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# **Cervical Cancer: Development of Targeted Therapies Beyond Molecular Pathogenesis**

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# **Abstract**

It is well known that human papillomavirus (HPV) is the causative agent of cervical cancer. The integration of HPV genes into the host genome causes the upregulation of E6 and E7 oncogenes. E6 and E7 proteins inactivate and degrade tumor suppressors p53 and retinoblastoma, respectively, leading to malignant progression. HPV E6 and E7 antigens are ideal targets for the development of therapies for cervical cancer and precursor lesions because they are constitutively expressed in infected cells and malignant tumors but not in normal cells and they are essential for cell immortalization and transformation. Immunotherapies are being developed to target E6/E7 by eliciting antigen-specific immune responses. siRNA technologies target E6/E7 by modulating the expression of the oncoproteins. Proteasome inhibitors and histone deacetylase inhibitors are being developed to indirectly target E6/E7 by interfering with their oncogenic activities. The ultimate goal for HPV-targeted therapies is the progression through clinical trials to commercialization.

## **Keywords**

cervical cancer; targeted therapy; human papillomavirus; vaccine; proteasome inhibitor; histone deacetylase inhibitor; siRNA; gynecological cancer; HPV-targeted therapies

# **1. Introduction**

Cervical cancer is one of the leading causes of cancer death in women worldwide [1,2]. A recent estimate indicates that there are approximately 529,800 cases and 275,100 deaths due to cervical cancer annually [3]. The current standard of care for advanced cervical cancer includes the use of a chemotherapeutic drug, cisplatin, in conjunction with local radiation therapy [4]. In many cases, these treatments are responsible for significant adverse effects [5]. Despite improvements noted with combination therapy, five-year survival in most

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**Conflict of Interest**

Jayne Knoff and Benjamin Yang declare they have no conflict of interest.

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patients affected by advanced cervical cancer is approximately 30% [6]. Furthermore, analyses of conservative surgical treatments for cervical dysplasia also found that conization and LEEP had pregnancy-related morbidities such as pre-term delivery, low birth weight, vaginal hemorrhage, and post-birth infections [5]. Additionally, these treatments are associated with significant recurrence rates of up to 10% of cases. Hence, the efficacy of existing surgical treatments may drop due to the risk of relapse, possibly insufficient clearance of affected tissue and potentially undesirable side effects. Furthermore, although the commercialization of preventive HPV vaccines, Gardasil and Cervarix, are effective in preventing cervical cancer, they do not have therapeutic effects against preexisting precancer or cancer lesions. Thus, there is an urgent need for innovative therapies that can reduce the number of cervical cancer cases as well as improve patients' lives and outcomes.

It is well known that human papillomavirus (HPV) is the causative agent of cervical cancer (for review, see [7,8]). More than 100 types of HPV have been identified [9] and among those, 12 are considered 'definite carcinogens' due to their oncogenic potential [10-12]. These high-risk HPV types are associated with the development of high-grade lesions and malignant tumors. Persistent infection with a high-risk HPV type has been proven to be causative and necessary for the development of squamous intraepithelial lesions (SIL) (also known as cervical intraepithelial neoplasia or CIN), and malignant cervical cancer [13]. HPV-16 and HPV-18 are the most common high-risk HPV types associated with cervical cancer and are responsible for about 62.6% and 15.7% of cervical cancers, respectively [14]. Therefore, HPV-16 and HPV-18 have been the primary focus of targeted therapies for cervical cancer.

HPV is a non-enveloped, circular, double-stranded DNA virus that belongs to the Papillomaviridae family. HPV is comprised of an icosahedral capsid enclosing an approximately 8 kilobase pair genome that encodes early proteins, including the oncoproteins E6 and E7, and late proteins that form the capsid of the virion (for review, see [15]). HPV infects basal cells of the cervical epithelium and other epithelial tissues upon tissue microtrauma [16] (Figure 1). While most HPV infections are self-limiting and transient (for review, see [1]), in a productive infection of HPV, expression of the HPV genome correlates with the maturation of the infected cell. Immature epithelial cells in the basal layer of the epithelium allow expression of the HPV early genes while the late genes are expressed in terminally differentiated cells, allowing encapsidated virions to be released from the superficial epithelial layers. This localization of HPV results in pathological lowgrade squamous intraepithelial lesions (LSIL, also known as CIN 1). In LSIL, the HPV genome is typically in the episomal form. High-grade squamous intraepithelial lesions (HSIL, also known as CIN 2/3) are of greater concern because they can progress to malignancies. In some HSILs, high-risk HPV types may integrate into the host genome. This integration of HPV genes causes the upregulation of E6 and E7 oncogenes because E2, a transcriptional repressor of E6 and E7, is deleted by viral integration into the host genome. These E6 and E7 proteins inactivate and degrade tumor suppressors p53 and retinoblastoma (Rb), respectively, leading to cell cycle deregulation, genomic instability, and uncontrolled proliferation of the host cell (for review, see [13] [17,18]).

HPV E6 and E7 antigens are ideal targets for the development of targeted therapies against cervical cancer and precursor lesion because they 1) are constitutively expressed in infected cells and malignant tumors but not in normal cells [19], 2) are essential for cell immortalization and transformation. Therefore, many cervical cancer targeted therapeutic strategies have mainly focused on HPV E6 and E7 oncogenic proteins (Figure 2). Therapies targeting HPV E6 and E7 are being explored by various approaches in preclinical studies as well as different phases of clinical trials. Immunotherapies are being developed to target E6 and E7 by eliciting antigen-specific immune responses. siRNA technologies also target E6

and E7 by modulating the expression of the oncoproteins. Proteasome inhibitors and histone deacetylase inhibitors are also being developed to indirectly target E6 and E7 by interfering with their oncogenic activities. The ultimate goal for HPV-targeted therapies is the progression through clinical trials to commercialization.

# **2. Targeted therapeutic strategies**

### **2.1 Immunotherapies**

HPV E6 and E7 oncogenic proteins are ideal targets for the development of immunotherapy against HPV-associated lesions. As mentioned above, they are constitutively expressed in infected cells and malignant tumors but not in normal cells. Furthermore, because they are foreign antigens, they do not have problem of central immune tolerance, which is commonly seen when endogenous antigens are used for vaccine development. In addition, because they are essential for cell immortalization and malignant progression, they do not have issues of antigenic loss. Therefore, many cervical cancer immunotherapeutic strategies have mainly focused on eliciting HPV E6 and E7-specific T cell immune responses to control and inhibit the progression of HPV-associated lesions. Therapeutic HPV vaccines have been evaluated in a variety of preclinical models and clinical trials using live vector, peptide, protein, DNA, RNA replicon, and dendritic cell (DC)-based vaccines targeting HPV E6 and E7 oncoproteins (for review, see [20,21]). Table 1 provides a summary of therapeutic HPV vaccines in clinical development.

**2.1.1. Live vector-based Therapeutic HPV vaccines—**The concept of live vector vaccines encompasses the use of bacterial and viral vectors, which replicate within the body and facilitate the spread of the encoded antigens. Live vector-based therapeutic HPV vaccines can deliver E6 and E7 antigens to antigen-presenting cells in order to stimulate antigen presentation through MHC class I to CD8+ cytotoxic T cells and MHC class II to CD4+ T helper cells. However, live vectors inherently pose a safety risk, particularly to immunocompromised individuals. Live vectors may also face limited capacity for repeated administration due to the stimulation of vector-specific neutralizing antibodies and/or preexisting vector-specific immunity.

**Bacterial vector-based vaccines:** Several bacterial vectors have been explored for therapeutic HPV vaccines including *Listeria monocytogenes* [22,23], *Lactobacillus casei* [24], and *Lactococcus lactis* [25]. Among these, *L. monocytogenes* has generated significant interest. *L. monocytogenes* is a gram-positive intracellular bacterium that invades macrophages and evades phagocytosis within the phagosome by using listeriolysin O (LLO), a pore-forming toxin. Upon its escape into the cytoplasm of antigen presenting cells (APCs), antigens are delivered and processed in both MHC class I and class II pathways that activate CD4+ and CD8+ T cells

*L. monocytogenes* bacterial vector-based vaccines have been used in clinical arena. ADXS11-001 is a live, attenuated *L. monocytogenes* bacterial vector secreting HPV-16 E7 fused to LLO. In Phase I trials in patients with advanced cervical cancer, ADXS11-001 was found to be safe and well tolerated with dose limiting toxicities of hypotension and flu-like symptoms [26,27]. Antigen-specific T cell responses and some clinical responses were observed, which will be further assessed in phase II trials. ADXS11-001 is currently being studied in three clinical trials with active enrollment. An open-label, Phase I dose escalation trial will assess the safety of ADXS11-001 in patients with HPV-associated oropharyngeal cancer (NCT01598792)[28]. An ongoing Phase II trial will assess the safety and efficacy of ADXS11-001 in treating women with persistent or recurrent cervical carcinoma (NCT01266460)[29]. A randomized, single-blind, placebo-controlled Phase II study is

testing whether three doses of ADXS11-001 in women with CIN2/3, for whom surgery is indicated, can safely reverse disease (NCT01116245)[30,31].

**Viral vector-based:** The high immunogenicity of viral vectors makes them attractive for use in therapeutic HPV vaccines. There have been several viral vectors used to deliver HPV E6 and E7 antigens, including adenoviruses [32,33], alphaviruses [34-36] and the vaccinia virus [37-39]. The vaccinia virus, an enveloped, double-stranded DNA virus within the Poxviridae family, is a promising viral vector because of its large genome and high infectivity.

Vaccinia vector-based therapeutic HPV vaccines have translated to clinical trials. A recombinant vaccinia virus expressing HPV-16/18 E6/E7 antigens (TA-HPV) has been evaluated in Phase I/II clinical trials in patients with early-stage cervical cancer [40], latestage cervical cancer [41] vulvar intraepithelial neoplasia [42] and vaginal intraepithelial neoplasia [43]. TA-HPV was found to be safe, well tolerated and potent in stimulating vaccinia-specific antibody responses and HPV antigen-specific CTL responses [40,41,44]. A Phase II trial in patients with early stage cervical cancer has been conducted to study the safety and immunological effects of TA-HPV in combination with surgery [40,45]. The results of this trial indicated that TA-HPV is safe, causing only mild to moderate toxicities, and was able to generate HPV-specific CTLs as well as serological responses in some patients [40].

Another vaccinia-based therapeutic HPV vaccine in clinical trials is the Ankara vector, a modified vaccinia vector expressing HPV-16 E6 and E7 antigens and the adjuvant IL-2 (MVA-HPV-IL2, also known as TG4001/R3484). TG4001/R3484 was designed to: (1) alert the immune system specifically to HPV-16-infected cells presenting HPV-16 E6 and E7 antigens and (2) further stimulate the immune system in clearing the infection using IL-2. In Phase II clinical trials, TG4001/R3484 was found to be both safe and effective in producing clinical responses in women with HPV-16-positive CIN 2/3 [46]. TG4001/R3484 was tested in a placebo controlled Phase IIb trial on patients with HPV-related CIN 2/3 lesions. Interim results demonstrated proof of concept for the therapeutic vaccine in HPV-16 monotherapy, but the trial did not reach its primary endpoint of six month resolution in the CIN 2/3 indication and is not progressing to a Phase III trial [47].

Vaccinia vector-based vaccines have also been assessed in clinical trials using heterologous prime boost regimens. In a Phase I clinical trial, 29 female patients with HPV-associated high-grade anogenital intraepithelial neoplasia received three doses of TA-CIN, a fusion protein-based HPV vaccine, followed by one dose of TA-HPV [48] [49]. The vaccine was safe and well tolerated in patients without any significant adverse side effects. Moreover, full and partial clinical responses were seen in 17% of patients while 62% had symptomatic improvement. A separate Phase I trial of a therapeutic HPV DNA vaccine with TA-HPV prime boost in combination with topical imiquimod in CIN3 patients is recruiting patients (NCT00788164)[50].

**2.1.2. Peptide-based Therapeutic HPV vaccines—**Direct administration of peptides derived from HPV antigens can lead to the uptake of peptides by dendritic cells for antigen processing and presentation, thus activating antigen-specific T cell immunity. Peptide-based vaccines are stable, easy to produce, and have a good safety profile. Their creation involves the identification of CTL epitopes and CD4+ T helper epitopes that stimulate potent antigenspecific CD8+ and CD4+ T cell immune responses, respectively. Research has focused on addressing the main limitations of peptide-based vaccines, namely their low immunogenicity and the obstacle of MHC restriction.

Adjuvants can be used to enhance the immunogenicity of peptide-based vaccines and have been explored in clinical studies. Because HLA-A\*0201 is the most common human MHC class I molecule carried by over 50% of the general population, peptide-based therapeutic HPV vaccine studies have focused on HPV-16 E7 CTL epitopes that are presented by HLA-A\*0201. Vaccination with CTL epitope peptide, strong adjuvant, and nonspecific help might prime CTLs against the weakly immunogenic CTL epitope and result in clearance of HPV in preclinical models. The combination of CTL epitope peptide with nonspecific T cell help has been explored in the form of a lipopeptide construct of HPV-16 E7 aa86-93 peptide linked to Pan HLA-DR epitope (PADRE) peptide in women with late stage cervical cancer [51]. Further similar studies have focused on combining two E7-derived epitopes (aa11-20 and aa 86-93) with Montanide ISA 51 as adjuvant in combination with PADRE universal T helper peptide [52] [53]. However, among these trials, only weak immune responses were detected without evidence of antitumor benefits [52] [53]. A Phase I clinical trial found that a vaccine consisting of HPV-16 E7 peptide (aa 12-20) administered along with a construct of Incomplete Freund's adjuvant, HPV E7 lipopeptide (aa 86-93) and PADRE, was able to stimulate an immune response in a significant proportion of eighteen HLA-A2-positive patients with CIN/VIN II/III [54]. The vaccine was well tolerated and led to complete regression of CIN lesions in 3 of 17 evaluable patients. However, as these epitopes are HLA-A2\*0201-restricted, these peptide vaccines would only have clinical benefit for the population of patients who are positive for this MHC class I.

The limitation of MHC restriction associated with peptide-based vaccines necessitates the identification of immunogenic epitopes corresponding to the polymorphic MHC molecules within the population. Long overlapping peptides circumvent MHC restriction by including a range of antigenic epitopes of HPV E6 and E7 proteins and have renewed interest in therapeutic HPV E6/E7 peptide-based vaccines. These larger peptides have been tested in recent clinical trials in end-stage cervical cancer patients. A vaccine comprised of 13 overlapping peptides representing HPV-16 E6 and E7 antigens mixed with Montanide ISA 51 adjuvant was safe and able to elicit broad T cell responses in end-stage cervical cancer patients [55]. A second clinical trial using a broad array of epitopes in early-stage cervical cancer patients generated increased HPV-16-specific CD4+ and CD8+ T cell responses compared to unvaccinated patients [56]. Phase II clinical trials of this vaccine demonstrated great efficacy in HPV-16-positive high-grade vulvar intraepithelial neoplasia (VIN) patients. Half of the patients with histologically confirmed HPV-16-positive VIN3 displayed complete regression of their lesion after 3 or 4 vaccinations with HPV-16 E6/E7 overlapping peptide [57]. Further investigation into vaccine-induced immune responses showed that a high ratio of the number of HPV16-specific effector T cells to the number of HPV16 specific  $CD4+CD25+F\alpha p3+T$  reg cells was predictive of clinical success [58]. Recently, a placebo-controlled randomized Phase II study demonstrated that this HPV-16 E6/E7 synthetic overlapping long-peptide vaccine increased numbers of circulating IFN-γproducing HPV-16 antigen-specific T cells in patients with HPV-16+ HSIL [59]. The vaccination had few side effects but no HPV clearance was observed at the time of lesion excision, nor could conclusions be drawn regarding vaccine-induced T cell lesional infiltration due to patient accrual problems. The focus remains on developing a welltolerated vaccine capable of generating strong immune responses in patients with precancer lesions.

**2.1.3. Protein-based vaccines—**Protein-based vaccines are promising forms of therapeutic HPV vaccines. They are safer than live-vector based vaccines and avoid MHC restriction since they include epitopes that bind to all haplotypes of MHC class I and II molecules. However, protein-based vaccines have some disadvantages, including relatively poor immunogenicity. Additionally, protein-based vaccines are processed through the endocytic pathway and are presented via the MHC class II pathway, generating

In clinical trials, several protein-based therapeutic HPV candidates have been explored. Namely, HspE7, a chimeric protein composed of bacille Calmette-Guerin heat shock protein (Hsp65) and HPV-16 E7, has generated considerable interest. HspE7 was found to be welltolerated as a single-agent therapy in both Phase I and Phase II clinical trials, promoting lesion regression in several HPV-associated diseases, including CIN 2/3 [60-62].

TA-CIN, a fusion protein-based vaccine expressing HPV-16 L2-E6-E7 conjugated proteins, has shown tremendous potential in clinical studies. TA-CIN has been shown to boost E6 and E7-specific CD8+ T cell immune responses in healthy volunteers [63]. TA-CIN has also been tested in conjunction with imiquimod in a Phase II clinical trial in patients with highgrade vulvar intraepithelial neoplasia (VIN). Intramuscular administration of TA-CIN and topical application of imiquimod was well tolerated without adverse effects. Antigenspecific antibody titers were generated; however, titers were not significantly different after imiquimod application or vaccination in "responders" versus "non-responders." "Responders" to the therapy demonstrated high levels of  $CD4^+$  and  $CD8^+$  T cells locally as well as within HPV-associated lesions. Imiquimod was shown to increase T cell infiltration, leading to the complete regression of VIN lesions in 63% of patients one year after treatment. Additionally, 36% of patients with VIN lesions showed complete HPV clearance and 79% of women remained symptom free. Phase III clinical trials will be needed to assess the comparative efficacy of this combinatorial approach. Additional clinical trials have tested TA-CIN with TA-HPV as mentioned in section 2.1.1 [48,49]. TA-CIN has been tested preclinically with GPI-0100, a semi-synthetic saponin adjuvant, and shown to generate significant HPV antigen-specific CD8+ T cell immune responses and therapeutic antitumor effects against HPV-16 E6/E7-expressing tumors [64]. The encouraging results have led to the preparation of several clinical trials (Roden, personal communication).

**2.1.4. DNA Vaccines—**Among the different forms of therapeutic HPV vaccines, DNA vaccines have become an attractive approach due to their stability, simplicity and safety. DNA vaccination involves direct injection of plasmid DNA encoding antigen of interest into host cells, promoting expression and presentation of the encoded antigen by transfected cells and stimulating cell-mediated and/or humoral immune responses against the encoded antigen. DNA vaccines do not pose the intrinsic safety risks associated with introducing a tumor cell, bacteria, or virus into a patient. Naked DNA is easy to manufacture and can sustain antigen expression by target cells for longer durations than RNA vaccines. Additionally, DNA vaccines do not elicit neutralizing antibodies *in vivo* as do live vectorbased vaccines, therefore allowing repeated administration. However, in general DNA vaccines have poor immunogenicity and require additional strategies to increase vaccine potency. Many strategies to enhance DNA vaccine potency focus on targeting DNA vaccines to dendritic cells (DCs), the potent activators of antigen-specific immune responses. In general, these strategies to improve the potency of therapeutic HPV DNA vaccines through DC modification can be classified by: 1) increasing the number of antigenexpressing/antigen-loaded DCs, (2) improving HPV antigen expression, processing and presentation in DCs, and 3) enhancing DC and T cell interaction (for review, see [21]). Encouraging data from preclinical studies have led to testing several therapeutic HPV DNA vaccines in clinical trials. Strategies to increase the number of antigen-expressing/antigenloaded DCs include enhanced vaccine delivery methods such as gene gun, microencapsulation, and electroporation (for review see [65]).

Another delivery technique that has been evaluated in several clinical trials is microencapsulation of DNA vaccines [66-68] (for review, see [69]). Of considerable interest is ZYC101, a plasmid encoding several HPV-16 E7-specific HLA-A2 restricted CTL epitopes encapsulated in 1-2 μm biopolymer microparticles composed of poly-lactide coglycolide (PLG). A Phase 1 trial examined the effects of ZYC101 on patients with highgrade CIN and found that five of 15 patients experienced complete histological regression and 11 showed substantial HPV-specific T cell responses with no serious adverse effects [67]. A newer version of the plasmid, amolimogene bepiplasmid (ZYC101a), encoding HPV-16 and -18 E6 and E7 protein fragments was tested in a Phase II clinical trial of 127 subjects with high-grade CIN. The vaccine was well tolerated and promoted resolution of CIN 2/3 in patients under 25 years of age (70% versus 23% in the placebo group) [68]. Amolimogene was recently explored in a Phase II/III double-blinded, randomized, placebocontrolled clinical trial examining its efficacy and safety in the treatment of patients with CIN 2/3. Eligible subjects were randomized to either drug or placebo groups and during the six-month study period and were monitored by colposcopic, cytologic, and HPV testing. Persistence or resolution of disease was determined by a loop electrosurgical excision procedure (LEEP) performed at study exit [70]. Results showed that 11 of 21 patients receiving amolimogene generated enhanced T cell responses to HLA-A2 restricted HPV 16/18 peptides compared to baseline and 6 of those subjects experienced resolution of CIN 2/3 lesions [70].

Electroporation following intramuscular injection has also been assessed as a potential vaccine delivery method [71]. For example, VGX-3100, a DNA vaccine incorporating plasmids targeting HPV-16/18 E6 and E7 proteins, is delivered via intramuscular injection followed by electroporation using a CELLECTRA constant current device to deliver a small electrical charge. In a Phase I clinical trial, subjects with a history of high-grade CIN were vaccinated with VGX-3100, which was well tolerated. VGX-3100 immunization elicited a T cell response in 14 out of 18 subjects and all subjects had antibody positivity to at least two antigens [72]. Currently, VGX-3100 is being tested in individuals with histologically confirmed HPV-16/18 associated high-grade CIN in a double-blind, randomized, placebocontrolled Phase II clinical trial (NCT01304524) [73].

Several clinical trials have been conducted to examine the effects of vaccines designed to improve MHC class I processing. A Phase I clinical trial showed that 8 of 15 patients with high-grade CIN had increased E7-specific T-cell responses after intramuscular injection with a therapeutic HPV DNA vaccine, pNGVL4a-Sig/E7(detox)/HSP70. The vaccine pNGVL4a-Sig/E7(detox)/HSP70 consists of DNA encoding a signal peptide (Sig), linked to an attenuated form of HPV-16 E7 (E7(detox)) fused to Mycobacterium tuberculosis heat shock protein 70 (HSP70). Although the DNA vaccine was well tolerated, subjects treated with the vaccine did not experience significantly improved therapeutic effects compared to unvaccinated subjects [74]. Despite poor immunogenicity, complete histologic regression did occur in 33% of patients vaccinated with the highest dose of pNGVL4a-Sig/E7(detox)/ HSP70 (3 mg per vaccination) [74]. The same vaccine has also been used in patients with HPV-16+ head and neck cancer (Gillison and Wu, personal communication). It is now clear that approximately 20% of head and neck cancers are associated with HPV, particularly HPV-16 (for review, see [75]). These early phase clinical trials with pNGVLA4a-Sig/ E7(detox)/HSP70 demonstrate great safety without significant side effects. The results of the early phase clinical trials suggest that the DNA, although safe, is not sufficient to generate impressive therapeutic effects on its own. Therefore, prime-boost regimens with different expression vectors or the adoption of different delivery methods, such as electroporation or intradermal administration via gene gun, may be necessary to increase DNA vaccine potency. For example, an ongoing Phase I clinical trial is examining three routes of administration (intradermal administration via gene gun, intramuscular administration, and

intralesional delivery) of a DNA vaccine encoding calreticulin (CRT) linked to HPV-16 E7 protein, pNGVL4a-CRT/E7(detox), to treat CIN 2/3 lesions in HPV-16 positive patients (NCT00988559) [76].

Other methods have also been explored for improving the immunogenicity of therapeutic HPV DNA vaccines. Of particular interest are toll-like receptor (TLR) agonists, which are immunomodulators that generate robust immune responses and increase the potency of therapeutic HPV vaccines. For example, imiquimod, a TLR7 agonist, has been shown to promote the activation of APCs leading to the production of the cytokines IFN-α, IL-6, and TNF-α [77]. Cytokines induce a robust and potent immune response by facilitating adaptive immune cell activation and differentiation. Thus, imiquimod is a promising adjuvant for therapeutic HPV DNA vaccines. An ongoing Phase I clinical trial investigates treatment of CIN3 with a DNA prime-vaccinia boost regimen combined with topical imiquimod. The subjects were intramuscularly primed with pNGVL4a-Sig/E7(detox)/HSP70 DNA vaccine and then boosted with a recombinant vaccinia virus encoding HPV-16 and 18 E6 and E7 (TA-HPV) with local application of imiquimod on the CIN lesion [50].

**2.1.5. Dendritic cell based-vaccines—**Dendritic cells are professional APCs and are thus able to induce the adaptive immune response, T cell-mediated immune responses in particular, by processing antigen and priming T cells in both the MHC class I and class II pathways. The process of pulsing DCs with HPV antigenic peptides, proteins, or DNA encoding antigens *ex vivo* enables loading of MHC class I and class II molecules with HPV epitopes and subsequently allows DCs to differentiate and mature. Upon re-introduction of the DCs into the body, T cells become primed and elicit a cell-mediated immune response. Understanding DC biology, such as differentiation and maturation, as well as antigen processing and presentation has been helpful in providing a rationale for improving DCbased vaccines. For example, siRNAs used to target pro-apoptotic proteins have been used to enhance the potency of therapeutic HPV DC-based vaccines. Specifically, a therapeutic HPV vaccine consisting of E7-loaded DCs transfected with BAK/BAX siRNA elicited potent E7-specific CD8+ T cell immune responses and antitumor effects in TC-1 tumorbearing mice by prolonging the life of DCs [78,79].

A Phase I clinical trial assessed the safety and immunogenicity of a DC-based vaccine using HPV-16 E7 and/or HPV-18 E7 in 10 patients diagnosed with early-stage cervical cancer [80]. All patients developed antibody and CD4+ T cell responses to the HPV E7-loaded DC vaccination and 8 of the 10 patients had increased E7-specific CD8+ T cell counts compared to prevaccination levels. Overall, the vaccine was determined to be safe and immunogenic. Although DC-based vaccines may be used in advanced cases of cervical cancer, it is unlikely that they will be used to treat CIN lesions because the procedures involved in this treatment are relatively labor-intensive and costly.

**2.1.6 Combinatorial Approaches—**Although strategies to enhance different types of therapeutic HPV vaccines have been developed, the combination of strategies may further increase the immunogenicity and efficacy of therapeutic HPV vaccines. Hence, the focus of therapeutic HPV vaccine strategies has shifted in the direction of combinatorial approaches to work toward commercialization. One combinatorial approach uses fusion protein antigens concurrently with low-dose radiation treatment of tumors in preclinical models [81]. Vaccination of mice with DNA vaccine encoding calreticulin (CRT) fused with HPV-16 E7 (CRT/E7) combined with radiation therapy showed an increase in therapeutic efficacy as compared to DNA vaccinated mice alone [81]. The combination of therapies elicited the highest frequency E7-specific CD8<sup>+</sup> T cell response and increased tumor susceptibility to E7-specific CTL activity in the tumor-microenvironment [81]. This response slowed and stabilized tumor growth, which ultimately led to an increase in mouse long-term survival

rates [81]. Additionally, radiation therapy was successful in causing apoptosis of tumor cells, indicating that radiation is a useful method in stabilizing tumor cell growth when applied with immunotherapy. The combination of chemotherapy, radiation therapy, and vaccination suggests effective antitumor effects.

The combination of chemotherapeutic agents with DNA-based vaccines is another emerging strategy that may be an effective HPV therapy as shown in preclinical models [82-86]. For example, the chemotherapeutic agent, apigenin, was used concurrently with a DNA encoding heat shock protein 70 (HSP70) and HPV-16 E7 [87]. Vaccination with E7-HSP70 DNA in conjunction with apigenin chemotherapy demonstrated the highest frequency of effector  $CD8^+$  T cells and memory  $CD8^+$  T cells. Vaccination and chemotherapy caused tumor susceptibility to E7-specific cytotoxic immune responses that led to a reduction in tumor size and an increase in survival rates. Apigenin treatment also proved to increase tumor cell apoptosis in a dose-dependent manner. Overall, the combination of chemotherapy and DNA-based HPV vaccination generated the greatest antitumor effect.

#### **2.2 Proteasome inhibitors**

A major oncogenic activity of HPV E6 is to bind to the E3 ubiquitin ligase E6-AP and redirect its activity to promote the rapid proteasomal degradation of p53 and PDZ family proteins (ex. hDlg, hScribble, and hMAGI) [88-90], thereby abolishing cell cycle regulation. This creates the opportunity to target ubiquitin-dependent protein degradation for the treatment of cervical cancer. Thus, treatment with proteasome inhibitors may restore near normal levels of p53 and in so doing, promote cell death of abnormally growing HPVassociated cervical cancer cells.

Bortezomib (marketed by Millennium Pharmaceuticals as Velcade) is a widely used proteasome inhibitor that has been recognized as a potent chemotherapeutic agent (for review, see [91]). It is currently approved for use in humans to treat relapsed multiple myeloma and mantle cell lymphoma (for review see [92]). Bortezomib-induced tumor cell apoptosis may enhance the immunogenicity of tumor cells and provide an opportunity for generating tumor-specific immunity [93]. The therapeutic effects of bortezomib have been examined in preclinical models of cervical cancer. Bortezomib has been found to dramatically decrease p53 degradation in HeLa cells and to a lesser extent in CaSki cells [94] as well as increase the expression of pRb in both cell lines [95]. Furthermore, bortezomib enhances E7-specific CD8+ T cell-mediated immune responses generated by therapeutic HPV DNA vaccine (CRT/E7(detox)) in mice bearing E7-expressing tumors [84]. Notably, bortezomib has been tested in a clinical trial in combination with another chemotherapeutic drug. A Phase I/II trial is testing bortezomib in combination with vandetanib, a chemotherapeutic drug for used for thyroid cancer treatment, in patients with a variety of cancers, including cervical cancer, for cancer reduction and duration (NCT00923247). The results of this study are pending.

Several different proteasome inhibitors have been explored for their potential as targeted therapeutics against HPV-expressing cervical cancer in cell culture systems. A collection of chalcone-derivatives lacking aminoacidic components (termed RAMBs), ubiquitinproteasome system stressors that inhibit ubiquitin-mediated protein degradation upstream of the 20S proteasomal catalytic activities, were found to selectively kill cervical cancer cells *in vitro* [96]. Furthermore, MG132 is a protein aldehyde that blocks the function of the 26S proteasome complex. MG132 was found to be capable of inhibiting the growth of HeLa cells by inducing cell cycle arrest as well as triggering apoptosis [97]. Thus, these proteasome inhibitors offer an opportunity to treat cervical cancer by indirectly targeting the oncogenic pathway contributed by HPV oncoprotein E6.

#### **2.3 Histone deacetylase inhibitors**

Histone deacetylases (HDACs) have been examined as targets for cervical cancer therapies. Class I HDACs deacetylate p53 and thereby repress p53-dependent transcriptional activation, apoptosis, and growth arrest [98]. This indicates that inhibition of p53 deacetylation may improve p53 levels, and consequently promote cell cycle arrest and apoptosis of tumor cells. Additionally, E7 oncoprotein binds indirectly to HDAC1 and HDAC2, leading to upregulation of the E2F2 promoter, and consequently HPV viral replication [99,100]. HDAC inhibitors (HDACi) have been shown to induce intrinsic apoptosis in HPV E7-expressing cells [101,102]. Furthermore, HDACi represent a great opportunity for boosting the potency of DNA vaccines. Previous studies have found that HDACi enhance the antitumor effects of therapeutic DNA vaccines in preclinical models by enhancing the expression of the protein/antigen encoded by the DNA vaccine [103,104]. HDACi can also lead to the upregulation of MHC class I and II molecules in treated tumor cells [105]. The upregulation of MHC class I and/or II molecules on tumor cells may render them more susceptible to T cell-mediated killing.

Suberoylanilide hydroxamic acid (SAHA), also known as Vorinostat, is an FDA approved HDACi used in the treatment of cutaneous T-cell lymphoma [106]. SAHA has been tested in human cervical cancer cell lines in combination with bortezomib and the combination of drugs was found to synergistically promote cancer cell apoptosis [94,107]. SAHA has also been tested in combination with cisplatin found similar synergistic killing effects on HeLa cells [108]. Although SAHA has not been clinically tested on cervical cancer patients, it is being tested in another HPV-associated malignancy. A Phase I study examining the effects of SAHA on patients with advanced stage oropharyngeal squamous cell carcinoma is currently recruiting participants and will include analysis of HPV-specific T cell immune responses in patients with HPV+ tumors (NCT01064921).

AR-42 is a novel HDACi that is similar in structure to SAHA and has recently received attention for its potential as an anticancer drug [109]. AR-42 has an optimized structure, which causes it to be a potent inhibitor of HDACs [109]. Indeed, AR-42 was shown to have greater potency and antitumor effects against various cancers in cell culture systems [110-112] as well as in hepatocellular carcinoma [113] *in vivo* compared to clinically available SAHA. Importantly, it has recently been shown that AR-42, but not other clinically available HDACi, can generate potent antigen-specific C8+ T cell-mediated immune responses and antitumor effects against a murine HPV-16 E6/E7-expressing tumor model when combined with a therapeutic HPV DNA vaccine [114]. While the effects of AR-42 on cervical cancer has only been tested in preclinical models to date, patients with acute myeloid leukemia (NCT01798901) and with advanced or relapsed multiple myeloma, chronic lymphocytic leukemia or lymphoma (NCT01129193) are being recruited for testing AR-42 in clinical trials. If these clinical trials demonstrate a high safety profile for AR-42, it potentially can be used in conjunction with a therapeutic HPV vaccine to further improve HPV therapeutic vaccine potency.

HDACi, such as Trichostatin A, have been used in combination with proteasome inhibitors, such as bortezomib, in HeLa xenografts and were found to have more potent antitumor effects than either drug alone [94]. Thus, it will potentially be rewarding to identify the ideal HDACi and proteasome inhibitors for their combined usage for the control of HPVassociated malignancies.

#### **2.4 siRNA technologies targeting E6/E7**

Small interfering RNA (siRNA) technologies have been widely employed in cancer gene therapy to modulate the expression of targeted proteins (for review, see [115]). Notably,

siRNA has been used to induce selective silencing of E6 and E7 in mammalian cells [116]. Although siRNA therapeutic strategies have been tested quite successfully in cell culture systems, data is limited on the effects of siRNA on HPV-16 E6/E7-expressing tumors in animal models. Nevertheless, a few studies have generated interesting results. HPV16 E6 targeting siRNA was found to significantly reduce tumor growth in CaSki tumor-bearing mice compared to non-specific siRNA [117]. Additionally, Chang et al developed potent siRNAs targeting HPV18 E6 and E7 and found that they substantially suppressed tumor growth when intratumorally injected in nude mice bearing HeLa xenografts [118]. Fuji et al also tested HPV18 E6 and E7-targeting siRNA in SKC-II tumor-bearing nude mice in combination with atelocollagen, which served a carrier [119]. They found that HPV18 E6/ E7 siRNA decreased tumor volume and tumor cell proliferation. A novel method of intravenous delivery of siRNA to cervical tumors within lipid particles protects siRNA from nuclease degradation and was shown to reduce target gene expression by 50% in TC-1 tumor-bearing mice [120].

There are not yet clinical trials testing the efficacy of siRNA against cervical cancer. Currently, siRNA technology remains limited by specific delivery and efficient biofunctionality and repeated administration is often necessary. As such, it is uncertain that clinical translation of this targeted treatment will occur soon.

# **3. Conclusion**

HPV oncoproteins E6 and E7 are the most promising targets for the development of targeted therapy against HPV-associated cervical cancer. Numerous immunotherapeutic and other targeted therapeutic strategies are in development. Furthermore, indirect targeting of E6 and E7 by obstructing and/or circumventing their oncogenic functions is also a promising strategy. Indeed, multiple proteasome inhibitors and HDACi are progressing toward clinical translation for the treatment of cervical cancer.

As some of the targeted therapeutic strategies such as immunotherapy targeting E6/E7 are translating to the clinic at different rates, it is worth noting that therapeutic efficacy of these strategies might improve by addressing the immunosuppressive tumor microenvironment in order to improve the antitumor immune response and clinical outcomes. Furthermore, these strategies could be used in conjunction with the current standard of care for cervical cancer, chemotherapy and/or radiation, to more effectively control cervical tumors. As research in the development of targeted therapies for cervical cancer continues, optimal strategies will be created that will drive cervical cancer closer to eradication.

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Knoff et al. Page 18

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**Figure 1. Cervical squamous intraepithelial lesions (SILs) and HPV-associated pathogenesis**

**A**. The normal cervical squamocolumnar junction. The layer of basal cells rests on the basement membrane is the normal barrier between the epithelium and the underlying stromal tissue. The parabasal cells form layers of one to two cells thick just above the basal cell layer. Normal squamous epithelium differentiates as shown, with the nuclear/ cytoplasmic ratio decreasing closer to the surface. The squamocolumnar junction is the most common site for cervical cancer to develop. **B**. Productive infections produce low-grade squamous intraepithelial lesions (LSILs), in which the basaloid cells occupy the lower third of the epithelium. **C**. The cancerous precursor pathway is usually initiated by high-risk HPV infections and produces high-grade squamous intraepithelial lesions (HSILs). HSILs show less cellular differentiation and the basaloid cells occupy at least the lower two thirds and up to the full thickness of the epithelium. Pap smears and HPV tests can be used to detect SILs. **D**. If untreated, premalignant lesions can progress into microinvasive or invasive cancer, in which tumor cells breach the basement membrane. This process is associated with integration of the HPV genome into the host chromosomes, loss of E2 and upregulation of viral oncogene expression and genomic instability.



HPV-infected tumor cell

#### **Figure 2. Summary of targeted therapies**

**A**. Therapeutic HPV vaccines: Vector, peptide, protein, DNA and dendritic cell vaccines generate E6/E7-specific cytotoxic T lymphocyte (CTL) immune responses, which result in the killing of tumor cells presenting antigen on MHC class I molecules. **B**. Proteasome inhibitors: HPV E6 induces ubiquitination of p53 and PDZ family proteins by the E6-AP ubiquitin ligase. HPV E7 induces ubiquitination of Rb by the CUL2 ubiquitin ligase complex. Proteasome inhibitors, such as bortezomib, impede proteasome-mediated degradation of p53 and Rb. **C**. siRNA targeting E6/E7: siRNA specific to E6 and/or E7 lead to the degradiation of E6/E7 mRNA, thereby inhibiting E6/E7 expression. **D**. HDACi: HDACi alleviate HDAC-mediated repression of p53-dependent transcriptional activation, apoptosis, and growth arrest. Additionally, HDACi prevent E7/HDAC1/2-mediated upregulation of HPV viral replication.

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**Table 1**







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Knoff et al. Page 23



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