00156] The enzymatic digestion method for plant biomass described herein has the potential to be applied to extracting of a broad variety of HA-VLPs. Adding to the extraction of HA-VLPs comprising H5/Indo, H1/Cal WT VLP, H3/Bris and B/Flo shown in previous examples, the method described herein was also shown to be suitable for the extraction of HA-VLPs from seasonal H1/Bris and H1/NC, as shown in Table 15.

Table 15 : Release of seasonal H1/Bris and H1/NC VLP from digestion of agroinfiltrated *N. benthamiana* leaves. (concentration in HA measured by hemagglutination activity, dil : reciprocal of dilution)

HA strain	Concentration in HA (dil /ml)
H1/Bri	1536
H1/NC	384

[00157] All citations are herein incorporated by reference, as if each individual publication was specifically and individually indicated to be incorporated by reference herein and as though it were fully set forth herein. Citation of references herein is not to be construed nor considered as an admission that such references are prior art to the present invention.

[00158] One or more currently preferred embodiments of the invention have been described by way of example. The invention includes all embodiments, modifications and variations substantially as hereinbefore described and with reference to the examples and figures.

It will be apparent to persons skilled in the art that a number of variations and modifications can be made without departing from the scope of the invention as defined in the claims. Examples of such modifications include the substitution of known equivalents for any aspect of the invention in order to achieve the same result in substantially the same way.

WHAT IS CLAIMED IS:

1. A method of preparing plant derived virus-like particles (VLPs), comprising:
   a. obtaining a plant or plant matter comprising apoplast-localized VLPs;
   b. producing a protoplast/spheroplast fraction and an apoplast fraction by treating the plant or plant matter with a cell wall degrading enzyme composition; and,
   c. recovering the apoplast fraction, the apoplast fraction comprising the plant-derived VLPs.

2. The method of claim 1 wherein the cell wall degrading enzyme composition comprises one or more than one pectinase, one or more than one cellulase, or one or more than one pectinase and one or more than one cellulase.

3. The method of claim 2, wherein the cell wall degrading enzyme composition does not include one or more of a lipase, a protease or a pectinase.

4. The method of claim 1 wherein in the step of obtaining (step a), the plant is transformed with a nucleic acid sequence encoding a protein selected from the group of a viral envelope protein, a viral structural protein, a viral capsid protein, and a viral coat protein, and the plant or plant matter is harvested.

5. The method of claim 4, wherein the nucleic acid is introduced into the plant in a transient manner.

6. The method of claim 4, wherein, the nucleic acid is stably integrated within a genome of the plant.

7. The method of claim 1 wherein in the step of obtaining (step a), the plant is grown and the plant or plant matter harvested.

8. The method of claim 4 wherein the nucleic acid encodes influenza hemagglutinin.

9. The method of claim 1 wherein the plant derived virus-like particles (VLPs) do not include neuraminidase or M protein.

10. The method of claim 1 wherein the plant matter is selected from the group of leaves, and cultured plant cells.

11. The method of claim 1 further comprising a step of d) purifying the plant derived virus-like particles (VLPs) from the apoplast fraction.

12. The method of claim 11, wherein the step of purifying comprises filtering the apoplast fraction using depth filtration to produce a clarified extract, followed by chromatography of the clarified extract using a cation exchange resin.

13. A method of preparing plant-derived virus-like particles (VLPs) comprising a plant-derived lipid envelope, comprising:

a. obtaining a plant, or plant matter comprising apoplast-localized VLPs;
b. treating the plant or plant matter with an enzyme composition to produce a protoplast/spheroplast fraction and one or more than one apoplastic protein complex that comprises the VLPs; and
c. separating the one or more than one apoplastic protein complex from the protoplast/spheroplast fraction.

14. The method of claim 13 wherein the enzyme composition comprises one or more than one pectinase, one or more than one cellulase, or one or more than one pectinase and one or more than one cellulase.

15. The method of claim 13 wherein the enzyme composition does not include one or more of a lipase, a protease or a pectinase.

16. The method of claim 13 further comprising the step of d) purifying the plant derived virus-like particles (VLPs) from the one or more than one apoplastic protein complex.

17. The method of claim 16, wherein the step of purifying comprises depth filtration to produce a clarified extract, followed by chromatography of the clarified extract using a cation exchange resin.

18. The method of claim 13 wherein in the step of obtaining (step a), the plant is transformed with a nucleic acid sequence encoding a protein selected from the group of a viral envelope

protein, a viral structural protein, a viral capsid protein, and a viral coat proteins, and the plant or plant matter is harvested.

19. The method of claim 18, wherein the nucleic acid is introduced into the plant in a transient manner.

20. The method of claim 18, wherein, the nucleic acid is stably integrated within a genome of the plant.

21. The method of claim 13 wherein in the step of obtaining (step a), the plant is grown and the plant or plant matter harvested.

22. The method of claim 18 wherein the virus-like particles (VLPs) comprise influenza hemagglutinin.

23. The method of claim 13 wherein the plant matter is selected from the group of leaves, and cultured plant cells.

24. The method of claim 13 wherein the virus-like particles (VLPs) do not include neuraminidase or M protein.

25. A method of preparing plant derived virus-like particles (VLPs), comprising:

a. obtaining a plant or plant matter comprising plant-derived VLPs;

b. digesting the plant or plant matter using a cell wall degrading enzyme composition to produce a digested fraction;

c. filtering the digested fraction to produce a filtered fraction and recovering the plant-derived VLPs from the filtered fraction.

26. The method of claim 25 wherein the cell wall degrading enzyme composition comprises one or more than one pectinase, one or more than one cellulase, or one or more than one pectinase and one or more than one cellulase.

27. The method of claim 25, wherein the cell wall degrading enzyme composition does not include one or more of a lipase, a protease or a pectinase.

28. The method of claim 25 wherein in the step of obtaining (step a), the plant is transformed with a nucleic acid sequence encoding a protein selected from the group of a viral envelope protein, a viral structural protein, a viral capsid protein, and a viral coat proteins, and the plant or plant matter is harvested.

29. The method of claim 28, wherein the nucleic acid is introduced into the plant in a transient manner.

30. The method of claim 28, wherein, the nucleic acid is stably integrated within a genome of the plant.

31. The method of claim 28, wherein the plant derived virus-like particles (VLPs) comprise influenza hemagglutinin.

32. The method of claim 25, wherein the plant matter is selected from the group of leaves, and cultured plant cells.

33. The method of claim 25 further comprising a step d) of separating the virus-like particles (VLPs) in the filtered fraction from cellular debris and insoluble materials.

34. The method of claim 33, wherein the step of separating is performed by centrifugation.

35. The method of claim 33 wherein the step of separating is performed by depth filtration.

36. The method of claim 25, further comprising the step of d) purifying the plant derived virus-like particles (VLPs) from the filtered fraction.

37. The method of claim 36, wherein the step of purifying comprises depth filtration of the filtered fraction to produce a clarified extract, followed by chromatography of the clarified extract using a cation exchange resin.

SEQUENCE LISTING

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<120> METHOD OF PREPARING PLANT-DERIVED VLPS

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Asp Thr Ile Met Glu Lys Asn Val Thr Val Thr His Ala Gln Asp Ile
35 40 45

Leu Glu Lys Thr His Asn Gly Lys Leu Cys Asp Leu Asp Gly Val Lys
50 55 60

Pro Leu Ile Leu Arg Asp Cys Ser Val Ala Gly Trp Leu Leu Gly Asn
65 70 75 80

Pro Met Cys Asp Glu Phe Ile Asn Val Pro Glu Trp Ser Tyr Ile Val
85 90 95

Glu Lys Ala Asn Pro Thr Asn Asp Leu Cys Tyr Pro Gly Ser Phe Asn
100 105 110

Asp Tyr Glu Glu Leu Lys His Leu Leu Ser Arg Ile Asn His Phe Glu
115 120 125

Lys Ile Gln Ile Ile Pro Lys Ser Ser Trp Ser Asp His Glu Ala Ser
130 135 140

Ser Gly Val Ser Ser Ala Cys Pro Tyr Leu Gly Ser Pro Ser Phe Phe
145 150 155 160

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Arg Asn Val Val Trp Leu Ile Lys Lys Asn Ser Thr Tyr Pro Thr Ile
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Lys Lys Ser Tyr Asn Asn Thr Asn Gln Glu Asp Leu Leu Val Leu Trp
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Gly Ile His His Pro Asn Asp Ala Ala Glu Gln Thr Arg Leu Tyr Gln
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Asn Pro Thr Thr Tyr Ile Ser Ile Gly Thr Ser Thr Leu Asn Gln Arg
    210                 215                 220

Leu Val Pro Lys Ile Ala Thr Arg Ser Lys Val Asn Gly Gln Ser Gly
225                 230                 235                 240

Arg Met Glu Phe Phe Trp Thr Ile Leu Lys Pro Asn Asp Ala Ile Asn
                245                 250                 255

Phe Glu Ser Asn Gly Asn Phe Ile Ala Pro Glu Tyr Ala Tyr Lys Ile
            260                 265                 270

Val Lys Lys Gly Asp Ser Ala Ile Met Lys Ser Glu Leu Glu Tyr Gly
        275                 280                 285

Asn Cys Asn Thr Lys Cys Gln Thr Pro Met Gly Ala Ile Asn Ser Ser
    290                 295                 300

Met Pro Phe His Asn Ile His Pro Leu Thr Ile Gly Glu Cys Pro Lys
305                 310                 315                 320

Tyr Val Lys Ser Asn Arg Leu Val Leu Ala Thr Gly Leu Arg Asn Ser
                325                 330                 335

Pro Gln Arg Glu Ser Arg Arg Lys Lys Arg Gly Leu Phe Gly Ala Ile
            340                 345                 350

Ala Gly Phe Ile Glu Gly Gly Trp Gln Gly Met Val Asp Gly Trp Tyr
        355                 360                 365

Gly Tyr His His Ser Asn Glu Gln Gly Ser Gly Tyr Ala Ala Asp Lys
    370                 375                 380

Glu Ser Thr Gln Lys Ala Ile Asp Gly Val Thr Asn Lys Val Asn Ser
385                 390                 395                 400
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Ile Ile Asp Lys Met Asn Thr Gln Phe Glu Ala Val Gly Arg Glu Phe
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Asn Asn Leu Glu Arg Arg Ile Glu Asn Leu Asn Lys Lys Met Glu Asp
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Gly Phe Leu Asp Val Trp Thr Tyr Asn Ala Glu Leu Leu Val Leu Met
        435                 440                 445

Glu Asn Glu Arg Thr Leu Asp Phe His Asp Ser Asn Val Lys Asn Leu
    450                 455                 460

Tyr Asp Lys Val Arg Leu Gln Leu Arg Asp Asn Ala Lys Glu Leu Gly
465                 470                 475                 480

Asn Gly Cys Phe Glu Phe Tyr His Lys Cys Asp Asn Glu Cys Met Glu
                485                 490                 495

Ser Ile Arg Asn Gly Thr Tyr Asn Tyr Pro Gln Tyr Ser Glu Glu Ala
            500                 505                 510

Arg Leu Lys Arg Glu Glu Ile Ser Gly Val Lys Leu Glu Ser Ile Gly
        515                 520                 525

Thr Tyr Gln Ile Leu Ser Ile Tyr Ser Thr Val Ala Ser Ser Leu Ala
    530                 535                 540

Leu Ala Ile Met Met Ala Gly Leu Ser Leu Trp Met Cys Ser Asn Gly
545                 550                 555                 560

Ser Leu Gln Cys Arg Ile Cys Ile
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<400> 3
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<210> 4
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<400> 6
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<210> 7
<211> 39
<212> DNA
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<223> Synthesized oligonucleotide ApaI-H5 (A-Indo).1c

<400> 7
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<210> 8
<211> 37
<212> DNA
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<220>
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atcattatta aacattagag taaagaaata tggatgataa gaacaagagt agtgatattt 180

tgacaacaat tttgttgcaa catttgagaa aattttgttg ttctctcttt tcattggtca 240

aaaacaatag agagagaaaa aggaagaggg agaataaaaa cataatgtga gtatgagaga 300

gaaagttgta caaaagttgt accaaaatag ttgtacaaat atcattgagg aatttgacaa 360

aagctacaca aataagggtt aattgctgta aataaataag gatgacgcat tagagagatg 420

taccattaga gaatttttgg caagtcatta aaaagaaaga ataaattatt tttaaaatta 480

aaagttgagt catttgatta aacatgtgat tatttaatga attgatgaaa gagttggatt 540

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cacgtaggag gataacagga tccccgtagg aggataacat ccaatccaac caatcacaac 780

aatcctgatg agataaccca ctttaagccc acgcatctgt ggcacatcta cattatctaa 840

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<211> 26
<212> PRT
<213> Medicago sativa

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<210> 11
<211> 25
<212> DNA
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<210> 12
<211> 34
<212> DNA
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<220>
<223> Synthesized oligonucleotide supP19-plasto.r

<400> 12
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<210> 13
<211> 20
<212> DNA
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<220>
<223> Synthesized oligonucleotide supP19-1c

<400> 13
atggaacgag ctatacaagg 20

<210> 14
<211> 32
<212> DNA
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<220>
<223> Synthesized oligonucleotide SupP19-SacI.r

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