



Advances and challenges in enveloped virus-like particle (VLP)-based vaccines

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ABSTRACT

Virus-like particles (VLPs) are highly organized particles that self-assemble from viral structural proteins. Like parental viruses, VLPs can be either non-enveloped or enveloped and can be produced in different expression systems depending on their complexity. Over the last three decades, VLPs have developed as a high-priority alternative to traditional vaccines against infectious pathogens due to their safety, simplicity and favorable immunological characteristics to induce both humoral and cellular immune responses. Most of emerging and re-emerging viruses that pose a continuous threat to human health are enveloped, but few vaccines are currently available. Advances in expression technology for complex, enveloped VLPs provide new possibilities to develop potent vaccines against pathogenic enveloped viruses. This review describes major progress and challenges in the production of enveloped VLPs, with respect to the main principles in the assembly and budding process, factors that need to be taken into account for the design strategies and choice of relevant production platforms.

Introduction

Vaccination remains the most cost-effective approach in preventing and controlling infectious viruses which pose a threat to global public health. The currently available preventative viral vaccines are mainly based on inactivated or live attenuated viruses and have shown good efficacy at inducing protective immunity and played an important role in the control or localized eradication of infectious diseases¹. Nevertheless, conventional vaccines have several drawbacks such as safety issues and deviation of the protective immune responses², so safer and more efficient vaccines are in urgent need.

Virus-like particles (VLPs) have been exploited as an alternative to traditional vaccines. VLPs are non-infectious nanostructures composed of viral structural proteins morphologically resembling authentic virions. Based on the structure of viruses, VLPs can be divided into non-enveloped VLPs and enveloped VLPs (eVLPs)³. VLP-based vaccines are considered safer than inactivated or attenuated viruses because a potential incomplete inactivation and reversion of the virus are avoided. They also offer advantages over other subunit vaccines because of their self-adjuvanting properties⁴. Firstly, VLPs have an optimal size (20 to 200 nm) and particulate structure, so they can be easily recognized and absorbed by antigen presenting cells. Secondly, VLPs can be produced in a range of

Table 1. Representative licensed recombinant VLP-based human vaccines.

Target virus	Vaccine name	Company/institution	Year of licensure	Expression system	Reference
HBV (Hepatitis B virus)	Engerix-B®	GlaxoSmithKline	1986	Yeast	Keating <i>et al.</i> (2003) ⁵
HPV (human papillomavirus)	Gardasi Cervarix®	Merck & Co. GlaxoSmithKline	2006	Yeast	Villa <i>et al.</i> (2005) ¹⁷
			2009	Insect	Harper <i>et al.</i> (2004) ²¹
HEV (hepatitis E virus)	Hecolin1	Xiamen Innovax Biotech Co.	2011	E.coli	Li <i>et al.</i> (2005) ²⁹

production systems and this provides flexibility in tailoring manufacturing conditions.

Since the first recombinant VLP-based vaccine, yeast-produced HBsAg vaccine (Engerix-B®), was approved by the Food and Drug Administration (FDA)⁵, hundreds of VLP-based vaccine candidates have been studied, and some have been shown to be efficacious in preclinical and clinical trials. However, only a few of VLP-based vaccines have reached the market as summarized in Table 1 and most of them are targeting non-enveloped viruses. In contrast to non-enveloped VLPs, eVLPs are much more complex in composition. The presence of a host cell-derived membrane provides additional possibilities to integrate heterogenous antigens and adjuvants, but also potential challenges in regulatory approval due to the uncertainty of host cell components. In addition, for different virus families, the origin and composition of envelope vary and usually depend on the virion assembly process in the natural host and the specific cell line used for production. Challenges remain to produce eVLPs with optimal quality, stability and good immunogenicity at high yield.

This review, therefore, discusses the achievements and challenges of the development of eVLP-based vaccines.

Assembly and budding of eVLPs

Formation of virus always requires the presence of all components including viral genome, inner structural proteins, and glycoproteins, but the minimal requirements that drive eVLP formation are not well understood⁶. Generally, self-assembly of eVLPs includes two steps, namely inner structural protein (core, capsid or matrix) formation and then membrane enclosure for further budding.

Assembly and budding of eVLPs may be dependent on viral inner structural protein, envelope glycoproteins or both⁷. In retroviruses, expression of the gag core particle alone can direct eVLP formation and budding⁸. For coronavirus or flavivirus, expression of glycoproteins leads to release of eVLPs morphologically similar to complete virions^{9,10}.

Viral surface antigens, mainly glycoproteins, will be present in the envelope of eVLP, possibly enhancing immunogenicity and acting as an adjuvant. The proper folding, especially glycosylation of these viral antigens plays an important role in the efficacy of eVLP-based vaccines as it is critical for stability, immune recognition,

and pathogenicity¹¹. For example, mutant Zika virus that lacks the N-glycosylation of E protein is attenuated at mammalian and mosquito host¹². Glycosylation of E protein also is a major determinant for viral pathogenesis of Dengue virus (DENV)¹³ and the secretion of tick-borne encephalitis virus (TBEV) VLPs¹⁴.

Like enveloped viruses, eVLPs get the lipid membrane from the host cells through budding. Many viruses including retroviruses, filoviruses, alphaviruses, ortho- and paramyxoviruses are released directly from the cell surface if budding occurs at the plasma membrane. However, for some viruses including flaviviruses and bunyaviruses, budding can also occur on intracellular membranes along the secretory pathway, resulting in the accumulation of virus particles in the lumen of cellular organelles⁷. Depending on the virion assembly process in the natural host, intracellular membrane compartments or some alternative pathway may be involved in secretory of eVLPs.

Efforts to investigate virus budding mechanisms will increase understanding of the formation of eVLP and accelerate the development of eVLP.

Expression platforms for eVLPs

The comparison of the bacteria, yeast, insect, mammalian and plant expression systems and production of several eVLP-based vaccine candidates tested in pre-clinical trials are summarized in Table 2 and Table 3.

Bacteria are the most widely used expression system for manufacturing recombinant proteins, but not a proper platform for eVLP production due to a lot of factors, including the lack of a post-translational modification (PTM) system and protein solubility problems¹⁵.

Eukaryotic expression systems are more suitable for the production of eVLP. Yeast expression systems, especially those based on *Saccharomyces cerevisiae* and *Pichia pastoris*, have advantages such as scalable fermentation, low production costs, and PTM process¹⁶. Two licensed vaccines, the HBV vaccine Engerix-B® and HPV vaccine Gardasil® have been manufactured using this system^{5,17}. Despite the successes, the system has some limitations, such as the different PTM of proteins from mammalian cells and it is generally used for the production of non-enveloped VLPs. Nevertheless, recently its application has been extended to the production of eVLPs. Secretory expression of enveloped human immunodeficiency virus-1 (HIV-1) Gag VLPs using *Saccharomyces cerevisiae*

Table 2. Comparison of different VLP production platforms.

VLP type	Bacteria	Yeast	Insect cells	Mammalian cells	Plants
	Non-enveoped	Non-enveoped Enveoped	Non-enveoped Enveoped	Non-enveoped Enveoped	Non-enveoped Enveoped
Secretory expression	-	+	++	++	+
Speed	+++	+++	++	++	++
Cost	+	+	++	+++	++
Scalability	+++	+++	++	++	++
Yield	+++	++	++	+	++
VLP complexity	+	++	+++	++	++

-, no successful cases reported; +, low; ++, medium; +++, high

Table 3. Production of eVLPs in different expression systems.

Family	Virus	Determinant protein	Expression system	Referrence
Retroviridae	Human immunodeficiency virus I	Gag, Env	Yeast (<i>Saccharomyces cerevisiae</i>) Insect (Sf9) Insect (S2) Mammalian cell (293T) Plant (<i>Nicotiana benthamiana</i>)	Sakuragi <i>et al.</i> (2002) ⁸ Lynch <i>et al.</i> (2010) ³⁰ Yang <i>et al.</i> (2012) ³¹ Crooks <i>et al.</i> (2007) ³² Meyers <i>et al.</i> (2008) ³³
Orthomyxoviridae	Influenza virus A	HA, NA, M1, M2	Insect (Sf9, High Five) Mammalian cell (Vero) Plant (<i>Nicotiana benthamiana</i>)	Krammer <i>et al.</i> (2007) ³⁴ Wu <i>et al.</i> (2010) ³⁵ D'Aoust <i>et al.</i> (2008) ³⁶
Togaviridae	Chikungunya virus (CHIKV)	C, E3, E2, 6k, E1	Yeast (<i>Pichia pastoris</i>) Insect (Sf21) Mammalian cell (293T)	Saraswat <i>et al.</i> (2016) ³⁷ Metz <i>et al.</i> (2013) ³⁸ Akahata <i>et al.</i> (2013) ³⁹
Coronaviridae	Severe acute respiratory syndrome (SARS)	S, E, M, N	Insect (Sf9)	Mortola <i>et al.</i> (2013) ⁹
Paramyxoviridae	Nipah virus	M, G, F	Yeast (<i>Pichia pastoris</i>) Mammalian cell (293T)	Joseph <i>et al.</i> (2016) ⁴⁰ Walpita <i>et al.</i> (2011) ⁴¹
Filoviridae	Ebola virus	VP40, GP, NP	Insect (Sf9) Mammalian cell (293T)	Sun <i>et al.</i> (2009) ⁴² Warfield <i>et al.</i> (2007) ⁴³
Flaviviridae	Dengue virus	prM, E	Yeast (<i>Pichia pastoris</i>) Insect (Sf9) Mammalian cell (COS-1) Mammalian cell (293T) Plant (lettuce)	Liu <i>et al.</i> (2009) ⁴⁴ Kuwahara <i>et al.</i> (2010) ⁴⁵ Chang <i>et al.</i> (2003) ²⁷ Zhang <i>et al.</i> (2011) ¹⁰ Kanagaraj <i>et al.</i> (2011) ⁴⁶
Phenuiviridae	Rift Valley fever virus (RVFV)	Gn, Gc, NP	Insect (S2) Insect (Sf21, Sf9) Mammalian cell (293T)	de Boer <i>et al.</i> (2010) ²² Liu <i>et al.</i> (2008) ⁴⁷ Habjan <i>et al.</i> (2009) ⁴⁸
Arenaviridae	Lassa virus (LASV)	GPC, NP, Z	Mammalian cell (293T/17) Mammalian cell (MDCK- II)	Branco <i>et al.</i> (2010) ⁴⁹ Schlie <i>et al.</i> (2010) ⁵⁰

spheroplasts has been achieved⁸. However, in the case of HIV-2, *Saccharomyces cerevisiae* cells fail to support the multimerization of the Gag protein into VLPs and particle budding from the membrane¹⁸. Therefore, it is important to note that yeast is not always suitable for eVLP production.

Higher eukaryotic expression systems, insect and mammalian cells, perform more complex PTM and support most aspects of the virus life cycle. So, they have been extensively used for both intracellular and secretory production of eVLPs.

The baculovirus-insect cell expression system (BVES) is the most commonly used system for producing a wide variety of eVLPs as summarized in Table 3. The most commonly used insect cell lines are derived from

Spodoptera frugiperda (Sf9 and Sf21) and *Trichoplusia ni* (Tn5, commercially known as High Five™), which grow optimally at 27°C and do not require CO₂, making scale up of protein production feasible^{19,20}. A licensed HPV vaccine Cervarix® has been manufactured using this system²¹. However, some issues limit the application of this system for eVLP production. One limitation is the yields of secreted and membrane-bound proteins are often low, probably because of the side effects of baculovirus infection on the secretory machinery of host insect cells. Furthermore, lepidopteran cells are unable to produce glycoproteins with structurally authentic mammalian N-glycans. Another main disadvantage is contamination of progeny baculovirus¹⁹. Stably transformed insect cells have emerged as attractive alternatives, particularly for the production of secreted

glycoproteins. The *Drosophila melanogaster* Schneider 2 (S2) cell line, has been widely used. It allows inducible expression and secretion of soluble glycoproteins under non-lytic conditions and serum-free medium²⁰. Rift Valley fever (RVFV) VLP composed of Gn and Gc glycoproteins has been obtained using S2 cell line²².

The mammalian cell is another platform that can produce both non-enveloped and enveloped VLPs. The ability to perform complete PTMs of recombinant proteins is the main advantage of mammalian cells. Several mammalian cell types, such as human embryonic kidney 293 (HEK293), Chinese hamster ovary (CHO) and baby hamster kidney (BHK-21) cell lines are extensively utilized for eVLP production. Based on these cell lines, most of the eVLP-based vaccines have been studied as summarized in Table 3. However, the high production cost, low protein yield, and potential safety concerns resulted from contamination by adventitious agents impose restrictions on commercialization of mammalian VLPs²³.

Plant expression systems have also been used for VLP production. A plant can produce large quantities of recombinant protein at low cost, provide complex but distinctive PTM, and bring a low-risk of introducing adventitious human pathogens²⁴. Many plant species have been used for recombinant protein production, e.g., tobacco, potato, carrot, and tomato. Although plant-specific glycans provide a unique advantage for producing eVLPs with high immunogenicity and the protein synthesis and folding pathway are conserved between animals and plants, there are some differences in protein glycosylation. The improper glycoforms may reduce efficacy or cause potential adverse effects.

Challenges and solutions in the development of eVLPs

Technical challenges still remain in the manufacture of eVLP-based vaccines, regarding the design, purification, and storage.

Firstly, the stability of the VLP-based vaccine is one of the most significant issues. VLPs are multimeric structures that are generally more stable than subunit vaccines, however, the lack of the viral genome makes them unstable when the conditions change, especially during downstream processing (DSP)²⁵. Generally, eVLPs having a host-derived envelop are more sensitive to the external environment than the protein-only VLPs. Variations in conditions, e.g., temperature, shear force and chemical treatment can destroy the integrity and stability of the particles, this structural destruction further leads to the reduction in immunogenicity of eVLPs. In some cases, VLPs have been modified in order to improve particle thermostability. One common technique is the introduction of the stabilizing mutations. A study of poliovirus type 3 VLPs shows that

VLPs having stabilizing mutations within the coat proteins retain the native antigenic conformation and the repetitive structure of the original virus particle and are more stable than the wild type VLPs²⁶.

Secondly, the expression level of viral proteins in different platforms vary considerably. In general, the secretory expression of glycoproteins is difficult. Since budding from cell membrane to get envelop is a key step during the eVLP formation process, if eVLP is not efficiently secreted, a cell lysis or other extraction step might be required, and these steps increase the difficulty for further purification. A common way to improve expression level of transmembrane glycoproteins is to delete or replace the transmembrane region which anchors the protein in the membrane. Replacement of the stem-transmembrane domain of DENV2 E protein which contains a strong ER retention signal with the corresponding region of Japanese encephalitis virus (JEV) provides extracellular secretion of eVLPs²⁷. This is not always useful, for instance, if membrane-integration dependent oligomerization is required for functionality or immunogenicity. Another way to improve the secretion potential for eVLP is the introduction of signal peptide. As for the expression of DENV VLP, cells co-expressing prM and E proteins fail to secrete eVLPs. When adding a JEV signal sequence at the N terminal of prME protein, cells can effectively secrete eVLPs, indicating signal peptide is one of the important factors that influence downstream protein translocation and topology¹⁰. It should be noted that extrinsic signal peptides will change the characteristic of the protein in some cases.

Finally, on the DSP, numerous impurities co-purified with eVLPs pose a daunting challenge. The process-related contaminants are predominantly attributed to host cell impurities such as cell debris, host cell proteins, DNA, and lipids. A typical example is the baculovirus from the BVES. The similar biophysical features, e.g., size, electrostatic and architecture between eVLPs and the enveloped baculoviral particles make the DSP complicate. Baculoviruses have shown adjuvant activity and if not removed or inactivated, would induce undesirable synergistic effects¹⁹. To meet the criteria on the security issue, purified eVLPs should be inactivated to eliminate baculovirus infectivity, but this strategy may affect the antigenicity of eVLPs. Alternatively, efforts to improve the purity of product by stepwise purification processes including clarification, intermediate purification, and polishing without sacrificing the immunogenicity of eVLPs are more significant. Diverse of purification methods such as centrifugation, precipitation, ultrafiltration and chromatography have been developed²⁸.

Conclusion

VLPs have been developed as an alternative to traditional vaccines based on their safety, flexibility and distinctive immunogenic properties. Challenges

encountered at the bioprocessing technology and dealing with regulatory issues to bring eVLPs to the market. The further understanding of VLP assemblies and advances in the technologies will lead to the development of a series of novel, stable and efficient eVLP-based vaccine candidates in the near future.

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