

EXPERIMENTAL STUDY

HPV16 L2 improves HPV16 L1 gene delivery as an important approach for vaccine design against cervical cancer

Namvar A¹, Bolhassani A¹, Hashemi M²Department of Hepatitis and AIDS, Pasteur Institute of Iran, Tehran, Iran. azam.bolhassani@yahoo.com**ABSTRACT**

BACKGROUND: High-risk human papillomavirus (HPV) infections have been associated with the development of cervical cancer. HPV16 is the most dominant high-risk types of HPV worldwide. L1 and L2 are the major and minor capsid proteins of HPV, respectively. Both proteins are able to self-assemble as a virus-like particle (VLP).

METHODS: In the current study, the human embryonic kidney cells were transfected with the plasmid DNA encoding HPV 16 L1 or L1–L2 genes and their expression was compared using different transfection reagents.

RESULTS: Our data showed that the recombinant L1–L2 DNAs were expressed in a high efficiency compared to L1 DNAs as detected by western blotting, fluorescent microscopy, and flow cytometry. In addition, Lipofectamine and Turbofect as the transfection reagents conferred more potent delivery than PEI 25 kDa indicating high toxicity of this system on HEK-293 cells. These results suggest the use of the full length of L2 as an efficient agent for overcoming the cell barriers and poor uptake of DNA *in vitro* and *in vivo*.

CONCLUSION: The high expression of HPV16 L1–L2 in HEK-293 cells using different delivery systems opens the way for new studies concerning to the use of L2 for DNA delivery via covalent linkage with the gene of interest (Fig. 5, Ref. 20). Text in PDF www.elis.sk.

KEY WORDS: HPV, cervical cancer, L1, L2, lipid and polymeric carriers.

Introduction

Human papillomaviruses (HPV) have been known as an original agent in the progress of cervical cancer (1, 2). Among the 15 oncogenic genital HPV types, HPV16 is the most common (~50%), followed by HPV18 (~20%) and HPV45 (~10%) of cervical cancer cases worldwide (3). Thus, the generation of a preventive or therapeutic HPV vaccine would be mostly helpful in these cases (1). Recently, the virus-like particles (VLPs) have been considered as the best candidate for vaccine development against HPV infections. HPV capsids are composed of 72 pentamers of the L1 major coat protein, and an unknown number of the L2 minor coat protein (4). VLPs can be produced by an expression and assembly of the L1 protein alone or its co-expression with the L2 protein (1). The major challenge for the use of L2 protein in VLP-based vaccine alone is its poor immunogenicity as compared to HPV L1 VLP. Regarding to the conserved epitopes of L2 protein, the efforts were done to enhance the L2 immunogenicity by the linkage of L2 to TLR agonists or the use of a concatenated N-terminal fragment of L2 (5). Furthermore, the addition of L2 to L1 VLPs increases the

number of neutralizing antibodies as well as better yields for the L1/L2 particles than for the L1 VLPs (1, 6). Indeed, the chimeric L1–L2 capsids may have higher stability than the L1 capsids (6). The studies showed that HPV L1 alone or accompanied by L2 expressed in cultured cells could be self-assembled in the nucleus (7, 8). The recent evidence has shown that VLPs harboring both the L1 and L2 capsid proteins may be more potent for DNA delivery than VLPs composed of L1 alone (9, 10). Despite current researches on using chimeric HPV VLPs as a vaccine or DNA/peptide delivery system, there are not any full reports comparing the quality and quantity of *in vitro* L1 protein expression alone, with L1–L2 fusion protein using lipid and polymeric delivery systems. Herein, the role of L2 in increasing L1 gene delivery and subsequently its *in vitro* expression was evaluated by fluorescent microscopy, flow cytometry, and western blot analysis. Our data indicated that L2 augmented the expression of L1 protein likely by facilitating its delivery. In addition, we evaluated the efficiency of three delivery systems such as lipofectamine, Turbofect, and polyethylenimine (PEI) for L1 and L1–L2 plasmid DNA delivery, *in vitro*.

Materials and methods*Preparation of pcDNA-L1–L2*

For the generation of eukaryotic expression vector harboring the L1 (pcDNA-L1), the L1 DNA, was amplified by PCR from pUF-L1 (kindly provided by Prof. Martin Muller, German Cancer Research Center) using primers designed to generate *Xba*I and *Not*I restriction sites at the 5' and 3' ends of the amplified fragments,

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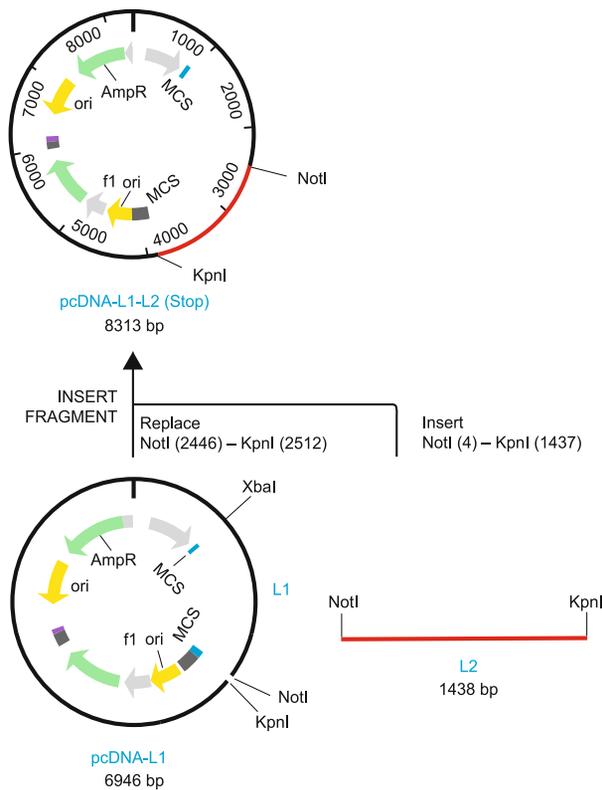


Fig. 1. Cloning of L2 into pcDNA-L1 vector.

48 h after transfecting HEK-293 cells. GFP fluorescence was observed in cells that received 2 µg of pEGFP-N1, pEGFP-L1 and pEGFP-L1-L2 vectors. The transfection efficiency using different delivery systems showed that the Turbofect and Lipofectamine were more potent than PEI 25 kDa (NrE=10). In addition, the level of GFP expression detected by pEGFP-L1-L2 delivery was significantly higher than that by pEGFP-L1 transfection ($p < 0.05$). The levels of protein expression were 89.14 %, 25.17 %, 84.91 % for pEGFP-L1-L2; 65.05 %, 10.73 %, 45.35 % for pEGFP-L1; 86.53 %, 33.76 %, 82.43 % for pEGFP-N1, using Lipofectamine, PEI, and TurboFect, respectively. The transfection efficiency of L1 and L1-L2 genes using three methods has been shown at 48 h after cell transfection by flow cytometry and fluorescent microscopy in Figs 3 and 4, respectively.

Detection of L1 and L1-L2 protein expression using western blot analysis

Western blot analysis was performed using anti-L1 monoclonal antibody to ensure the proper expression of L1 and L1-L2. The specific bands with expected size of 82 kDa, 106 kDa, and 133 kDa were detected for L1-GFP, L1-L2, and L1-L2-GFP expressed from pEGFP-L1, pcDNA-L1-L2, and pEGFP-L1-L2 vectors in the transfected cells as shown in Figure 5. Indeed, L1 or L1-L2 expression was detectable in transfected cells as compared to untransfected cell extracts by western blotting.

Discussion

The prevention of cervical cancer would need to induce immune responses against at least 7 high-risk HPV types (e.g., 16, 18, 31, 33, 45, 52, and 58), which increases the cost and complexity of prophylactic vaccines (5). The recent L1 protein-based HPV vaccines stimulate the neutralizing antibodies against infections and offer type-restricted protection (5). The HPV L2 protein is a good candidate for development of prophylactic vaccine, because L2-specific antibodies have cross-neutralizing activity against various HPV types (11, 12). L2 polypeptide vaccines could provide a broad range of protection; however, the L2 protein vaccines are poorly immunogenic compared to L1 vaccines. Thus, the chimeric VLPs showed the potential to use as a vaccine candidate for a broad spectrum of high-risk HPVs (13). Different studies demonstrated that L2 (especially the amino terminal of L2) has a potential as a protective antigen, although it does not make VLP (14, 15). Indeed, L2 confers more stability to the VLP and is also necessary for HPV infections (8).

In this study, the human embryonic kidney cells (HEK-293) were transfected with the DNA constructs expressing HPV16 L1, and HPV16 L1-L2 proteins. Our major goal was the evaluation of L1-L2 expression compared to L1, *in vitro*. Furthermore, we compared the efficiency of different cationic polymers and lipids to deliver plasmid DNAs. Recombinant L1-L2 DNAs were expressed in HEK-293 cells in a high potency, detected by fluorescent microscopy, flow cytometry, and western blotting. High GFP fluorescence was observed in the cells that received pEGFP-L1-L2

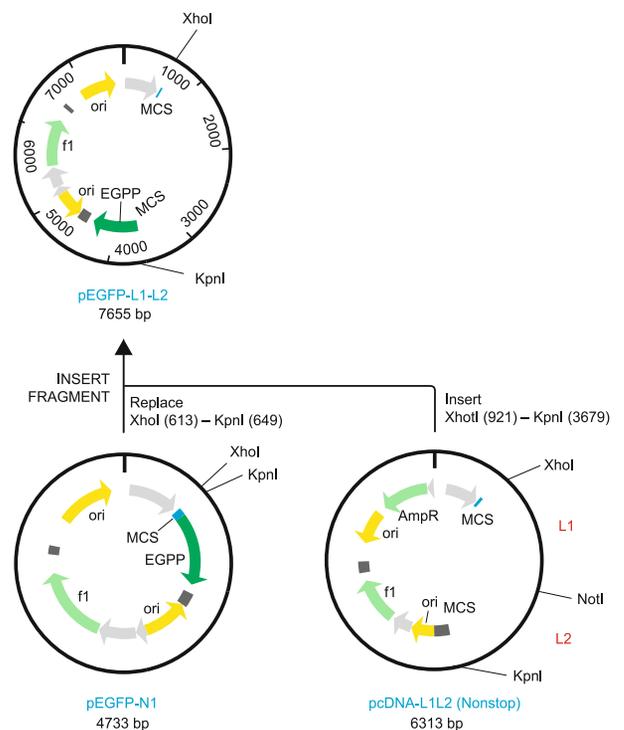


Fig. 2. Cloning of L1-L2 into pEGFP-N1 vector.

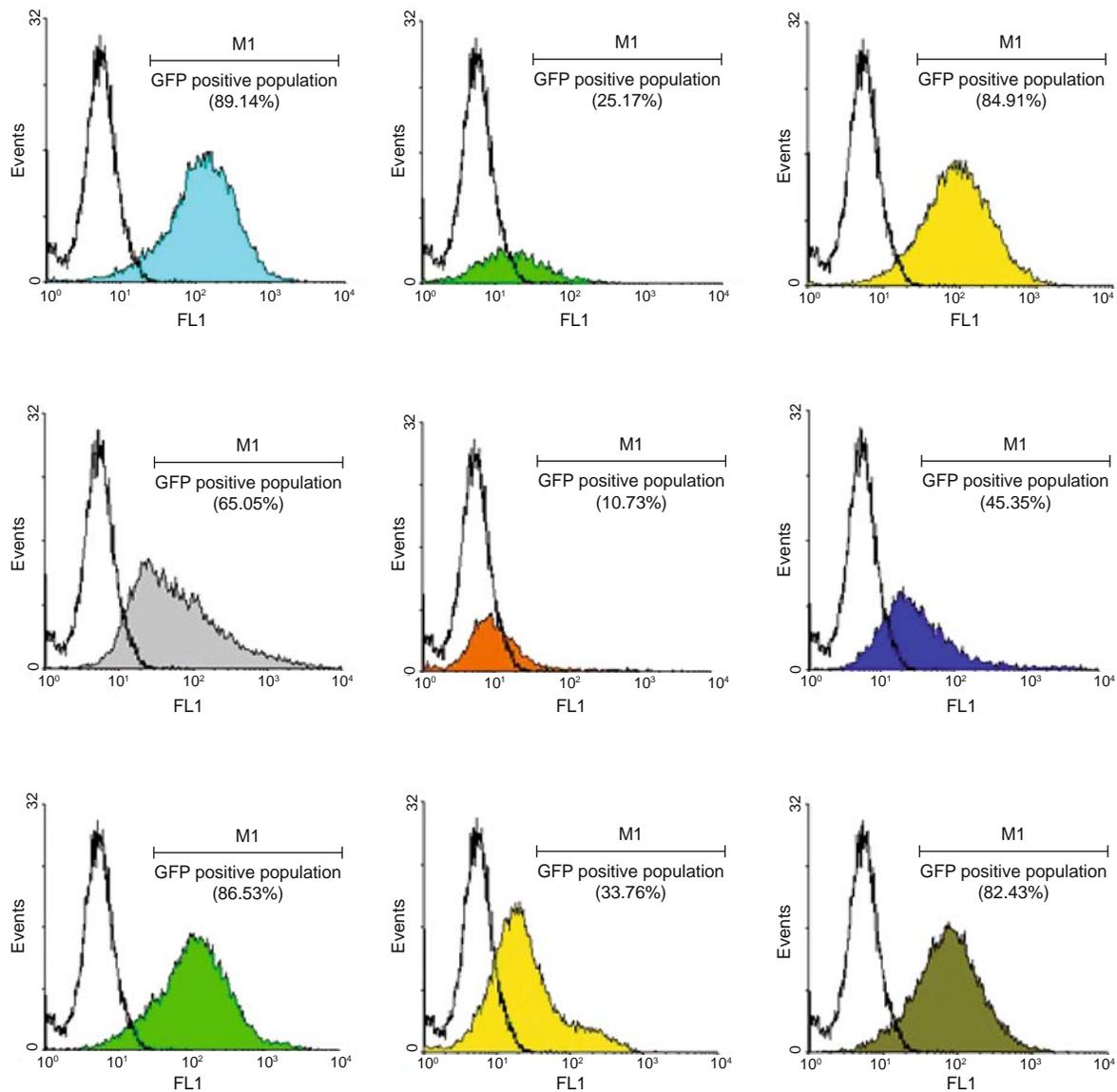


Fig. 3. The transfection efficiency of L1 and L1-L2 genes using flow cytometry analysis: Delivery of pEGFP-L1-L2 (A-C), pEGFP-L1 (D-F), and pEGFP-N1 (G-I) by Lipofectamine, PEI, and Turbofect, respectively. High GFP expression level was observed for pEGFP-L1-L2 delivered by Lipofectamine (89.14%) and then TurboFect (84.91%) as compared to pEGFP-L1 delivered by these systems (65.05%, 45.35%, respectively)

vector as compared to pEGFP-L1 vector. In addition, the transfection efficiency observed by Lipofectamine and then Turbofect was higher than that by PEI. Flow cytometry analysis indicated a clear and significant quantitative separation between L1-GFP or L1-L2-GFP expressing transfected cells and untransfected cells (negative control). L1-L2-GFP expression was detectable in 84–89% of the transfected cells as compared to L1-GFP expression (~45–65%) using Lipofectamine and TurboFect transfection systems. Using PEI reagent, higher expression of L1-L2-GFP protein (~25%) was observed in comparison with L1-GFP protein (~10%). However, PEI 25 kDa showed more toxicity on HEK-293 than two other delivery systems suggesting a lower percentage of GFP

fluorescent. Furthermore, L1 or L1-L2 expression was also detectable in the transfected cell extracts compared to untransfected cells by western blotting. The dominant bands of ~82 kDa, 106 kDa, and 133 kDa were detected in transfected cells expressing L1-GFP, L1-L2, and L1-L2-GFP using Anti-L1 antibody. No such corresponding band was revealed in the untransfected cells. These results confirmed the expression of L1-L2 protein as fused to GFP or not. Regarding the obtained data, it seems to consider the L2 gene as facilitating agent of DNA delivery followed by its protein expression.

Other studies also showed that L2 is necessary for intracellular encapsidation of papillomavirus genomes. During the primary

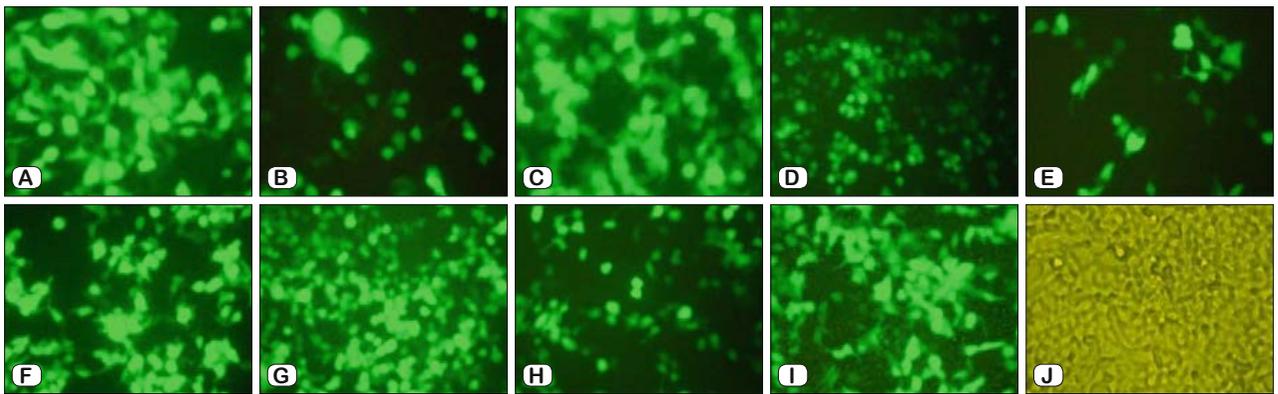


Fig. 4. The transfection efficiency of L1 and L1-L2 genes using fluorescent microscopy: Delivery of pEGFP-L1-L2 (A-C), pEGFP-L1 (D-F), and pEGFP-N1 (G-I) by Lipofectamine, PEI, and Turbofect, respectively against untransfected cells (negative control, J).

infection, the L2 protein localizes in sub-nuclear domains known as nuclear domain 10 (ND10). The targeting of L2 to ND10 may facilitate the delivery of the viral genome to ND10 for initiating viral transcription (11, 16–18). L2 has also been shown to mediate co-localization of L1 and DNA within the nucleus in promyelocytic leukemia oncogenic domains (POD) (19, 20). Moreover, it was recently indicated that VLPs containing both the L1 and L2 capsid proteins might be more efficient for DNA delivery than VLPs consisting of L1 alone (9). The studies showed that DNA co-delivered with L1 VLPs was retained within endosomes, and that efficient endosomal escape was dependent on a 23 amino acid sequence located within the C-terminal region of L2 (10). Generally, L2 may facilitate expression of co-delivered DNA not only by mediating endosomal escape, but also by mediating localization

of DNA to transcription sites (10). All these experiments utilized L2 VLP associated with L1 VLP (L1/L2 VLP) for DNA delivery or VLP-based vaccination. In the current study, the full length of L2 gene fused to the full length of L1 gene without linker could increase the level of L1 expression as compared to the L1 DNA. This is the first investigation on the effects of L2 DNA as linked to L1 gene as well as the comparison of three transfection reagent. The similar results were obtained by Lipofectamine, Turbofect, and PEI for high expression of L1-L2 DNA.

Conclusion

In summary, our findings support the use of L2 as an efficient DNA delivery system to overcome cell barriers and enhance protein expression for development of HPV vaccines.

References

1. Wakabayashi MT, Da Silva DM, Potkul RK, Kasta WM. Comparison of human papillomavirus type 16 L1 chimeric virus-like particles versus L1/L2 chimeric virus-like particles in tumor prevention. *Intervirology* 2002; 45: 300–307.
2. Ki Rnbauer R, Taub J, Greenstone H, Roden R, Durst M, Gissmann L, Lowy DR, Schiller JT. Efficient self-assembly of human papillomavirus type 16 L1 and L1-L2 into virus-like particles. *J Virol* 1993; 67: 6929–6936.
3. Alphas HH, Gambhira R, Karanam B, Roberts JN, Jagu S, Schiller JT, Zeng W, Jackson DC, Roden RBS. Protection against heterologous human papillomavirus challenge by a synthetic lipopeptide vaccine containing a broadly cross-neutralizing epitope of L2. *PNAS* 2008; 105: 5850–5855.
4. Conway MJ, Cruz L, Alam S, Christensen ND, Meyers C. Cross-neutralization potential of native human papillomavirus N-terminal L2 epitopes. *PLoS ONE* 2011; 6: e16405.
5. Nieto K, Weghofer M, Sehr P, Ritter M, Sedlmeier S, Karanam B, Seitz H, Muller M, Kellner M, Horer M, Michaelis U, Roden RBS, Gissmann L, Kleinschmidt JA. Development of AAVLP (HPV16/31L2) particles as broadly protective HPV vaccine candidate. *PLoS ONE* 2012; 7: e39741.

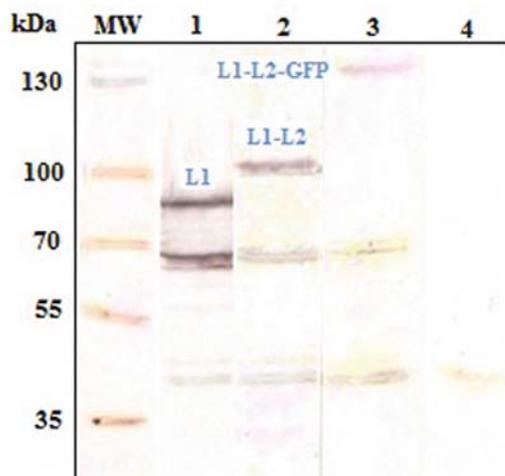


Fig. 5. Western blot analysis using an anti-L1 monoclonal antibody showed the expression of the full-length L1-GFP (line 1, 82 kDa), L1-L2 (line 2, 106 kDa), and L1-L2-GFP (line 3, 133 kDa) proteins, respectively. Any detectable band was not observed in untransfected cells as a negative control (line 4)

6. **Hernandez BY, Ton T, Shvetsov YB, Goodman MT, Zhu X.** Human papillomavirus (HPV) L1 and L1–L2 virus-like particle-based multiplex assays for measurement of HPV virion antibodies. *Clinical and Vaccine Immunology* 2012; 19: 1348–1352.
7. **Nakao S, Mori S, Kondo K, Matsumoto K, Yoshikawa H, Kanda T.** Monoclonal antibodies recognizing cross-neutralization epitopes in human papillomavirus 16 minor capsid protein L2. *Virology* 2012; 434: 110–117.
8. **Cianciarullo AM, Szulcowski V, Alencar Muniz A, Boschi Bazan S, Araújo Aires K, Müller M, Armbruster-Moraes E.** Production of HPV16 L1L2 VLPs in cultures of human epithelial cells. *FORMATEX* 2010; 1073–1082.
9. **Kamper N, Day PM, Nowak T, Selinka HC, Florin L, Bolscher J.** A membrane-destabilizing peptide in capsid protein L2 is required for egress of papillomavirus genomes from endosomes. *J Virol* 2006; 80: 759–768.
10. **Malboeuf CM, Simon DAL, Lee YEE, Lankes HA, Dewhurst S, Frelinger JG, Rose RC.** Human papillomavirus-like particles mediate functional delivery of plasmid DNA to antigen presenting cells *in vivo*. *Vaccine* 2007; 25: 3270–3276.
11. **Yoon SW, Lee TY, Kim SJ, Lee IH, Sung MH, Park JS, Poo H.** Oral administration of HPV-16 L2 displayed on *Lactobacillus casei* induces systematic and mucosal cross-neutralizing effects in Balb/c mice. *Vaccine* 2012; 30: 3286–3294.
12. **Gambhira R, Karanam B, Jagu S, Roberts JN, Buck CB, Bossis I, Alphas H, Culp T, Christensen ND, Roden RBS.** A protective and broadly cross-neutralizing epitope of human papillomavirus L2. *J Virol* 2007; 81: 13927–13931.
13. **Kondo K, Ochi H, Matsumoto T, Yoshikawa H, Kanda T.** Modification of human papillomavirus-like particle vaccine by insertion of the cross-reactive L2-epitopes. *J Med Virol* 2008; 80: 841–846.
14. **Jagu S, Kwak K, Karanam B, Huh WK, Damotharan V, Chivukula SV, Roden RBS.** Optimization of multimeric human papillomavirus L2 vaccines. *PLoS ONE* 2013; 8: e55538.
15. **Rubio I, Seitz H, Canali E, Sehr P, Bolchi A, Tommasino M, Ottonello S, Müller M.** The N-terminal region of the human papillomavirus L2 protein contains overlapping binding sites for neutralizing, cross-neutralizing and non-neutralizing antibodies. *Virology* 2011; 409: 348–359.
16. **Bowsera BS, Chena HS, Conway MJ, Christensen ND, Meyersa C.** Human papillomavirus type 18 chimeras containing the L2/L1 capsid genes from evolutionarily diverse papillomavirus types generate infectious virus. *Virus Res* 2011; 160: 246–255.
17. **Becker KA, Florin L, Sapp C, Sapp M.** Dissection of human papillomavirus type 33 L2 domains involved in nuclear domains (ND) 10 homing and reorganization. *Virology* 2003; 314: 161–167.
18. **Kieback E, Muller M.** Factors influencing subcellular localization of the human papillomavirus L2 minor structural protein. *Virology* 2006; 345: 199–208.
19. **Day PM, Baker CC, Lowy DR, Schiller JT.** Establishment of papillomavirus infection is enhanced by promyelocytic leukemia protein (PML) expression. *Proc Natl Acad Sci USA* 2011; 101: 14252–14257.
20. **Kiesslich A, von Mikecz A, Hemmerich P.** Cell cycle-dependent association of PML bodies with sites of active transcription in nuclei of mammalian cells. *J Struct Biol* 2002; 140: 167–179.

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