

## Human Papillomaviruses: Some Genital-Mucosal Types

CAS No.: none assigned

Known to be human carcinogens

First listed in the *Eleventh Report on Carcinogens* (2004)

Also known as HPVs

### Carcinogenicity

Some human papillomaviruses (HPVs) of the genital-mucosal type are *known to be human carcinogens* based on sufficient evidence of carcinogenicity from studies in humans.

### Cancer Studies in Humans

In epidemiological research, numerous case-control studies have consistently reported strong associations between cervical cancer and infection with HPV-16, HPV-18, or “high-risk” HPVs as a class (discussed under Properties, below). Moreover, several case-control studies have provided strong evidence of positive associations between cervical cancer and other individual HPVs, including HPV types 31, 33, 35, 39, 45, 51, 52, 58, and 59 (Muñoz 2000). Cohort studies have demonstrated that infection with HPV-16 or with high-risk HPVs as a class occurs before the development of high-grade cervical intraepithelial neoplasia (CIN), which is thought to be a precursor of invasive cancer. The evidence from cohort studies is weaker for individual high-risk viruses, possibly because they are less common; among these, the evidence for an association with cervical cancer appears to be strongest for HPV-18 (NTP 2003). It is unlikely that the association between HPV infection and cervical cancer is due to other factors that could increase the risk of cancer, because many studies included these factors in their analysis, and because of the large magnitude of the odds ratios estimated in the case-control studies. Thus, these studies demonstrate that some genital-mucosal HPVs cause cervical cancer. In addition to the association with cervical cancer, there is strong evidence that HPV-16 infection is associated with other anogenital cancers, especially cancer of the vulva (NTP 2003). Evidence also suggests associations between HPV infection and some cancers of the head and neck and, especially, the soft palate (oropharynx), tonsils, and back of the tongue and throat (NTP 2003).

Based on testing of tissue specimens from more than 1,000 invasive malignant cervical tumors from women from 22 countries (collected for the International Biological Study of Cervical Cancer), it was estimated that HPV is present in 99.7% of all malignant cervical tumors, suggesting that HPV infection may be necessary for development of cervical cancer (Walboomers *et al.* 1999). Nonetheless, not all individuals infected with HPV develop cervical cancer. Most HPV infections (about 70%) clear within one to two years, and thus confer little risk of cancer. The specific risk factor for cervical cancer appears to be persistent infection with HPV-16 or other high-risk HPVs. Whether HPV infections persist probably depends both on viral characteristics, such as greater persistence of specific HPV types or variants, and on characteristics of the patient, such as sex-hormone levels, smoking behavior, or immune-system status.

Since human papillomaviruses (some genital-mucosal types) were listed in the *Eleventh Report on Carcinogens*, numerous human cancer studies on HPVs have been published. The International Agency for Research on Cancer concluded that HPV types 16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, and 66 were carcinogenic in humans based on sufficient evidence for the carcinogenicity of HPV-16 in the cervix, vulva, vagina, penis, anus, oral cavity, and oropharynx and sufficient evidence for the carcinogenicity of HPV types 18, 31, 33, 35, 39, 45, 51,

52, 56, 58, 59, and 66 in the cervix. IARC also concluded that there was limited evidence for the carcinogenicity of HPV types 6, 11, and 18 in the vulva, penis, and anus; HPV types 6, 11, 16, and 18 in the larynx; HPV-18 in the vagina; and HPV-16 in the periungual skin (the skin around the fingernails or toenails) (IARC 2007).

### Cancer Studies in Experimental Animals

Because HPV infections are specific to humans, experimental animals cannot be infected with them. Many studies have investigated the carcinogenicity of various animal papillomaviruses both in their natural host species and in other species. Studies in monkeys, cattle, rabbits, and sheep have shown that animal papillomaviruses cause cancer in their natural hosts. Studies in transgenic mice carrying HPV genes demonstrated that HPV proteins play a role in the development of abnormal tissue growth (dysplasia) and progression to tumor formation. Transgenic mice expressing some HPV type 16 or 18 genes and producing the corresponding viral proteins developed tumors of the cervix and other tissues (Arbeit *et al.* 1994, Comerford *et al.* 1995).

### Studies on Mechanisms of Carcinogenesis

Infection with high-risk HPVs is associated with chromosomal aberrations, including abnormal centrosome numbers, chromosomal imbalances at specific chromosomal regions, and changes in chromosome number, including tetrasomy and other types of aneuploidy (NTP 2003).

HPV can integrate into the DNA of the host cell and can immortalize and transform cells, enabling them to proliferate and form tumors. Most studies on the mechanisms of HPV carcinogenesis have investigated HPV-16 and HPV-18. HPV types 16, 18, 31, and 33 have been shown to transform cells, types 16, 18, and 31 to immortalize cells, and types 16 and 18 to produce proteins that bind to regulatory proteins of the host cell. The HPV proteins E2 and E5 and the long control region of the HPV genome (discussed under Properties, below) play a role in HPV-induced cell transformation. However, the HPV proteins primarily responsible for immortalization and transformation are E6 and E7, as shown in studies with human and rodent cell cultures. Studies with transgenic mice expressing the E6 or E7 gene further support the notion that the E6 and E7 proteins are important in HPV-associated neoplasia. Both the E6 and E7 proteins alter the pathways that regulate tissue growth, by interfering with growth receptors or growth factors; production of cytokines has been shown to be altered in cells infected with HPV-16. The E6 protein increases degradation of the p53 tumor-suppressor protein, thereby interfering with apoptosis. The E7 protein disrupts complexes of the transcription factor E2F with the tumor-suppressor protein pRb and related proteins involved in control of the cell cycle and causes their degradation, altering control of transcription and progression of the cell cycle. The E7 protein has been shown to cause abnormal synthesis and duplication of centrosomes, resulting in abnormal mitotic division.

### Properties

HPVs of the genital-mucosal type are DNA viruses that infect the genital skin and genital and non-genital mucosa, sometimes causing genital warts or cervical abnormalities. They are members of the family Papillomaviridae, which consists of species-specific non-enveloped viruses that infect the squamous epithelium of the skin and mucosal membranes of animals. More than 100 different HPVs had been identified by 2004, including viruses that cause skin warts as well as the genital-mucosal type (Howley and Lowy 2001). The over 40 genital-mucosal HPVs have been classified as either “high risk” or “low risk”; high-risk viruses have been associated with cervical cancer in human

epidemiological studies, whereas low-risk viruses have been associated with genital warts or low-grade CIN (abnormal tissue growth in the cervical epithelium that is unlikely to progress to cancer). Most studies have considered HPV types 16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, and 68 to be high-risk viruses; some studies also include other HPVs, most notably HPV-66. Classification of HPVs is based also on phylogenetic and mechanistic considerations. Most high-risk viruses have DNA sequences highly similar to those of either HPV-16 or HPV-18, suggesting that they are closely related to these types. Studies on the mechanisms of carcinogenesis have shown that high-risk but not low-risk viruses immortalize human keratinocytes (skin cells), interact with the tumor-suppressor proteins pRb and p53, and cause chromosomal aberrations. However, most mechanistic studies have evaluated only a few HPVs, the majority focusing on HPV-16 or HPV-18 and a few on HPV-31 or HPV-33.

HPVs are small (about 52 to 55 nm in diameter), consisting of about 8,000 base pairs of covalently closed, double-stranded DNA. The viral genome consists of a series of open reading frames, each of which is a DNA sequence that codes for an HPV protein, and a long control region, which contains elements that regulate DNA replication and protein synthesis. Productive infection of cells (leading to replication of the virus) is linked to their stages of differentiation. Viral replication can be divided into early and late stages, which occur in cells at different stages of differentiation. Early stages of replication (including attachment of the virus to the cell, entry and uncoating, early gene expression, protein production, and DNA replication) occur in basal cells. These cells are the youngest, least differentiated cells and are located in the lower layers of the epithelium; they are the only dividing cells in the squamous epithelium. Late stages of viral replication, which include the events leading to production of viral particles (late gene expression, production of capsid proteins, vegetative viral DNA replication, and virus assembly and release), occur in the terminally differentiating squamous epithelial cells, which are the oldest, most differentiated cells, in the upper layers of the epithelium. The genes expressed in the early stages of viral replication, designated E1 through E8, are associated with regulation of transcription (e.g., E2) and cellular proliferation (e.g., E6 and E7). The genes expressed in the late stages, designated L1 and L2, encode the two proteins that make up the viral capsid (Howley and Lowy 2001).

### Infection, Prevention, and Treatment

Genital-mucosal HPVs infect the cervix, causing lesions of varying severity, including genital warts, low- and high-grade CIN, and invasive cervical cancer (Einstein and Burk 2001). Low-grade CIN (CIN I) is a well-differentiated lesion in which the squamous epithelial cells show alterations characteristic of the cytopathogenic effects of a replicative viral infection, such as the presence of two nuclei or other nuclear abnormalities and koilocytosis (the presence of cells with abnormal nuclei and a hollow appearance resulting from collapse of the cell's internal structure). The alterations seen in CIN I are not usually considered to be precursors of cancer. The majority of CIN I lesions are transient and resolve spontaneously, but a small percentage may progress to high-grade CIN or invasive cancer (Jastreboff and Cymet 2002). Both high-risk and low-risk HPVs can cause low-grade CIN (IARC 1995). High-grade CIN (CIN II or CIN III) is characterized by the presence of undifferentiated cells above the lower third of the epithelium (extending into the upper layers) and by nuclear crowding, substantial pleomorphism, loss of tissue organization and cellular polarity, abnormal mitotic figures, and larger numbers of atypical cells than observed in low-grade CIN (IARC 1995). High-grade CIN probably results from persistent HPV infection, and it is more likely than low-grade CIN to progress to invasive cancer. (CIN III is also

known as carcinoma *in situ*, or noninvasive cancer.) Microinvasive squamous-cell cervical cancer usually arises from high-grade CIN.

Two HPV vaccines are licensed by the U.S. Food and Drug Administration and recommended by the Centers for Disease Control and Prevention (CDC 2010). Both vaccines are effective against HPV types 16 and 18, which are responsible for most cervical cancer, and one of the vaccines is also effective against HPV 6 and 11, which cause genital warts. Both vaccines are given in three doses, with the second dose given one to two months after the first and the third dose six months after the first (CDC 2009). Treatment of HPV infection depends on the severity of the disease and may involve topical applications, interferon-related therapies, or excision of the lesion via laser methods, surgery, or cryotherapy.

### Detection

HPV infection is detected by observation of visible lesions or microscopic changes in cells, by detection of HPV DNA, or by detection of antibodies against HPV proteins in the blood. Genital warts (condylomata acuminata) are genital lesions visible to the naked eye; they have a fleshy red appearance and a raised surface that usually extends in papillae. Flat condylomata are flat, nonpapillary lesions; they are more difficult to detect and may be apparent only after swabbing with acetic acid and colposcopic examination, in which they appear as white, flat, shiny lesions. The Papanicolaou (Pap) smear, which involves microscopic examination of stained exfoliated genital cells, detects koilocytosis and other signs of CIN; it is used to screen for cervical cancer by detecting high-grade CIN (Trofatter 1997).

The most sensitive and specific method for detecting HPV infection is to test for HPV DNA. DNA testing can be used to detect a broad spectrum of HPV genotypes (Trofatter 1997). Detection of HPV DNA signifies present exposure or persistent infection resulting from a past exposure. The most sensitive HPV DNA tests are (1) those based on the polymerase chain reaction and (2) the Hybrid Capture assay, which is based on the formation of hybrids between HPV DNA and RNA probes. The most commonly used serological tests for HPV infection measure antibodies (immunoglobulin G) against capsid antigens (most often tested as virus-like particles). Several validation studies have estimated the sensitivity of such serological tests to be approximately 50%, using detection of HPV DNA as a standard (Dillner 2000). Because of their low sensitivity, serological assays are not recommended for diagnostic use, but they are useful for comparison of groups in epidemiological studies, which also commonly use HPV DNA testing. Clinical diagnosis of HPV is most commonly based on the Hybrid Capture 2 assay.

### Exposure

Genital-mucosal HPVs are transmitted primarily through sexual contact with infected cervical, vaginal, vulvar, penile, or anal epithelium (IARC 1995). This finding is supported by numerous epidemiological studies demonstrating that HPV infection is associated with behaviors related to sexual activity. Numerous studies of HPV in women have reported a positive association between lifetime number of sex partners and HPV seropositivity (Sun *et al.* 1999, Silins *et al.* 2000) or the presence of HPV DNA (Franco *et al.* 1995, Kjær *et al.* 1997, Lazcano-Ponce *et al.* 2001). Recent sexual activity, the number of sex partners, frequency of sexual intercourse, and presence of genital warts on sex partners are strong predictors of HPV infection, as indicated by HPV DNA testing (Franco *et al.* 1995, Ho *et al.* 1998). The role of men in carrying HPV infection from one woman to another has been demonstrated in studies showing that cervical cancer is relatively more frequent among wives whose husbands have detectable HPV DNA in their penis or whose husbands have had more

extramarital partners (Bosch *et al.* 1996). Penile lesions containing the DNA of high-risk HPV are frequent among male sex partners of women with CIN (Bleeker *et al.* 2002). There are conflicting reports as to whether HPV is transmitted at birth or perinatally. Infants exposed perinatally to HPV-11, or less commonly to HPV-6, may develop a rare benign tumor of the airway called juvenile-onset recurrent respiratory papillomatosis (Shoultz *et al.* 1997).

HPV infection is one of the most common sexually transmitted diseases. It appears that the majority of those infected have no symptoms, and it is estimated that 20 million people in the United States are infected with HPV (CDC 2001). The percentage of infected individuals (prevalence) is highest among those who are young and sexually active. U.S. epidemiological studies based on HPV DNA testing indicate that between 25% and 40% of sexually active women aged 15 to 25 are infected (Lowy and Howley 2001). Among all U.S. men and women aged 15 to 49, the estimated prevalence of HPV infection (based on HPV DNA testing) is 10% to 20%, whereas only 1% have genital warts, and 4% show cellular abnormalities associated with HPV infection (Koutsky 1997). For most populations of mixed age groups, the prevalence of HPV infection has been estimated at 5% to 15%. HPV-16 appears to be the most prevalent type worldwide (Jastreboff and Cymet 2002). In a study of women aged 18 to 40 with no history of high-grade CIN, among whom the prevalence of HPV was 39%, high-risk HPV were more common (occurring in 26.7% of women) than low-risk HPV (occurring in 14.7%) (Peyton *et al.* 2001).

In 2000, CDC estimated the number of new genital HPV cases per year (incidence) to be 5.5 million (CDC 2001). In the general population of Rochester, Minnesota, the average age- and gender-adjusted incidence of genital warts increased from 13 per 100,000 in the early 1950s to 106 per 100,000 in the late 1970s. During this period, the U.S. incidence of other sexually transmitted diseases also increased dramatically (IARC 1995, Shoultz *et al.* 1997). Several follow-up studies reported very high incidences of HPV infection (as detected by HPV DNA testing) among young, sexually active individuals, with three-year cumulative incidences ranging from 43% to 55% (Ho *et al.* 1998, Moscicki *et al.* 2001).

In most women infected with HPV (70%), the infection clears within 12 to 24 months (Franco *et al.* 1999, Dillner 2000). Some studies have suggested that low-risk HPV infections are more likely to regress than are high-risk HPV infections (Franco *et al.* 1999, Elfgrén *et al.* 2000). The immune system plays an important role in HPV infection; immunocompromised patients are at increased risk for persistent HPV infection (Lowy and Howley 2001).

## Regulations

No specific regulations or guidelines relevant to reduction of exposure to HPV were identified.

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