



Published in final edited form as:

Immunol Cell Biol. 2009 ; 87(4): 287–299. doi:10.1038/icb.2009.13.

Developing vaccines against minor capsid antigen L2 to prevent papillomavirus infection

Balasubramanyam Karanam^{1,*}, Subhashini Jagu^{1,*}, Warner K. Huh², and Richard B.S. Roden^{1,#}

¹ Department of Pathology, Johns Hopkins University, Baltimore, MD 21231, USA

² Department of Obstetrics and Gynecology, University of Alabama at Birmingham, Birmingham, AL 35233

Abstract

A subset of human papillomavirus (HPV) genotypes is responsible for ~5% of all cancer deaths globally, and uterine cervical carcinoma accounts for the majority of these cases. Their impact is greatest for women who do not have access to effective secondary preventive measures, and consequently over 80% of cervical cancer deaths worldwide occur in Developing nations. The understanding that persistent infection by this ‘oncogenic’ subset of HPV genotypes is necessary for the development of cervical carcinoma has driven the development of preventive vaccines. Two preventive vaccines comprising recombinant HPV L1 virus-like particles (VLPs) have been licensed. However, the current cost of these vaccines precludes global delivery, and they target only two of the ~15 known oncogenic HPV types, although ~70% of cervical cancer cases are attributed to these two types and there is evidence for some degree of cross-protection against other closely related types. A possible approach to broader immunity at lower cost is to consider vaccination against L2. L2 vaccines can be produced inexpensively and they also have the promise of conferring much broader cross-type protective immunity than observed with L1 VLP immunization. However, L2 vaccine development lags behind L1 VLP vaccines and several technical hurdles remain.

THE CASE FOR PROPHYLACTIC VACCINATION AGAINST HPV

The precursor of cervical cancer is high grade cervical intraepithelial neoplasia (CIN, also known as squamous intraepithelial lesion SIL). Ablation or cold knife excision of these precursors is generally curative, thereby halting progression to invasive cervical cancer. Since the ‘Pap’ smear was introduced by Dr. Georgios N. Papanicolaou in the 1940s to detect high grade CIN, numerous studies have demonstrated higher mortality rates in unscreened or under-screened women and the efficacy of secondary prevention for the reduction of cervical cancer incidence. Unfortunately, cervical cancer therefore disproportionately impacts under-developed nations that lack such cytologic screening programs wherein 83% of cervical cancer cases occur. National screening programs have shown that Pap screening for high grade CIN decreases the occurrence of cervical carcinoma

[#]Correspondence to: Richard Roden, Department of Pathology, Johns Hopkins University, Room 308, Cancer Research Building 2, 1550 Orleans St, Baltimore, MD 21231 USA, roden@jhmi.edu, telephone: (410) 502-5161, fax: (443) 287-4295.

^{*}Equal contributors

Conflict of Interest Statement: RBSR is a paid consultant of Merck & Co, and Knobbe, Martens, Olson and Bear LLC. SJ and RBSR have received unrestricted educational grant funding from GSK. RBSR is an inventor on L2 patents licensed to PaxVax, Inc. and Acambis, Inc. The terms of these arrangements are being managed by Johns Hopkins University in accordance with its conflict of interest policies. WKH is a paid consultant and speaker for Merck Inc and a paid consultant for GlaxoSmithKline. He has received research support from both companies.

by over 60%. The effectiveness of such programs could be further increased by reducing the testing interval and improve the performance characteristics of testing. The Pap test has remained virtually unchanged for 50 years as part of national screening programs until the advent of molecular testing for HPV. Molecular testing for HPV significantly improves the performance of cytologic screening and full implementation of such a combined screening approach can further reduce rates of cervical cancer. However, such programs are expensive and impractical in large parts of the world. Furthermore, HPV infection causes significant morbidity in some patients including other anogenital and some head and neck cancers. Therefore prevention of HPV infection with a low cost vaccine is preferable to large screening programs, particularly in low resource settings. Maximum benefit is obtained by combining both interventions, but this has significant cost implications and reduces the performance of screening, necessitating careful consideration of the optimal approaches to implementation.

The strength and diversity of evidence has led to general recognition of oncogenic HPV infection as the necessary cause of cervical cancer. Dr. Harald zur Hausen had first postulated this link during the 1970s, and in the early 1980s his group identified the two most important oncogenic HPV genotypes, HPV16 and HPV18 (1,2). For this seminal work and his continued contributions, Dr zur Hausen was awarded the 2008 Nobel prize in physiology or medicine. Building of the case for the full recognition of oncogenic HPV as the central etiologic agent for cervical cancer took many years of effort by researchers worldwide. The evidence implicating oncogenic HPV infection includes basic laboratory studies of viral pathogenesis, large epidemiologic studies, and successful screening and preventive vaccine interventions.

It is important to recognize that the majority of HPV infections do not lead to cervical cancer, i.e. HPV infection *per se* is not a sufficient cause and that additional changes and co-factors contribute to carcinogenesis. HPV is a small, nonenveloped DNA virus that induces generally benign epithelial tumors (warts) of skin or mucosa. Most such HPV-induced lesions show limited growth and often regress spontaneously. Of the nearly 200 genotypes of HPV that have been identified, more than fifteen have been shown to induce tumors that may eventually progress to carcinomas (3). In particular, 53.5%, 17.2%, and 6.7% of cervical cancer cases worldwide are attributed to the genital types HPV16, HPV18 and HPV45 respectively (4). The remaining 12+ oncogenic genotypes (HPV31, HPV33, HPV52, HPV58, HPV35, HPV59, HPV56, HPV51, HPV39, HPV68, HPV73, and, HPV82 in descending order of importance, and probably additional less well characterized types) are considered to cause the remaining 30% of cervical cancer cases in total, with less than 3% for any one of these minor types (4). It should be noted that this type distribution does not hold for all HPV-associated cancers. For example the contribution of HPV16 is much greater for head and neck cancer (5), as well as vaginal or vulval cancers (6), whereas HPV18 and HPV45 show greater importance in cervical adenocarcinoma (7). Papillomaviruses exhibit strict tropism, and therefore preclinical studies of vaccines have been limited mainly to a few animal papillomavirus models, notably bovine papillomavirus (BPV) infection of cows, cottontail rabbit papillomavirus (CRPV) or rabbit oral papillomavirus (ROPV) infection of rabbits and canine oral papillomavirus (COPV) infection of dogs.

Over 99% of cervical cancers and their precursor lesions CIN, contain human papillomavirus (HPV) DNA (8,9). The HPV genotypes found in cancer cells have immortalizing activity for primary human keratinocytes (10), and the viral oncoproteins, E6 and E7, are consistently expressed both in cervical cancer cell lines (11) and in HPV-associated cancers (12). Knockdown of E6 and E7 expression in cervical cancer cells triggers cell death, indicating addiction to their expression. These properties make E6 and

E7 ideal tumor-associated antigens to target for therapeutic vaccination. The HPV genome frequently integrates into the host genome during the cervical carcinogenesis. This integration frequently results in the loss of particular regions of viral DNA and is correlated with the upregulation of E6 and E7 expression. E6 and E7 expression overcomes the regulation of cell cycle and apoptotic signals, to drive proliferation. E6 causes the degradation of p53, PDZ proteins and activates *c-myc* and *hTERT* expression whereas E7 inactivates Rb, events that synergize to drive malignant transformation (13,14). Late genes encoding the capsid proteins (L1 and L2) and some early genes are commonly lost, notably E2 which de-represses E6 and E7 expression. Therefore, in contrast to E6 and E7, vaccination against the capsid proteins L1 and L2 is unlikely to confer therapeutic benefit (14).

Formalin-inactivated BPV has been successfully used in cattle to protect herds from bovine papillomas, whereas denatured BPV was not, and formalin-inactivated COPV is also protective for dogs against oral papillomas (15–17). The inability to grow large quantities of HPV in culture and its oncogenic genome precluded the development of an inactivated virus as a preventive vaccine for human use (18). However, the success of the inactivated virus vaccines in animals suggested that the capsid conferred protection if its structure was maintained.

In the early 1990's several groups demonstrated that empty capsids, termed virus-like particles (VLPs), were formed when L1 was over-expressed in insect cells, mammalian cells and yeast (19). Significantly vaccination with L1 VLP, but not denatured L1, induced high serum titers of type-specific neutralizing antibodies, similar to those induced by vaccination with native virions (20). Studies in several animal models demonstrated that vaccination with L1 VLPs was highly protective, albeit only for the type used to derive the vaccine (21–24).

Despite the successful induction of immunity in animal models by L1 VLP vaccination, it was not clear that this could be translated into patients because they do not mimic sexual transmission of HPV. Early phase trials demonstrated that as in the animal models, vaccination with L1 VLP induced high titers of neutralizing antibody, even without the use of adjuvant, with no apparent significant side effects (25,26). The first evidence that L1 VLP vaccines are highly protective against sexual transmission of an oncogenic HPV was a landmark efficacy trial of HPV16 L1 VLPs (27), in which 100% (95% confidence interval, 90–100; $P < 0.0001$) protection against acquisition of persistent HPV16 infection was reported (Table 1). Furthermore, the women vaccinated with placebo acquired HPV16+ CIN, while those vaccinated with HPV16 VLPs did not. However, both groups acquired equal numbers of CIN containing genotypes other than HPV16 (Table 1). This suggests that HPV L1 VLP vaccines provide protection against infection by the homologous papillomavirus type and is consistent with the type-restricted *in vitro* neutralization by sera of vaccinated animals and patients. These early findings have been extended in several Phase III efficacy trials (6,28–31). The data indicate near complete protection for more than five years against the acquisition of infection by the HPV types used to develop the vaccine. Furthermore, there is evidence of partial protection for the types most closely related to those used to generate the L1 VLP vaccine, e.g. partial protection against HPV31 is observed in those vaccinated with HPV16 L1 VLPs, and protection against HPV45 is also observed in those vaccinated with HPV18 L1 VLPs. Interesting evidence of cross-neutralization is also observed for these types *in vitro*, although at titers 10–100-fold lower than observed for homologous type neutralization (32–35).

Such type-restricted (not type-specific) immunity complicates comprehensive vaccination against all oncogenic HPV types. One approach is to increase the number of L1 VLP types

utilized, but highly multivalent vaccines are likely to be more expensive than the currently licensed vaccines (32,33,36). Current formulations of both licensed L1 VLP vaccines, called CERVARIX[®] (GSK) and GARDASIL[®] (MERCK & Co), contain two oncogenic HPV genotypes, HPV16 and HPV18 which together account for ~70% of cervical cancers (28,30). Merck includes HPV6 and HPV11 L1 VLPs in their licensed vaccine GARDASIL[®], to protect against benign genital warts (30) and provide additional rationale for vaccination of men. Efforts to produce octa- or nona-valent L1 VLP vaccines are underway to extend protection to more HPV types and thus potential protect against at least 90% of all cervical cancer cases based upon the current contribution of each oncogenic HPV type (4,7).

WHY DO WE NEED SECOND GENERATION HPV VACCINES?

The current HPV vaccines targeting HPV16 and HPV18 have the potential to produce a dramatic drop in cervical cancer rates if they are widely used (4). Nevertheless, the licensed vaccines are currently too costly for global implementation and thus reach those populations that would benefit most from such a vaccine (37). It is important to note that over 80% of cervical cancer cases occur in Developing nations (38). This reflects the lack of cytologic screening programs and intervention in these countries. Furthermore, as a significant fraction of cervical cancer (~30%) is caused by oncogenic HPV types not included in current HPV vaccine formulations, the type-restricted protection induced by the L1 VLP vaccines means that cervical cancer screening must remain in place. In these settings, therefore, the cost of HPV vaccination must be borne in addition to that of screening. In this situation cytologic screening will become much less cost effective, although cost savings might be achieved by altering the interval and/or modality (i.e. using HPV testing as a first line approach) by which vaccinated women are screened (39). Thus it would be highly desirable to develop a more affordable vaccine for the Developing world as well as to broaden protection against oncogenic HPV infections to a degree that would permit a major reduction in screening in the industrialized world. While increasing the valency of the current L1 VLP vaccines represents a logical approach to overcome the latter problem (51), it seems likely that such a second generation vaccine would, for many years, continue to be too expensive for widespread use in low resource settings. Furthermore, the development of an alternative inexpensive approach to HPV vaccination could drive down the cost of all HPV vaccines and increase their global penetration.

WHY L2?

Papillomaviruses have one other capsid protein beside L1, which is called L2. L2 is a minor component of the viral capsid (Figure 1), but it is necessary for papillomavirus infection, possibly by binding to a secondary viral receptor to facilitate exit from the endosomes and delivery of the viral genome to appropriate domains within the nucleus (40–43). Studies in animal papillomavirus models have indicated that L2, like L1, is a promising antigen for preventive vaccination. These pioneering studies examined immunization of cows or rabbits with the L2 polypeptides produced in bacteria for protection from experimental papillomavirus challenge at both mucosal and cutaneous sites respectively (Table 2). Several studies from Campo's group showed that vaccination with full length and polypeptides of BPV4 L2 were protective against experimental viral challenge (Figure 2) (44–47). They demonstrated that vaccination with a polypeptide comprising residues 11-200 of BPV4 L2, but not a C-terminal peptide thereof, protects cattle against papillomas resulting from experimental challenge with BPV4 virions (Figure 2). Near complete protection against warts after BPV4 challenge was obtained by vaccination with either BPV4 L2 11-200 and full length BPV4 L2, similar to that obtained after vaccination with BPV4 L1 VLP (23,46,48). While the correct conformation of L1 VLPs is critical for the induction of

neutralizing antibodies and protection after vaccination, immunization with L2 purified under denaturing conditions both induced neutralizing antibodies and protected. The L2-vaccinated cattle were challenged shortly after completion of the vaccination regimen (Figure 1), and additional studies to address the long term efficacy of L2 vaccination suggested that protection lasts at least one year (49). However, important caveats remain; the titers of neutralizing antibodies induced by vaccination with L2 are much lower than those produced by L1 VLP vaccination (50–53), and that the cattle vaccinated with BPV4 L2 in these studies were not tested for cross-protection against warts produced by a different papillomavirus type. Furthermore, it should be noted that vaccination with BPV2 L2 did not effectively protect against BPV2 challenge, but rather induced rapid clearance of papillomas that was associated with massive T cell infiltration (54). Interestingly, vaccination with BPV2 L2 did not induce detectable neutralizing antibodies in these animals (54), which contrasts the findings for BPV1 and BPV4 (50,51,55), and may explain the discrepancy in protective efficacy in these studies. It is possible that the data reflect a potent T cell response to L2 vaccination that triggered rejection, but we favor the notion that a weak neutralizing antibody response triggered by BPV2 L2 vaccination reduced the size of the inoculum and thereby facilitated spontaneous clearance of the BPV2 papillomas in this study (54).

Vaccination with L2 polypeptide has also conferred immunity to challenge in the rabbit model (56), although the C-terminus was also protective in this model (52). Vaccination with full length L2 or its polypeptides provided near complete protection of rabbits from cutaneous challenge with the cognate virus type, CRPV. The N-terminal region of L2 that confers protection exhibits considerable amino acid sequence conservation among the medically significant HPV genotypes, suggesting the potential for cross-reactive immune responses to vaccination with this polypeptide (Figure 3). Indeed, animals immunized with HPV L2 produce antibodies that cross-neutralize a broad range of HPV genotypes (53,57–60) and cross-neutralizing L2-specific monoclonal antibodies have been generated (59,61,62), and mapped (Figure 4). This contrasts the type-restricted neutralizing antibodies and protection induced by L1 VLP vaccination. This important distinction suggests the possibility of a simple pan-HPV prophylactic vaccine derived from L2 sequences.

Much of the early work on L2 vaccine development utilized animal papillomaviruses as models because of an inability to readily propagate HPV in tissue culture, the lack of *in vitro* HPV neutralization assays, and the strict tropism of papillomaviruses, i.e. HPV did not produce disease in animals (18). However, the value of this approach has been borne out during the development of L1 VLP vaccines which utilized the same animal papillomavirus models, notably BPV1 or BPV4 infection of cattle, CRPV or ROPV infection of rabbits and COPV infection of dogs. Unfortunately animal papillomavirus models of infection are of limited value in testing L2 vaccines for their ability to induce immunity to a broad range of HPV types as they do not involve genital challenge and the L2 sequences of animal papillomaviruses are evolutionarily distant from the genital HPV types. Many of the technical limitations in production of HPV and testing for *in vitro* neutralization have been overcome by the development of HPV pseudovirion technologies (32,63,64). These HPV pseudovirions can also be used in challenge models in which mice are infected intravaginally or subcutaneously using pseudovirions carrying markers such as RFP or luciferase (62,65–67). These HPV pseudovirion challenge models are consistent with the animal papillomavirus models in indicating that L2 vaccines do induce more broadly cross-protective immune responses than observed for L1 VLP (66).

One possible approach to extend protection to more genotypes might be to vaccinate with L1/L2 VLP rather than L1 only VLPs, since L2 co-assembles with L1 into VLPs when co-expressed. Unfortunately, L2 is immunologically subdominant to L1 in the context of an L1/L2 VLP or even virion vaccine as well as in serologic studies of natural infection (53), and

consequently L1-specific antibodies result almost exclusively. Thus, no more cross-protection is seen in animals vaccinated with L1/L2 VLPs as compared to L1 only VLPs because they generate only L1-specific neutralizing antibodies (Figure 5) (21). This probably reflects the close packed, regular array of L1 (68,69), whereas L2 present at a least a five-fold lower level, and more widely spaced in the capsid (Figure 1A) (70). In addition there is evidence that the principle neutralizing epitopes on L2 may be partially buried and only revealed during infection (71). It is likely that the virus evolved such that L2 is poorly immunogenic given its critical and conserved role in infection, otherwise papillomavirus types would have remained a single serotype.

These studies, when taken together, suggest that L2 contains protective epitopes mostly near its N-terminus, and that not all regions of L2 confer immunity to infection. Immunization with some of these protective L2 polypeptides can trigger the production of broadly neutralizing antibodies which likely mediate protection against both the papillomavirus type from which the L2 vaccine was derived, as well as challenge with divergent papillomavirus types (72). However, the L2 polypeptide vaccine must be presented to the immune system in the appropriate context to trigger immunity because it is subdominant to L1 within the capsid (53). These principles have driven studies to better define the neutralizing epitopes within L2, to understand how L2-specific antibodies prevent infection, and efforts to optimize the approach for L2 vaccine delivery through multimerization, epitope display and/or adjuvant technologies.

Mechanism of vaccine-induced protection from new HPV infection

Vaccines have had a profound impact on infectious diseases, yet much remains to be learned about how they work. T cells and antibodies are typically central to protection, and immunological memory is necessary, but there is much controversy over the details. The relative importance of cell-mediated and antibody responses in preventing viral infection is hotly debated. Opinions range from the notion that antibodies are required primarily to control bacterial, rather than viral, infection and are dispensable for the control of some viral infections, to the polar opposite view that antibodies are the only identified agent of successful vaccine protection (73,74). Study of the unique biology of papillomavirus and the remarkable efficacy of HPV vaccines, as detailed below, can instruct this debate and further the development of second generation HPV vaccines, and approaches to vaccination against other pathogens.

Passive transfer of L1 VLP or L2-specific neutralizing antibodies is sufficient to confer protection to naïve animals suggesting that cell-mediated immunity to capsid proteins is not required for protection (21,22,62). However, these studies do not exclude a contribution of cellular immunity in protecting from natural infection, and indeed there is evidence of L1 and L2-specific cytotoxic T cell responses do occur in response to vaccination with capsid antigen (54,75–77). There have been some reports that vaccination with either L1 VLP or L2 can enhance clearance of pre-existing disease (54,75,78), indicating a possible contribution of capsid protein-specific cytotoxic T cells in controlling infection. However, these effects have not been demonstrated in any randomized and controlled clinical trial to date (79–82). In addition, vaccination with L2 peptides protects rabbits against challenge with CRPV virions but not CRPV viral DNA. Furthermore, papillomas generated by inoculation with wild type CRPV DNA and CRPV DNA defective for L2 expression grow at the same rate in L2 vaccinated animals that are immune to challenge with CRPV virions (72,83).

Neither the L1 nor the L2 capsid proteins are detectably expressed by the basal epithelial cells that harbor the viral infection, but only by terminally differentiated epithelial cells about to slough off (84). Probably as a consequence of this unusual biology, protective

immunity upon vaccination with papillomavirus capsid protein is likely mediated primarily by neutralizing antibodies. This might occur by sterilizing neutralization at the site of infection via antibody in secretions and/or exudates at the site of microtrauma that is typically associated with transmission, e.g. during intercourse. Alternatively, as indicated for immunity to Hepatitis B after vaccination, the rapid induction of immune memory may play a key role in limiting the very early spread and establishment of disease, particularly when neutralizing titers have waned (85,86). Patients vaccinated with GARDASIL[®] remain protected against HPV18 despite loss of detectable HPV18-specific serum antibody in 40% of patients, as measured by a competitive Luminex-based immune assay (86). However, it is unclear if this reflects the ability to rapidly induce immune memory, or the insensitivity of the antibody assay (85). For example, the competitive Luminex-based immune assay used to detect the HPV18 antibody response might fail to detect all neutralizing antibodies because it is based on blockade of binding of a single monoclonal antibody to the capsid, or the cutpoint may be set too high. Further studies using sensitive *in vitro* neutralization assays are needed to address this issue.

Serum neutralizing antibody titer is the most robust immune correlate of protection, both for papillomavirus and numerous other human infectious diseases (25,64,73,87). Importantly, the degree of cross-protection that occurs in patients receiving the HPV vaccines inversely correlates with the cross-neutralization data from *in vitro* assays (33,88). Taken together, the current findings suggest that neutralizing antibodies, not cytotoxic T cell responses, are the principle effectors of protection of patients from acquisition of new HPV infection after vaccination, and this is consistent with the failure of infected basal epithelial cells to express capsid antigens and therefore to be recognized by capsid antigen-specific cytotoxic T cells.

Mechanisms of viral neutralization

The complex events of HPV infection have recently been reviewed elsewhere (89). How papillomavirus infects and how antibodies block the infectious process have been the subject of much debate (71,74). Leading hypotheses are that viruses are aggregated and/or neutralized by preventing their interaction with the cell upon the binding of capsid to one or many antibody molecules; or alternatively that antibody binding prevents necessary conformational changes in the capsid or induces inappropriate conformational states; or that the infectious process is halted by antibody after virion entry to infected cells by, for example, blocking its release of nucleic acid or re-routing the virions along a non-infectious pathway (55,90). The simple occupancy model proposes that neutralization occurs when a significant proportion of available epitopes on the virion are bound by antibody, thereby preventing the attachment of virus to host cells or interfering with the process of entry (74). Indeed in three dimensional reconstruction of the L1-specific neutralizing monoclonal antibody (Mab)9 bound under saturating conditions to BPV1 reveals that the antibody completely coats the surface of the virion (Figure 1B) (91). Consistent with this occupancy model, Mab9 prevents binding of BPV1 to cell monolayers (55). By contrast, the L1-specific antibody 5B6 neutralizes but does not block binding to cell monolayers. Structural analysis reveals that the 5B6 antibody does not cover the pentavalent capsomers on the 5-fold axes, suggesting that they might be sufficient to mediate surface binding (91). Importantly, no evidence of structural changes in L1 was apparent in virions coated with antibody in these moderate resolution reconstructions (Figure 1B). However, there is clear evidence of different states of capsid assembly including closed/mature and open/immature forms that have been associated with differing degrees of intra- and inter- L1 subunit disulphide bond formation (92,93). Structural analysis reveals that the 5B6 antibody bridges the capsomers (91), and between 14 to 72 antibodies per virion were necessary for neutralization, suggesting that 5B6 antibody might block virus uncoating or conformational changes in the capsid (55,94).

The HPV16 L1-specific neutralizing Mabs H16.V5 and H16.E70 prevent endocytosis of virions but not the ability of the virions to bind to the surface of HaCaT cells (71,95). Interestingly they do block binding of HPV16 particles to erythrocytes and to extracellular matrix (33,95,96), suggesting two distinct modes of binding to cell surfaces. These two antibodies also prevent a conformational change in the virion which allows during cleavage of approximately 13 amino acids from the amino terminus of L2 by furin (71). This cleavage of L2 by furin occurs on the cell surface and is critical to infection (97). Neutralizing antibodies to L2, including Mab RG-1 that recognizes residues 17-36 of HPV16 L2, do not prevent the virus from binding to the cell surface (71,94,98). Furin cleavage enhances the exposure of L2 on the virion surface, notably for the residue 17-36 epitope bound by RG-1 (71). Residues within the RG-1 epitope are critical for viral infection (41). Since L2 plays a necessary role for intracellular trafficking in infection and several groups have identified numerous non-neutralizing L2 Mabs of the same isotype that bind to the virus surface (62,99), these findings suggest L2 antibody-dependent neutralization occurs by preventing conformational changes and/or interaction between L2 and key cellular proteins that mediate the complex events of uptake and intracellular papillomavirus trafficking.

Neutralization and protective epitope identification in L2

Many studies have attempted to define the relevant epitopes using vaccination with synthetic peptides or by production of neutralizing monoclonal antibodies. Vaccination with the N-terminus of L2, notably residues 11-200 or 1-88, is protective in multiple animal models (45,46,50,72), and much effort to identify neutralizing epitopes has focus on these regions. Vaccination of rabbits with BPV1 L2 1-88 or 45-173 induces cross-neutralizing antibodies (55,57). Vaccination with L2 residues 94-122 provides relatively type specific protection in rabbits (83,100), and studies in cattle suggest the value of L2 residues 101-120, 131-151 and 151-170 for protection against BPV4 (47). Other studies support residues 108-120 as a cross-neutralizing epitope (58-61). We recently defined L2 residues 17-36 as an important protective epitope (62,66), and others have defined neutralizing epitopes between residues 36-49 (101) and 69-81 (59). Studies examining the remainder of L2 for neutralizing epitopes have generally been negative, although one study demonstrated protection of rabbits after vaccination with the C-terminus (52). There is also evidence for critical functions of L2 during infection within this region (102). Denatured L2 was utilized to elicit neutralizing antibodies in all of the above studies, and this clearly contrasts the conformational dependence of the L1-specific neutralizing antibodies (20). These findings are summarized in Figure 4.

PROOF OF CONCEPT

Synthetic Peptide-based vaccines

Immunization with short synthetic L2 peptides coupled to Keyhole Limpet Hemocyanin (KLH) and administered in adjuvant to animal models protects them from experimental papillomavirus infection at both mucosal and cutaneous sites (83). Embers *et al* used two overlapping peptides within L2 residues 94-122 coupled to KLH and administered in Freund's adjuvant to protect rabbits from experimental challenge with CRPV and ROPV. Campo's laboratory observed the best protection from BPV4 challenge after vaccination of cattle with three immunodominant L2 peptides comprising residues 101-120, 131-151 and 151-170 together in alum, although the 131-151 peptide was effective alone (47). Kawana *et al* found that intra-nasal vaccination with HPV16 L2 108-120 induced IgG and IgA antibodies that neutralized HPV16 and several other HPV genotypes without the need to couple the peptide to KLH (58). This effect was observed in Balb/c but not C57BL/10 mice, suggesting the importance of recognition of an MHC class II epitope within this peptide

(58). No challenge studies were performed, although this epitope overlaps the protective epitopes described by Embers *et al* for the CRPV and ROPV rabbit challenge models (83).

Passive transfer of monoclonal antibody RG-1, which is specific for L2 residues 17-36 and neutralizes both HPV16 and HPV18, protects naïve mice from cutaneous challenge with HPV16 pseudovirions carrying a luciferase reporter gene (62). This suggests that the 17-36 region is a promising candidate for a preventive vaccine. Although coupling of this peptide to KLH to provide T-help, and using an adjuvant to enhance the immune response was effective at inducing cross-neutralizing antibodies (62), Alphas *et al* sought to assemble a totally synthetic and self-adjuvanting vaccine candidate containing the target epitope (HPV16 L2 17-36) together with the TLR2 ligand Pam₂Cys (P2C) and a promiscuous T helper epitope (P25) in a simple branched configuration (66). This approach has demonstrated promise as a vaccine strategy even for poorly immunogenic self-epitopes vaccine(103). Direct chemical synthesis of vaccines has potential advantages over current biological systems for the preparation of antigen, in aspects relating to scale-up of manufacture, purification, reproducibility, stability and regulatory requirements (104). The lipopeptide is able to act as its own adjuvant for the neutralizing epitope, facilitating the generation of an antibody response by triggering dendritic cell maturation via the TLR2-MyD88 pathway and enhancing the class II MHC epitope presentation to and activation of CD4 T cells (66).

HPV L1 VLP vaccines are injected *i.m.* and efforts at needle-free intra-nasal immunization have resulted in variable immune responses and require larger doses (105). Furthermore, licensed HPV L1 VLP vaccines utilize an adjuvant (either alum or alum+ monophosphoryl lipid A), although this may not be required (25). In contrast, *s.c.* and *i.n.* administration of the self-adjuvanting P25-P2C-HPV vaccine each generate similarly potent and consistent serum neutralizing antibody responses in mice suggesting that this approach could be tested in patients. Activation of the antibody response is dependent upon CD40 signaling, consistent with the requirement for class II MHC signaling and a CD4 T cell-dependent humoral response. Vaccination of mice with the lipopeptide construct derived from HPV16 L2 provided effective protection against HPV16 and cross-protection against HPV45, and protected against cutaneous or intra-vaginal challenge. Thus this lipopeptide approach has advantages of direct chemical synthesis, stability and potential to be delivered without needles and generate cross-protection which would facilitate delivery in resource limited settings. The complication for clinical translation of this approach is the need for a class II epitope in the lipopeptide that is broadly recognized in the diverse human population.

Recombinant polypeptide-based vaccines

Immunization with N-terminal L2 polypeptides comprising residues 11-200 protects cattle and rabbits from experimental viral challenge at mucosal or cutaneous sites (46,50,72). Antisera raised to these L2 polypeptides neutralize not only the same type but also the diverse range of heterologous papillomavirus types (50,57,72). However, neutralizing antibody titers against the cognate type are higher than against heterologous types, and there is no clear relationship between titers and evolutionary distance of the heterologous types from the type used to derive the L2 vaccinogen (57). Vaccination of rabbits with L2 1-88 or 11-200 protected rabbits from challenge with even high divergent papillomaviruses, although protection against the homologous type tended to be better. Remarkably, vaccination with HPV16 L2 11-200 provided effective protection from CRPV challenge (72).

As an alternative to vaccination with a mixture of L2 polypeptides derived from multiple types, which would greatly increase the cost and complexity of vaccine development, Jagu *et al* examined concatenated fusion proteins, consisting of several homologous L2 peptides

derived from different medically-significant HPV genotypes, as candidates for a broadly protective HPV vaccine (106). For example, L2 polypeptides corresponding to HPV L2 17-36, 11-88, and 11-200 were chosen for fusion constructs. Three multi-type constructs were tested: one comprising 3 copies of 11-200 (termed 11-200x3, with L2 peptides from HPV6, 16 and 18), one with 5 copies of 11-88 (termed 11-88x5, from HPV1, 5, 6, 16, and 18), and one with 22 copies of 17-36 (termed 17-36x22), with each L2 peptide being derived from different HPV types (107). The proteins were expressed in bacteria and purified under denaturing conditions, and used to immunize rabbits and mice.

The L2 11-88x5 construct generated higher HPV16 neutralizing antibody titers than the HPV16 L2 1-88 peptide, but 11-200x3 did not induce a greater response than HPV16 L2 11-200. The 17-36x22 was even less effective, although strong responses were seen in rabbits, consistent with weak murine T helper epitopes within this peptide (66). The cross-neutralizing antibody responses observed in mice vaccinated with HPV16 L2 polypeptides were consistently weaker than those generated in rabbits receiving the same vaccines, possibly reflecting their differing mechanisms to generate antibody diversity. However, when comparing the multi-type L2 versus the mono-type HPV16 L2 1-88 and 11-200 constructs, immunization with 11-200x3 and 11-88x5 was far more effective in inducing cross-neutralizing antibodies against HPV45, and HPV58, although the fusion proteins do not contain sequences from either of these HPV types, in addition to inducing neutralizing antibodies against HPV6 and HPV18, which are represented in the fusion proteins.

Mice were challenged with HPV16 pseudovirions at 4 months after vaccination with the L2 11-200x3 in a variety of different adjuvants, including alum, immunostimulatory sequence (ISS) 1018, and the modified saponin GPI-0100. Vaccination with 11-200x3 in any of the adjuvants tested induced significant protection. In particular, immunization with 11-200x3 formulated in alum+ISS1018 was more effective than 11-200x3 alone. All of the GPI-0100 adjuvant formulations tested with 11-200x3 were more effective than 11-200x3 alone or 11-200x3 formulated in alum. These findings suggest the need for a potent adjuvant to maintain long-term protection after vaccination with multi-type L2, although this may confer an increased risk for unwanted reactogenicity.

Capsid display vaccines

The VLP vaccines induce very high titer neutralizing antibody that persists but the responses to denatured L2 antigen are much weaker (108). This may partially reflect the ability of L1, but not L2-specific antibody to bind bivalently to virions providing a very high avidity, although very low avidity antibodies are protective (109). Consequently, linear L2 protein-based vaccines may require the use of a potent adjuvant, increasing the possibility of unwanted reactogenicity. Alternatively, it might be possible to confer the immunogenicity of a VLP upon L2-based epitopes by capsid display (69,110). The immunogenicity of L1 VLPs is mediated by the close-packed array of the neutralizing epitopes on the virion surface that potentiates B cell activation (68). L1 VLPs are also able to directly activate dendritic cells and this may also contribute to their immunogenicity (111,112). Since L1 VLP are clearly potent immunogens, several groups have replaced the surface loops comprising the L1 immunodominant neutralizing epitopes with L2 epitopes, including HPV16 L2 65-81 and 108-120 (113). However, the L1 structure is very sensitive to alteration, and this also disrupts some of the major L1 immunodominant epitopes. Varsani et al were able to insert HPV16 L2 108-120 at several sites, and vaccination with these chimeras produced detectable L2-specific antibody, although in vitro neutralization titers were not measured (113). Insertion of either HPV16 L2 65-81 and 108-120 into BPV1 L1 VLP between residues 133 and 134 produced chimeric particles (114). These chimeras were immunogenic and triggered low, but measurable L2-specific neutralizing antibody titers. Thus it is

challenging to display a linear L2 neutralizing epitope by replacing or inserting into a conformation L1 neutralizing epitope and render it immunodominant (114).

Others have examined different viral particles whose structure is well characterized for display of L2 epitopes. Palmer *et al* utilized recombinant tobacco mosaic virus (TMV) to display the L2 94-122 epitope of ROPV and CRPV by insertion between residues 155 and 156 of the U1 coat protein (100). Vaccination of rabbits with these constructs in RIBI adjuvant triggered low titers of CRPV neutralizing antibodies and afforded protection against experimental CRPV challenge (100). These studies demonstrate the potential of this approach, but it is not clear that the optimal selection and placement of the L2 protective epitope has been achieved.

CLINICAL TRIAL DATA

Several L2-based vaccines have been tested in patients. The promising immunogenicity of the HPV16 L2 108-120 peptide vaccine upon intranasal administration in pre-clinical studies led to a placebo-controlled phase I trial (58). Kawana *et al.* tested intranasal inoculation with placebo (n=3), 0.1 mg (n=5) or 0.5 mg (n=5) of HPV16 L2 108-120 peptide at weeks 0, 4, and 12 (60). The immunization caused no serious local or systemic reactions, but the serum antibody response was very weak. Four of the five patients vaccinated with 0.5 mg of peptide generated detectable HPV16 and HPV52 neutralizing antibody titers, whereas this did not occur in the low dose or placebo cohorts (60). It is likely that the immune response to such L2 protective epitopes after intra-nasal inoculation would be greatly improved by coupling to KLH or using the synthetic lipopeptide approach with an appropriately broadly reactive HLA class II epitope (66,115,116).

Two vaccines comprising L2 fused to early viral oncogenes and expressed in bacteria have been tested in patients; TA-CIN is a single fusion protein comprising full length HPV16 L2, E7 and E6, and TA-GW is a single fusion protein comprising HPV6 L2 and E7. Each has potential as a candidate preventive and therapeutic HPV vaccine as they contain capsid antigen and the early viral oncogenes respectively. Both TA-CIN and TA-GW were produced by expression in bacteria and purified under denaturing conditions. A placebo-controlled, double-blinded phase I dose escalation study provided preliminary evidence that three immunizations with 533 µg of TA-CIN without an adjuvant is well-tolerated and immunogenic in healthy volunteers (117). However, this vaccination regimen produced only low titers of L2-specific cross-neutralizing antibodies and weak E6/E7-specific interferon γ and proliferative T cell responses and required high doses of antigen (117,118). In phase II studies in 29 predominantly HPV16+ high grade vaginal or vulval intraepithelial neoplasia (VAIN or VIN) patients, vaccination with TA-CIN followed by a boost with recombinant vaccinia virus expressing E6 and E7 of both HPV16 and HPV18 (TA-HPV) failed to induce an improved rate of lesion regression as compared to earlier studies utilizing TA-HPV alone in 30 VIN patients (82,119). Switching the order of vaccination did not provide improved responses (81,82). The weak antibody and T cell responses to vaccination with TA-CIN protein alone strongly suggested the need to formulate TA-CIN with a potent adjuvant. Studies in primates and mice suggest that the saponin-based adjuvant GPI-0100 greatly enhances the responses to TA-CIN (67). In mice this combination of TA-CIN in GPI-0100 protects against experimental skin infection with HPV16 pseudovirions and challenge with HPV16 E6/E7-transformed tumor challenge (67).

TA-GW was administered at 0, 3, 30 or 300 µg in 1.2 mg of alum to healthy males in a phase I, randomized placebo controlled dose escalation trial (120) and the vaccination was well tolerated. Robust antibody responses were consistently triggered with only the 30 and 300 µg doses. Antigen-specific T cell proliferation and the production of IL5 and interferon

gamma were also observed (120). This led to a Phase IIa open-label study in which 27 genital wart patients received 3 immunizations with TA-GW in alum over 4 weeks (121). The TA-GW vaccine was again well-tolerated. All of the patients generated specific serum IgG antibody responses, predominantly IgG1, against L2E7 after vaccination, and 19/25 made L2E7-specific T cell proliferative responses, producing both interferon gamma and IL5. Five of the 27 subjects completely cleared warts within 8 weeks, and the remainder were offered conventional therapy (121). Recurrence of warts was not seen in any of the 13 persons whose warts cleared by vaccine alone or with conventional therapy. These findings led to a large randomized controlled efficacy trial in patients with genital warts. In this trial the L2-E7 antigen was formulated in the AS02A adjuvant which contains the TLR4 agonist monophosphoryl lipid A, the saponin-based adjuvant QS21, and an oil-in-water emulsion (80). AS02A has been shown to induce high antibody titers, a strong CD4+ cell response, and cytotoxic T lymphocyte activity in humans and has an acceptable safety profile. Two placebo-controlled substudies were also performed to evaluate the value of the TA-GW in AS02A adjuvant when administered in addition to standard therapy. Unfortunately, there was no evidence for benefit of vaccination upon time to recurrence (80). There have been no studies addressing the value of vaccination with TA-GW for prevention of acquisition of HPV6 or HPV11 infection or genital warts to date.

Studies to date generally suggest that vaccination with L1 VLPs or L2 does not influence pre-existing infection (23,72,79,80,83), although there are exceptions (54,78,122). The lack of an obvious therapeutic effect likely reflects the absence of detectable capsid gene expression in HPV-infected basal cells. However, these cells do express E6 and E7 and thus E6 and E7-specific T cell responses could potentially eliminate HPV infection and disease. However, vaccination of women with persistent HPV16+ high grade VIN/VAIN with TA-CIN (without adjuvant) in prime boost combinations with recombinant vaccinia virus expressing HPV16/18 E6 and E7 (TA-HPV) failed to enhance the rate of disease clearance (81,82,119). Similarly, vaccination with HPV6 L2-E7 in ASO2 adjuvant had no impact on pre-existing genital warts (80). These outcomes may reflect induction of an ineffective type of immune response, a failure of the observed E6/E7-specific T cell responses to traffic to the site of the lesion, or their suppression within the lesion microenvironment, or some combination of these issues (118,123).

Vaccination using TA-CIN without adjuvant has not been demonstrated to impact the clinical course of pre-existing lesions in randomized trials. However, it is possible that the induction of E6/E7-specific T cells responses prior to HPV infection could prevent the onset of disease from new infections or possibly the reactivation of previously subclinical disease. In the prophylactic setting, issues of tolerance and immune suppression within the lesion microenvironment might be reduced or negated. Indeed, preventive vaccination with TA-CIN+GPI-0100 completely protected mice from TC-1 tumor challenge, whereas vaccination with TA-CIN alone was poorly effective (67). Furthermore, the GPI-0100 adjuvant dramatically boosted the E7-specific IFN- γ secreting CD8+ T cell response in mice as compared to vaccination with TA-CIN alone (67). Vaccination of macaques induced systemic E6 and E7-specific CD4+ and CD8+ T cell responses. Thus it is possible that any protective efficacy against HPV16-related disease by TA-CIN vaccination via the induction of L2-specific neutralizing antibody could potentially be augmented by HPV16 E6/E7-specific T cell responses.

DISCUSSION

Here we have attempted to lay out the case that second generation preventive HPV vaccines are needed. The principle arguments are that the licensed HPV vaccines are too expensive for global implementation and that they are not targeted against all medically relevant HPV

genotypes. Cost is particularly significant since over 80% of cervical cancer cases occur in Developing countries that currently also lack comprehensive cytologic screening and secondary prevention programs (38). In Developed nations, expensive national screening programs have impacted the incidence of cervical cancer. Unfortunately, despite these efforts significant numbers of cases still occur, particularly in poor and marginalized populations who may not access the costly licensed vaccines. In addition, because the licensed vaccines only target a subset of oncogenic HPV genotypes, it is likely that the current screening programs will remain in place. Thus the cost of vaccination will be borne in addition to the costs of the screening programs. Further, the reduction in disease burden by implementation of the licensed HPV vaccines will dramatically reduce the predictive value and cost effectiveness of cytologic screening.

How might these issues be addressed? It is unlikely that the licensed HPV vaccines can be produced sustainably at the costs required for global implementation without a switch to local production, and this has intellectual property issues. Local production by companies in Developing countries can achieve dramatic cost savings following the model of numerous childhood immunizations, e.g. the robust biotechnology sectors in India and China. Furthermore, new vaccines would increase the competition and help drive down costs. The production of multiple L1 VLP types in insect cells or even yeast, and formulation of all of these components is complex. It is likely that the production in bacteria of a single polypeptide, as with an L2 fusion protein, could be substantially less expensive to manufacture than a polyvalent L1 VLP vaccine prepared in eukaryotic cells. Indeed, licensed vaccines contain two or four L1 VLP types and eight or nine type vaccines are being tested. Thus if an L2 vaccine were proven effective in people for broad protection against oncogenic and other HPV types, its simpler manufacturing process could make local production of such a vaccine highly feasible. This might achieve the goal of producing an HPV vaccine at sustainable prices in emerging countries and lead to more global implementation of cervical cancer prevention. Broad and long lasting protection against all oncogenic types could also eliminate the need for cytologic screening programs in Developed nations, and L2 vaccines might also prevent the morbidity and costs associated with other non-malignant HPV disease. Unlike genital HPV infection, non-genital HPV infection is typically not transmitted sexually and frequently occurs in children. If future clinical testing of an L2-based vaccine demonstrated protection against infection at non-genital cutaneous sites, in addition to protection against genital HPV infection, it might provide a rationale for administration with other childhood vaccines. This could remove both the logistical challenges of vaccinating adolescents and the theoretical issue that vaccination of adolescents against HPV might influence their sexual behavior.

Acknowledgments

The authors gratefully acknowledge the support of grants to RBSR from the National Institutes of Health (National Cancer Institute, SPORE in Cervical Cancer, P50 CA098252 and RO1 CA118790), WKH (RO1 CA133749) and to SJ with a fellowship from the Prevent Cancer Foundation, Alexandria, VA.

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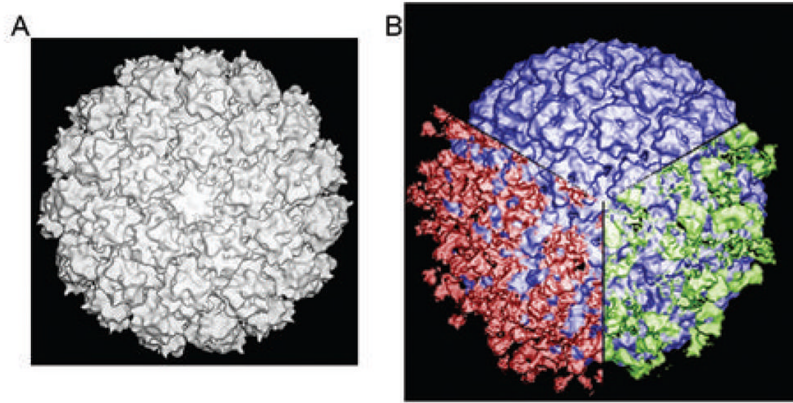


Figure 1. Image reconstruction of bovine papillomavirus type 1 alone or bound to neutralizing L1-specific monoclonal antibody

A. An 18 Å resolution image reconstruction of a BPV1 virion purified from a bovine papilloma and viewed down the 5-fold axis of symmetry. The structure was obtained using image reconstruction of cryoelectron micrographs (70). Note that the capsid has T=7 icosahedral symmetry. It comprises 72 capsomers, each containing 5 L1 molecules and up to one L2 molecule (124). 60 capsomers are coordinated with 6 adjacent capsomers (hexavalent) and 12 with five neighboring capsomers (pentavalent). A pentavalent capsomer can be seen in the center of this image. The button of density present in the center of the pentavalent capsomers likely corresponds to the minor capsid protein L2 (124), a small portion of which is exposed on the capsid surface and available for binding by neutralizing antibodies. Image reconstructions of HPV1, HPV16 and CRPV suggest that all papillomaviruses have a very similar structure (92,124–126). The viral genome forms a nucleohistone core with no detectible icosahedral symmetry. **B.** An image reconstruction of a BPV1 virion (blue) bound to either neutralizing monoclonal antibody MAb#9 (red) or 5B6 (green) (91) and viewed along the 3-fold axis of symmetry. MAb#9 covers the surface of the virion and binds to all of the capsomers. 5B6 covers the hexavalent capsomers, but not the pentavalent capsomers along the 5-fold axes of symmetry. Both images were generated by Benes Trus (CIT, NIH).

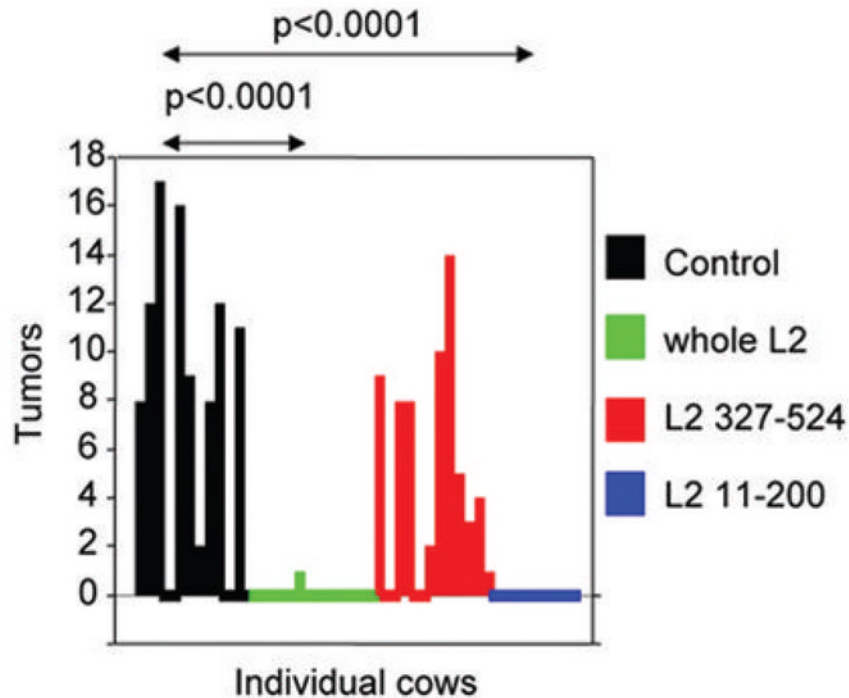


Figure 2. Vaccination of calves with BPV4 L2 or polypeptide comprising residues 11-200 of BPV4 L2 protects against experimental challenge at a mucosal site with BPV4

The data is taken from (46) and redrawn. Recombinant polypeptides comprising the full length BPV4 L2 or residues 11-200 or 327-524 were expressed in *E. coli*, prepared as inclusion bodies and formulated in alum. Each animal (12/group) was inoculated twice at monthly intervals with either 50 μg of polypeptides encompassing full length L2, or 330 μg of L2 residues 11-200 or 330 μg of residues 327-524 and the control group was not vaccinated. The cattle generated antibody responses to each antigen. BPV4 challenge was performed in the upper alimentary tract two weeks after completion of vaccination. Only the control group and the group vaccinated with the C-terminal L2 peptide developed tumors (papillomas).

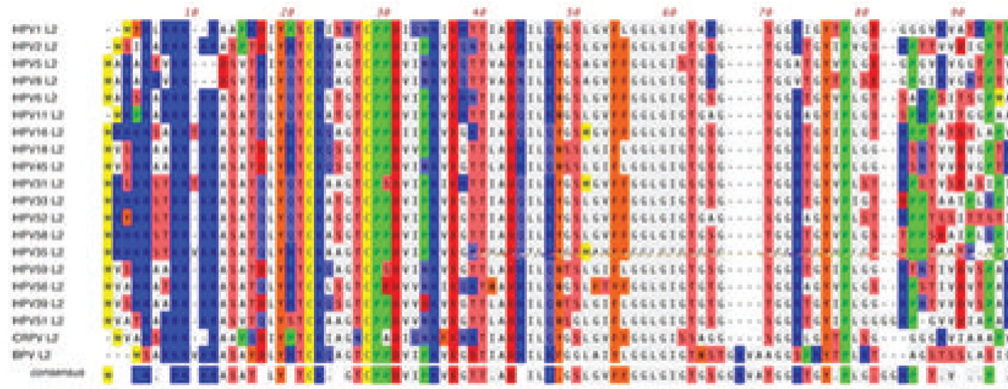


Figure 3.
Amino acid sequence conservation at the N-terminus of L2.

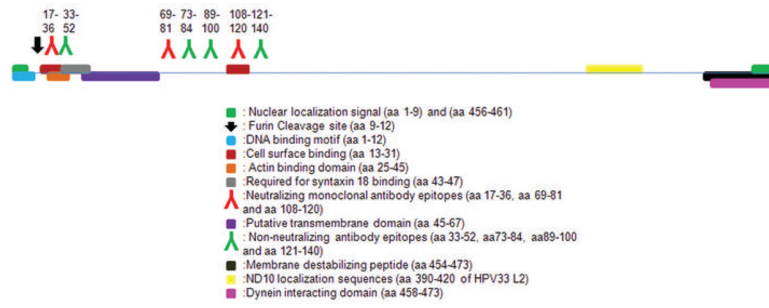


Figure 4.
 Diagram of the known functions and monoclonal antibodies binding sites in HPV16 L2.

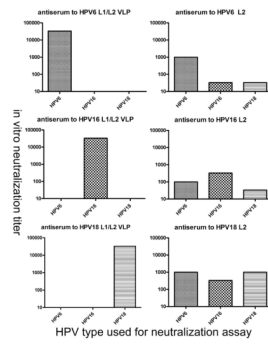


Figure 5. Vaccination with L1/L2 VLPs induces type specific neutralizing antibodies, vaccination with L2 protein alone generates a lower titer, but broadly neutralizing antibody response Rabbits were immunized with L1/L2 VLPs or L2 protein alone derived from HPV6, HPV16 or HPV18. Sera were tested for neutralizing titer against HPV6, HPV16 or HPV18 pseudovirions (53). These HPV types represent prototypic members of three distinct evolutionary species (107).

Table 1

Summary of the data from a landmark efficacy trial of an HPV16 L1 VLP vaccine (27).

	Vaccine	Placebo
Women enrolled	768	765
Persistent HPV16 infections	0	41 *
Incident CINI HPV16+	0	5
Incident CINII HPV16+	0	4
Incident CIN attributed to a genotype other than HPV16	22	22

* In this prophylactic vaccination trial to assess the safety and efficacy of HPV16 L1 VLPs, women without evidence of HPV16 infection were vaccinated three times with HPV16 L1 VLPs or placebo and followed for new persistent HPV16 infection or development of CIN over a median of 17.4 months. Vaccination provided 100% (95% confidence interval, 90–100; $P < 0.0001$) protection against acquisition of persistent HPV16 infection, although 41 incident persistent HPV16 infections and HPV16 DNA+ 9 Cervical Intraepithelial Neoplasias (grades I or II) occurred in the placebo recipients. However, the same number of cases of non-HPV16 CIN occurred in women who received placebo and those who received the HPV16 L1 VLP vaccine (27).

Table 2

Summary of published L2 vaccine studies in animal models.

Study (<i>et al</i>), date	Virus - Host	Antigen	Dose-Adjuvant	Vaccination Schedule	Challenge	Neutralization Peak Titer	Protected
Jarret, 1991	BPV2-Cattle	BPV2 L2 full length	2x 500 µg-IFA	0, 2 wks	+2 wk	Below detection	Enhanced clearance ¹
Campo, 1993; Chandrachud, 1995; Gaukroger, 1996; McGarvie, 1994	BPV4- Cattle	BPV4 L2 full length or 11-200	2x 100 µg-Alum 2x 50 µg-None 2x 1 mg-Alum	0, 4 wks 0, 4 wks 0, 4 wks	+2 wk +2 wk +2 wk +1 year	5	Yes Yes Yes Yes
Christensen, 1991	CRPV-Rabbit	CRPV L2	4x gelpure-CFA	0, 4, 8, 12 wks	+2wk	32	Yes
Lin, 1992	CRPV-Rabbit	CRPV L2	4x 250 µg-RIBI	0,3,6,9 wks	+2 wk	10	Yes
Embers, 2002	ROPV, CRPV-Rabbit	CRPV/ROPV L2 peptide	3x 500 µg peptide/ KLH-CFA/IFA	0, 4, 8 wks	+4-12 wks	10	Yes
Gambhira, 2007	ROPV, CRPV-Rabbit	L2 1-88 or 11-200	3x 500 µg -RIBI	0, 4, 8 wks	+4 wks	8000	Yes, Cross-type
Palmer, 2006	ROPV, CRPV-Rabbit	TMV-L2 recombinant particles	3x 200 µg-RIBI	0, 33, 63 days	+1 wk	500	Yes, Cross-type
Karanam, 2009	HPV16-mouse	TA-CIN (full length HPV16 L2)	3x 20 µg-GPI-0100	0, 2, 4 wks	+2 wk	400,000	Yes
Jagu, 2009	HPV16-mouse	Polymeric L2	3x 25 µg-GPI-0100, or alum, or ISS 1018 or adjuvant combinations	0, 2, 4 wks	+4 mo	100,000	Yes
Alphas, 2008	HPV16/45-mouse	HPV16 L2 17-36 lipopeptide	3x 20 nmol, self- adjuvanting	0, 4, 8 wks	+2 wk	10,000	Yes, Cross-type

¹ Cattle were not protected, but warts were rapidly cleared in the vaccine group. This may reflect reduced inoculum due to partial neutralization or clearance, perhaps as a bystander effect. This enhanced clearance was not described in any other L2 vaccine study.

Table 3

Summary of published studies in patients utilizing vaccines that contain L2.

Study (<i>et al</i>), date	Virus - Disease	Dose-Adjuvant	Vaccination Schedule	Trial type
Thompson, 1999	Healthy volunteer	3 × 0.3, 30, 300 µg HPV6 L2-E7 in alum	0, 1, 4 wks 0, 4, 8 wks	Dose escalation (randomized, controlled)
Lacey, 1999	HPV6/11+ Genital warts	3 × 300 µg HPV6 L2-E7 in alum	0, 1, 4 wks	Therapeutic trial (open label)
Vandepapeliere, 2005	HPV6/11+ Genital warts	3 × ? µg HPV6 L2-E7 in ASO2A	0, 2, 4 wks	Therapeutic trial (randomized, controlled)
De Jong, 2002	Healthy volunteer	3 × HPV16 L2-E7/E6 26, 128, 533 µg, no adjuvant	0, 4, 8 wks	Dose escalation
Smyth, 2004	HPV16+ AGIN	3 × 533 µg HPV16 L2-E7/E6 no adjuvant	Vaccinia E6E7 then at 0, 4, 8 wks	Therapeutic trial
Davidson, 2004	HPV16+ AGIN	3 × 533 µg HPV16 L2-E7/E6 no adjuvant	0, 4, 8 wks then Vaccinia E6E7 boost	Therapeutic trial
Fiander, 2006	HPV16+ AGIN	3 × 533 µg HPV16 L2-E7/E6 no adjuvant	0, 4, 8 wks then Vaccinia E6E7 boost	Therapeutic trial
Kawana, 2003	Healthy volunteer	HPV16 L2 108-120 peptide at 500 µg, 100 µg, no adjuvant	0, 4, 12 wks	Dose escalation