

## **Papillomavirus virus-like particles as vehicles for the delivery of epitopes or genes**

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**Summary.** Papillomaviruses (PVs) are simple double-strand DNA viruses whose virion shells are  $T = 7$  icosahedrons and composed of major capsid protein L1 and minor capsid protein L2. L1 alone or together with L2 can self-assemble into virus-like particles (VLPs) when expressed in eukaryotic or prokaryotic expression systems. Although the VLPs lack the virus genome DNA, their morphological and immunological characteristics are very similar to those of nature papillomaviruses. PV VLP vaccination can induce high titers of neutralizing antibodies and can effectively protect animals or humans from PV infection. Moreover, PV VLPs have been good candidates for vehicles to deliver epitopes or genes to target cells. They are widely used in the fields of vaccine development, neutralizing antibody detection, basic virologic research on papillomaviruses, and human papillomavirus (HPV) screening. Besides the structural biology and immunological basis for PV VLPs used as vehicles to deliver epitopes or genes, this review details the latest findings on chimeric papillomavirus VLPs and papillomavirus pseudoviruses, which are two important forms of PV VLPs used to transfer epitopes or genes.

### **Abbreviations**

*PV* Papillomavirus; *VLPs* virus-like particles; *cVLPs* chimeric virus-like particles; *HPV* human papillomavirus; *DCs* dendritic cells; *LCs* Langerhans cells; *IL* interleukin; *TNF* tumor necrosis factor; *INF* interferon; *CTLs* cytotoxic T lymphocytes

### **Introduction**

Papillomaviruses (PVs), members of the family *Papillomaviridae*, are simple double-strand DNA viruses that infect squamous or mucosal epithelia and produce

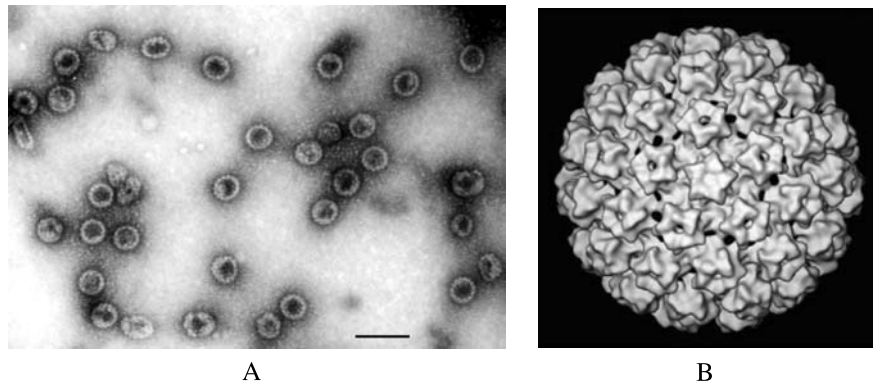
a range of epithelial neoplasms, both benign and malignant, in most animals and humans. Until now, 118 PV types have been completely described, and many new types have been detected by preliminary data such as subgenomic amplicons. More than 80 human papillomavirus (HPV) types have been defined, and many of these HPV types are ubiquitous and globally distributed ([http://hpv-web.lanl.gov/stdgen/virus/cgi-bin/hpv\\_organisms.cgi?dbname=hpv](http://hpv-web.lanl.gov/stdgen/virus/cgi-bin/hpv_organisms.cgi?dbname=hpv)). The detailed information on PV classification has been reviewed by de Villers EM et al. [15]. The papillomavirus virion shells consist of L1 protein (the major capsid protein) and L2 protein (the minor capsid protein). L1 protein alone or together with L2 protein can self-assemble into virus-like particles (VLPs) when expressed in eukaryotic or prokaryotic expression systems. zur Hausen et al. [95] observed that infection with HPVs might be responsible for human cervical cancer, which was confirmed by numerous subsequent studies. Because cervical cancer is the second-most common cancer in women, more attention has been paid to HPV research, especially on PV VLP-based epitope or gene delivery [73, 82], effective prophylactic and therapeutic vaccine development [28, 58], the life cycle of PVs [17], the mechanism of virus evasion from host-cell control [96], and the epidemiology of HPV infections [2, 53]. We will focus on the structural biology and immunological characteristics of PV VLPs and the latest studies on chimeric papillomavirus VLPs and papillomavirus pseudoviruses, which are two important forms of PV VLPs used as vehicles to deliver epitopes or genes.

### **Structural biology and immunological characteristics of papillomavirus VLPs**

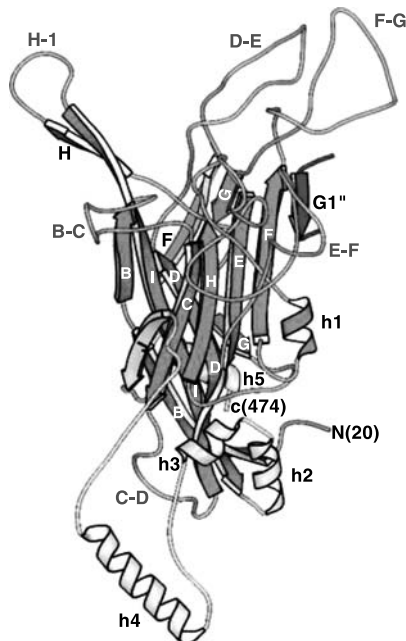
#### *Structural biology characteristics of papillomavirus VLPs*

The papillomavirus virion contains 72 pentamers of L1 protein, centered on the vertices of a  $T = 7$  icosahedral lattice [83], while L2 protein is present at about 1/30 of the abundance of L1 [38]. The diameter of the PV particles is about 50–60 nm. Although the PV VLPs lack the virus genome DNA, they are also composed of 72 pentamers of L1 protein, and their morphological characteristics are very similar to those of the PV virions as well (Fig. 1).

Because it is very difficult to obtain large amounts of PV viruses by *in vitro* culture and to obtain suitable crystals of PV virions or VLPs, the crystallographic study of PV particles has been limited. Chen et al. [13] truncated the 10 N-terminal residues of HPV16 L1 to overcome the irregularity of the shells formed by the intact L1 for successful crystallization and expressed the truncated HPV16 L1 in *E. coli*. The truncated L1 could assemble into a 12-pentamer,  $T = 1$  icosahedron, named “small VLP”, whose outer diameter was 318 Å, and the X-ray crystallographic analysis was performed at 3.5 Å resolution. The L1 monomer contains 12  $\beta$ -strands, 6 loops, and 5 helices (Fig. 2). These loops extend towards the outer surface of the capsid, and the loop sequences are quite variable among different papillomavirus types [13]. Mutagenesis of the hypervariable loop domains has revealed the binding sites of several neutralizing monoclonal antibodies against



**Fig. 1.** Structure of papillomaviruses (PVs) and PV virus-like particles (VLPs). **A.** Electron micrograph of negative-stained human papillomavirus (HPV)16 L1 VLPs with 1% uranyl acetate solution. Bar represents 100 nm. **B.** The structure model of bovine papillomavirus(BPV) [83]



**Fig. 2.** The human papillomavirus (HPV) 16 L1 monomer structure [13]. Secondary structural elements are labeled, with letters *B–J* for  $\beta$ -strands and *h1–h5* for the 5  $\alpha$ -helices. Loops between strands are labeled *B–C*, *C–D*, etc. The first and the last residues are marked *N* (20) and *C* (474), respectively. A primed label (*F'*) denotes a segment of polypeptide chain from the clockwise neighbor within the pentamer; a double-primed label (*G''*), one from the anticlockwise neighbor

HPV11 [47–49] or HPV6 [50], showing that many neutralizing epitopes lie in the loop domains.

#### *Immunological characteristics of papillomavirus VLPs*

Though papillomavirus VLPs lack the virus genome DNA, they are structurally and immunologically similar to the infectious viruses as judged by electron microscopic imaging studies and their ability to bind type-specific and conformation-dependent monoclonal antibodies. Many studies have demonstrated that

vaccination with PV VLPs by different methods can induce high titers of type-specific neutralizing antibodies (IgG and IgA) and effectively protect animals [5, 37, 38, 41, 44, 49, 57, 60, 79] and humans [1, 6, 7, 19, 21, 27, 32, 40, 54, 64, 65, 71, 87, 99] from papillomaviruses infection. The phase II randomized, multicentre, double-blind placebo-controlled clinical trials of bivalent (HPV16 and HPV18) VLPs vaccine and quadrivalent (HPV6, 11, 16 and 18) VLPs vaccine, which were conducted by GlaxoSmithKline and Merck, respectively, have been finished. Both vaccines were very effective in protecting women against incident and persistent HPV infections [32, 87]. What's more, since the VLPs do not contain the viral genome, they are not harmful. So PV VLPs are considered good candidates for both prophylactic and therapeutic vaccines against PV infection. Besides their safety and high immunogenicity, the following characteristics make PV VLPs good candidates for vehicles to deliver epitopes or genes.

Papillomavirus VLPs themselves are good “adjuvants” and can induce strong T-helper cell responses. A lot of studies have demonstrated that PV VLPs can activate dendritic cells (DCs) [3, 42, 70, 90–93]. DCs are a family of professional antigen-presenting cells (APCs) with a unique capacity to initiate and modulate cell-mediated immune responses. A wide range of stimuli, such as infectious virus and inflammatory cytokines, can induce DC maturation, which is associated with the up-regulation of co-stimulatory molecules (reviewed in [39, 66]). The activated DCs then migrate to regional lymph nodes and induce T cell activation. Immature DCs can efficiently bind and rapidly internalize PV VLPs, which is followed by DC maturation. DCs activated by PV VLPs can induce the secretion of the proinflammatory cytokines interleukin (IL)-6 and tumor necrosis factor (TNF)- $\alpha$ , IL-8, and interferon(INF)- $\alpha$  [42, 43]. Although Langerhans cells (LCs) can effectively bind and take up HPV VLPs, they can not be activated by HPV VLPs [22–24]. But a recent study showed that heterologous PV VLPs immune complexes could activate human Langerhans cells [25]. Both DCs and LCs incubated with the VLPs immune complexes could up-regulate surface activation markers and increase secretion of IL-12 p70 and IL-15. So both DCs and LCs can be activated by PV VLPs using certain immunization strategies. In addition, papillomavirus VLPs could bind to a wide variety of cell types [51, 88] through certain molecules, such as CD16 [90], heparan sulfate [14, 18, 30, 34, 68], and  $\alpha_6$  integrin [20, 94]. Therefore, the epitopes or genes encapsidated in the VLPs will be well delivered to certain immunocytes.

#### *Chimeric papillomavirus VLPs*

Chimeric papillomavirus VLPs, which means PV VLPs composed of capsid proteins (L1 or L1/L2) fused with foreign epitopes or polypeptides in certain areas, are one of the important forms of PV VLPs used to deliver epitopes or polypeptides. Some of the different strategies for construction of chimeric PV VLPs are summarized in Table 1.

It is clear that mutants of PV L1 with some residues in the C-terminus truncated can also self-assemble into VLPs [59]. As a result, epitopes or short polypeptides,

**Table 1.** Different construction strategies of chimeric papillomavirus VLPs

Construction strategy	Epitopes or polypeptides	References
Fusing epitopes or polypeptides to the C terminus of the C-terminal truncated PVs L1	HPV16 E7 (partial and full length gene)*	[29, 52, 74]
	HPV16 E7 CTL epitope (RAHYNIVTF)	[62]
	HIV-III <sub>B</sub> gp160 CTL epitope (RGPGRAFVTI)	[62]
	A fusion polypeptide containing HPV16 E7 B cell epitope (QAEPD), HPV16 E7 CTL epitope (RAHYNIVTF), HIV-III <sub>B</sub> gp120 CTL epitope (RGPGRAFVTI), HIV-III <sub>B</sub> Nef CTL epitope (PLTFGWCFKL), and HIV-III <sub>B</sub> RT CTL epitope (VIYQYMDDL)	[45]
	P1A epitope (LPYLGWLVF)	[55]
Replacing certain sequences of PVs L1 by epitopes or polypeptides	HPV16 E7 (partial and full length gene)	[52]
	SV40 tag	[52]
	HPV16 L2 (LVEETSFIDAGAP)	[86]
	HIV-1 gp41 B cell epitope (LELDKWAS)	[76]
	HIV-1 gp41 epitope (ELDKWA)	[98]
Mouse CCR5 epitope (HYAANEWVFGNIMCKV)	[10]	
Inserting epitopes or polypeptides into some areas of the PVs L1	HPV16 E7 <sub>1-50</sub>	[52]
	HBc epitope (DPASRE)	[72]
	HIV-1 gp41 B cell epitope (LELDKWAS)	[76]
Fusing epitopes or polypeptides to the C terminus of PV L2	HPV16 E7 (partial length gene)	[16, 25, 31, 70, 89]
Replacing certain sequences of PVs L2 by epitopes or polypeptides	HPV16 E1E2E7 fusion protein	[81]

PVs papillomaviruses, HPV human papillomavirus, CTLs cytotoxic T lymphocytes, HIV human immunodeficiency virus, HBc hepatitis B core antigen

\*The description or amino acid sequence of the polypeptide is parenthesized

up to 60 amino acids, can be fused to the C-terminus of the truncated PV L1 without disrupting the assembly of VLPs [52]. Until now, many epitopes or short polypeptides have been added to the C-terminus of the truncated PV L1 (see Table 1), and these chimeric VLPs (cVLPs) can induce good immune responses against not only the epitopes or polypeptides but also the VLP shells. Because the size of epitopes or polypeptides fused to the C-terminus of the truncated PV L1 is limited, large proteins cannot be delivered using this strategy. However, these large proteins can be delivered well by replacing certain areas of L1 or L2 that are not important for VLPs assembly, or by fusing them to the C-terminus of L2 (see Table 1). L1 is the major capsid protein, and it has been shown that some residues in the L1 protein, such as Asp<sub>202</sub>, Cys<sub>175</sub>, and Cys<sub>428</sub> of HPV16 L1, are very important for VLP formation. Therefore, those important residues should not be included in the target replacing area. However, there may be a few potent important residues in L1. When some areas of L1 were replaced by large proteins, the efficiency of VLP assembly has been reduced to some extent [76]. While L2

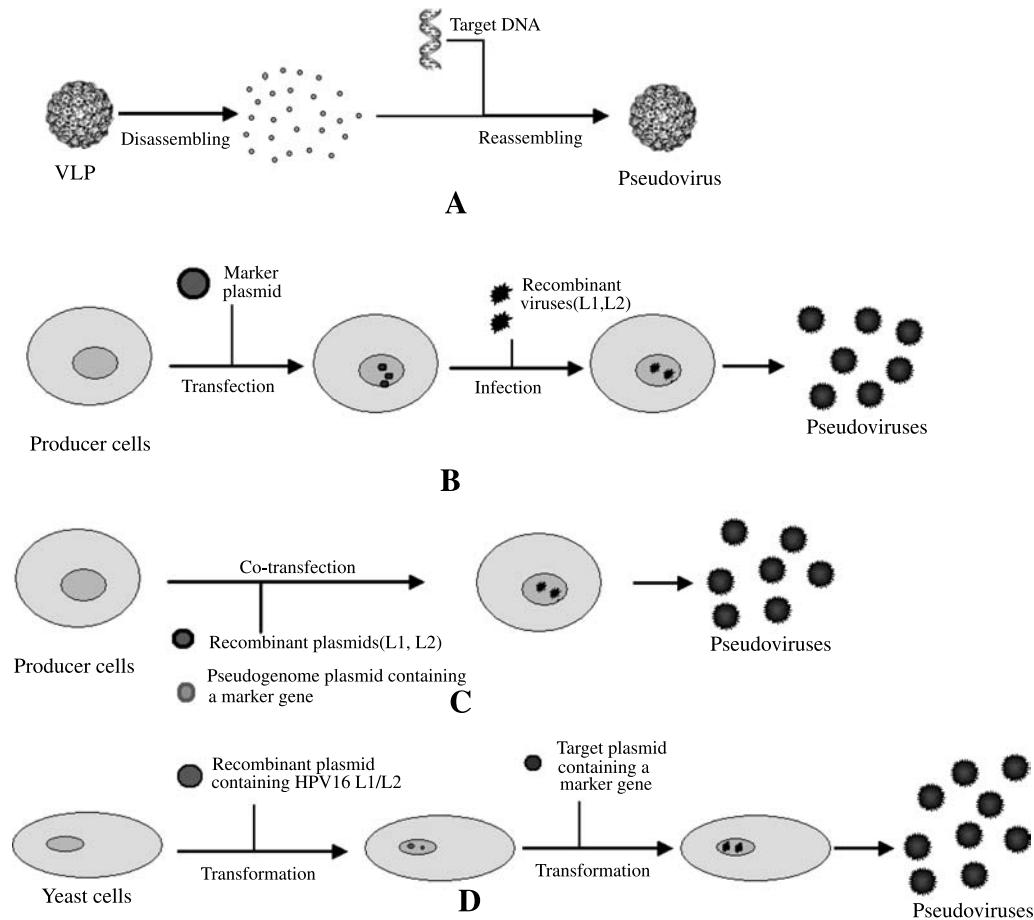
is the minor capsid protein, it is not as important as L1 for VLPs assembly. So replacing some sequences of L2 by large genes or fusing large proteins to the C-terminus of L2 may have less influence on VLP formation than that of L1. Tobery et al. [81] showed that replacing the area of HPV16 L2<sub>70–390</sub> by HPV16E1E2E7, approximately 130 kDa, did not disrupt VLP assembly and that the chimeric VLPs had good immunogenicity. In addition, co-expression of L1 and L2 can enhance the efficiency of VLP formation compared with L1 alone [38]. However, in HPV16 VLPs the ratio of L1 to L2 molecules is about 30:1 [38], that is to say an L1-epitope fusion would theoretically result in 360 copies of the epitopes being delivered to the cytosol of an APC, compared to the delivery of 12 copies of the epitopes by an L2-epitope fusion.

Besides the construction strategies described above, there is another L1-loop-based construction strategy. Structural biology research on PV VLPs has shown that the loop regions of L1 are exposed on the outer surface of VLPs. Some epitopes fused into these loop regions by insertion or replacement might be displayed on the outer surface of cVLPs, so these epitopes could be efficiently delivered and well recognized by the immunocytes, and could induce epitope-specific immune responses [72] (Table 1).

Given that cervical cancer is one of the leading causes of female cancer mortality worldwide, especially in developing countries, many studies are focused on the development of effective prophylactic and therapeutic vaccines. It has been well demonstrated that expression of early protein E7 is required for the maintenance of the proliferative state of HPV-infected cells [80, 85]. E7 is considered a tumor antigen and a potential target for activated T cells in the development of strategies for immune therapy, and many forms of E7 vaccines, such as purified protein; DNA vaccine; recombinant vaccinia virus and chimeric VLPs, have been developed as well. PV L1E7 or L1/L2E7 cVLPs are considered candidates for “dual-purpose” vaccines inducing neutralizing antibodies, which prevent infection, and cytotoxic T lymphocytes (CTLs) that can eradicate established infections. Many studies showed that immunization of a few different forms of L1E7 or L1/L2E7 cVLPs could induce both strong CTL responses and high-titer, type-specific neutralizing antibodies against the VLP shells (see the related references listed in Table 1).

### **Papillomavirus pseudoviruses**

Besides the chimeric papillomavirus VLPs, papillomavirus pseudoviruses are another important form to deliver genes using PV VLPs shells. Although the DNA molecules encapsidated in PV pseudoviruses are not PV genome DNA, PV pseudoviruses have very similar structural and immunological characteristics to native PVs. PV pseudoviruses could be produced by the following methods illustrated in Fig. 3. Firstly, PV pseudoviruses could be generated by a cell-free system *in vitro* [36, 56, 75, 82, 97] (Fig. 3A). PV VLPs were able to package unrelated plasmid containing target genes by the following process: The purified PV VLPs could be disassembled by different methods, such as incubation of VLPs with 5% 2-mercaptoethanol at 4 °C for 16 h or with 50 mM Tris–HCl buffer



**Fig. 3.** Different strategies of producing papillomavirus (PV) pseudoviruses. **A.** Using a cell-free system to produce pseudoviruses *in vitro*. Purified PV virus-like particles (VLPs) were disrupted under certain conditions. The disassembled VLPs were co-cultured with the plasmids containing target genes and could reassemble into pseudoviruses. **B.** Recombinant-virus-based production of pseudoviruses. The producer cells were transfected by the plasmid containing one marker gene. Then the transfected cells were infected by recombinant viruses containing capsid proteins L1 or L1/L2, which was followed by the formation of pseudoviruses. **C.** Producer cells were co-transfected by two recombinant plasmids containing capsid proteins L1 and L2, respectively, and the pseudogenome plasmid containing a marker gene, then the pseudoviruses were produced by the transfected producer cells. **D.** Producing pseudoviruses in *Saccharomyces cerevisiae*. Yeast cells were transformed with a recombinant plasmid containing HPV16 L1/L2 and subsequently with the target plasmid containing a marker gene. Pseudoviruses then could be generated by yeast cells transformed with both plasmids

(pH 7.5) containing 150 mM NaCl, 1 mM EGTA and 20 mM dithiothreitol at room temperature for 30 min. Then the plasmid containing the target gene was co-cultured with disassembled VLPs. The preparation was then incubated with CaCl<sub>2</sub> (25 mM) and 20% dimethyl sulfoxide at room temperature for 1 h to form

the pseudoviruses. The second method to produce pseudoviruses was based on certain recombinant viruses, and is illustrated in Fig. 3B. HPV16 pseudoviruses were generated in hamster BPHE-1 cells by using a recombinant Semliki Forest virus [67], and those of HPV 33 and BPV 1 were produced in COS-7 or COS-1 cells by using a recombinant vaccinia virus [73, 84, 46, 100, 101]. The third method has been reported very recently and it seems to be a better method with higher production efficiency [8, 9, 61] (Fig. 3C). Briefly, producer cells, including C127, HaCaT, and 293TT, were co-transfected by two recombinant plasmids containing L1 and L2, respectively, and the pseudogenome plasmid containing a marker gene, then the pseudoviruses were produced by the transfected producer cells. Buck et al. [8] showed that the titers of pseudoviruses produced by this method could reach the level of several billion transducing units per milliliter, which was at least 10-million-fold greater than that of the previous method [67]. Finally, infectious HPV16 pseudoviruses could also be generated in yeast using the two-step-transformation method [69] (Fig. 3D). *Saccharomyces cerevisiae* strain 1699 was transformed with a yeast expression plasmid containing HPV16 L1/L2 so that it could produce HPV16 VLPs, which was followed by being transformed with a target plasmid containing the green fluorescent protein (GFP) gene as a marker gene. Subsequently, the yeast cells transformed with both plasmids could generate HPV16 pseudoviruses.

Although PV VLPs composed of L1 alone can package unrelated plasmid DNA *in vitro* [4], the efficiency is enhanced to a great extent by co-expressing L1 and L2 in the procedure of pseudovirus production [36, 67, 84, 100]. In addition, Zhao et al. showed [101] that BPV1 E2 protein could enhance the packaging of plasmid DNA into BPV1 pseudoviruses. Pseudogenome encapsidation within L1/L2 capsids was largely sequence-independent, and plasmids completely lacking PV sequences could be packaged efficiently, providing they were less than 8 kb in size. PV pseudoviruses were structurally indistinguishable from native viruses and could efficiently infect many cell types [82]. Moreover, these infections of PV pseudoviruses could be inhibited by type-specific anti-serum against the same type of PV VLPs [36, 67, 82, 84]. So, if the plasmids encapsidated by the PV pseudoviruses contained a certain marker gene, such as GFP [73, 82, 69],  $\beta$ -galactosidase [82, 36, 84, 77], or secreted alkaline phosphatase (SEAP) [8, 61, 84], these pseudoviruses could be used to analyze the papillomavirus internalization pathways, to detect the level of neutralizing antibodies, and to test the reactivity of human sera in population-based HPV screening. What's more, Pastrana et al. [61] showed that the pseudovirus-based papillomavirus neutralization assay was high-throughput and appeared to be as sensitive as, but more specific than, a standard VLP-based enzyme-linked immunosorbent assay (ELISA).

It is clear that using appropriate adjuvants can enhance vaccine-induced specific immune responses. Oh et al. [56] showed that the mucosal and systemic immunogenicity of HPV16 VLPs could be enhanced by encapsidating IL-2 gene adjuvant. Vaccination of the pseudoviruses encapsidated with IL-2 gene could elicit much higher titers of vaginal and salivary HPV16 L1-specific IgA and serum IgG,



IgG<sub>1</sub>, and IgG<sub>2a</sub> antibodies than those of HPV16 VLPs alone. In addition, PV pseudoviruses encoding IL-2 could fully restore mucosal and systemic immune responses to vaccinations in aged mice [26]. In aged mice, immunization with PV pseudoviruses encapsidating the IL-2 gene could activate specific Th cells, induce specific antibodies and CTL responses, and protect against mucosal viral challenge.

PV VLPs were shown to induce strong T-helper responses and the concurrent T-helper responses to the VLPs might enhance the CTL responses against the antigen encoded by the plasmid in the pseudoviruses. Shi et al. [75] showed that PV pseudoviruses could efficiently pseudoinfect mucosal and system lymphoid tissues and reach Peyer's patches, lamina propria, and the spleen when administered orally. By systemic immunization, PV pseudoviruses induced stronger CTL responses than plasmid DNA vaccines alone, and by oral immunization, pseudoviruses could generate specific mucosal and systemic CTL responses and protect mice against mucosal challenge. Women infected with human immunodeficiency virus (HIV) are at a greater risk of being co-infected with HPV. HIV-positive women have a 3–5 times greater risk of developing intraepithelial lesions and a 3–4.5 times greater risk of developing invasive neoplasia [63]. Zhang et al. [97] showed that oral immunization with PV pseudoviruses encoding HIV-1 Gag could induce mucosal and systemic Gag-specific CTL responses which could effectively lyse Gag-expressing target cells. Furthermore, pseudovirus vaccination could generate Gag-specific IFN- $\gamma$ -producing T cells, serum IgG, and mucosal IgA. It would seem that PV pseudoviruses may be better candidates for prophylactic and therapeutic vaccines to some infectious diseases.

In summary, PV pseudoviruses are valuable tools for determination of neutralizing antibody and analysis of papillomavirus internalization pathways and are good candidates for vaccine development.

## Conclusions

Papillomavirus VLPs are good vehicles for the efficient delivery of epitopes or genes to target cells. Now, chimeric papillomavirus VLPs and papillomavirus pseudoviruses have been widely used in the fields of basic virological studies of papillomavirus, development of various vaccines, neutralizing antibody detection, and population-based HPV screening. In order to enhance vaccine immunogenicity, there are still other strategies based on PV VLPs, such as fusing adjuvant molecules to the capsid proteins to form chimeric VLPs (our unpublished data), encapsidating nonmethylated CpG motifs into VLPs derived from the hepatitis B core antigen or the bacteriophage Q $\beta$  [78], chemically binding adjuvant molecules (subunit B of cholera toxin [35]) or antigenic peptides (TNF- $\alpha$  [11, 12], epitope of influenza type A M2 protein [33]) to PV VLPs. What's more, DNA molecules can directly bind to PV VLPs to form a VLP/DNA complex, which is another way to transfer genes using PV VLPs [4]. It is expected that more types of PV VLP-based vaccines with higher immunogenicity and better epitopes or gene delivery systems will be developed in the future.

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