

PREVALENCE OF VULVOVAGINITIS IN WOMEN
WITH HUMAN PAPILLOMA VIRUS INFECTION

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List of Abbreviations

<i>BV</i>	<i>Bacterial Vaginosis</i>
<i>CIN</i>	<i>Cervical Intraepithelial Neoplasia</i>
<i>HPV 4</i>	<i>Quadrivalent HPV vaccine</i>
<i>HPV</i>	<i>Human Papilloma Virus</i>
<i>PAP</i>	<i>Papanicolaou</i>
<i>PCR</i>	<i>Polymerase Chain Reaction</i>
<i>SIL</i>	<i>Squamous Intraepithelial Lesion</i>
<i>STD</i>	<i>Sexually Transmitted Diseases</i>
<i>STI</i>	<i>Sexually Transmitted Infections</i>
<i>TV</i>	<i>Trichomonas Vaginalis</i>
<i>VVC</i>	<i>Vulvovaginal Candidiasis</i>



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INTRODUCTION

Infectious processes in the vagina caused by potentially pathogenic microorganisms are very common and are the result of growing profusion of infectious or saprophytic agents (*Edwards, 1994*).

Evaluation of the vaginal secretions under the microscope may assist in arriving at the correct diagnosis, but it is better to obtain confirmation by culturing, since non-infectious inflammatory processes may present similar symptomatology (*Edwards, 1994; Di Bartolomeo et al., 1997*).

Bacterial vaginosis is the most prevalent cause of abnormal vaginal discharge, affecting women of reproductive age (*Morris et al., 1991*).

Some risk factors have been associated with bacterial vaginosis, including cigarette smoking, use of intrauterine devices, frequent vaginal douches, multiple sexual partners, early age at first intercourse (*Cherpes et al., 1994; Fethers et al., 1994*).

Lesions induced by the human papillomavirus are usually associated with vaginal infections. Among the various microbiological studies regarding genital infections, few in the literature have related the presence of koilocytosis to infections by other microorganisms (*Alves et al., 1994*).

AIM OF THE WORK

To assess the prevalence of vulvovaginitis caused by *Trichomonas vaginalis* (TV), *Candida* species , and bacterial vaginosis (BV) in women with and without human papilloma virus (HPV) infection.

HUMAN PAPILLOMAVIRUS (HPV)

Human papilloma virus are small nonenveloped double stranded DNA virus in the family papovaviridae the circular genome is 8 kilobases in length (*Graziottin et al., 2009*).

HPV have a common gene organization: an early region encoding non-structural genes, the late region for structural genes, and a regulatory region (long control region: LCR). The functions of each viral protein are summarized as. E1 and E2 are cooperatively involved in the initiation of viral DNA replication. E2 also functions as a transcriptional transactivator. E3 and E4 modulate the cell cycle control and contribute to viral genome maintenance. They also contribute to cancer development though E5 and E6 are speculated to modulate the productive phase of the HPV life cycle, their biological roles remain unclear). Both L1 and L2 are capsid proteins (*Wilson, 2009*).

Types of HPV

More than 120 types of HPV have been identified (*de Villers et al., 2004*) and one-third of them target mucosal membranes, mucosa-tropic HPVs can be classified into two types based on their association with malignant

carcinomas: high risk types (16, 18, 31, 33, 39, 45, 52, 58 and 69) and no-or low risk types (6, 11, 42, 44, 53 ... ect) (*Reich et al., 2004*).

Life cycle

The target of a HPV infection is the stratified epithelium. In the normal stratified epithelium, the cell attached to the basal membrane (basal cell) is the only cell that has the potential to proliferate. The basal cell divides into a new basal cell and a daughter cell that is detached from the basal membrane, and the daughter cell launches its differentiation process. The daughter cells exit from the cell cycle and change their gene expression pattern, proceeding to terminal differentiation, then peel off from the epithelium (*Kines et al., 2009*).

The life cycle of HPV is tightly regulated by the differentiation program of the host cell. The duration of incubation for HPV disease is poorly known, for cutaneous warts, experimental inoculation leads to the development of lesions within 3 to 4 months on average, but it can occur as early as 6 weeks or as late as 5 years (*Schiffman et al., 2003*).

Immunity

HPV suppresses its replication to a “maintenance level” or “latent infection mode” in the basal cells, and maintains the DNA synthesis potential of the infected cells detached from the basal membrane to maintain viral genome replication. In terminally differentiated cells, a tremendous level of genome amplification and late gene expression takes place. After completion of virion assembly, the virions are released externally with the cornified cells. One of the reasons for this unique life cycle of HPV is escape from the immune- surveillance system (*Bodily and Laimins, 2011*).

Epidemiology and prevalence of HPV

The estimated prevalence of anogenital tract HPV infections in the United States is 20 million, with an annual incidence of 6,6 million. It has been estimated that 50 to 80% of sexually active adults will acquire a genital tract HPV infection before the age of 30 (*Workowski et al., 2011*).

Mode of transmission

Several groups of studies have clearly shown that HPV is predominantly and largely transmitted through sexual intercourse (*Kjaer et al., 2011*).

HPVs can also be transmitted in other ways especially from mother to child (*Burchel et al., 2007*).

The evidence for the non-sexual transmission of HPVs has been reviewed by several authors concluding that: 1) Genital HPV infections, including genital warts, may occur in sexually native populations such as virgins, infants, and children; 2) There is some evidence of horizontal transmission of low-risk HPVs; 3) Vertical and perinatal transmission of HPVs from mother to child, does exist, although rates are small and vary widely; 4) High-risk genital HPVs have been detected in non-genital mucosa, such as that in the mouth, oropharynx and conjunctiva, and they have been associated with a fraction of cancers of the oral cavity and oropharynx (*Castellsague et al., 2007*).

There are the possible non-sexual horizontal transmission of HPVs, particularly of the low-risk HPV types. Such as contaminated fomites (clothing, sheets, towels, objects, and instruments) but its impact in passing and inducing active infections is most likely small if any (*IARC, 2009*).

Risk factor

The main factors associated with HPV infection are sex, age, race, sociodemographic characteristics, prior sexually transmitted infections (STI), parity, contraceptive methods and smoking (*IARC, 2007*).

The key determinants among women are the number of sexual partners, the age at which sexual intercourse was initiated (*Castellsague et al., 2007*).

Clinical features

Studies have shown that subclinical infections are the most common manifestation of HPV infection. The vast majority of HPV infections regress spontaneously within 2 years in immunocompetent host (*Parkin and Bray, 2007*).

Cervical HPV infection

They are recognized as papillary epithelial proliferations, the lesion may be single or multiple, scattered or confluent, usually detected on the cervical transformation zone (*Reich et al., 2004*).

Anogenital manifestation

The most common clinically apparent manifestation of genital HPV infection is genital warts (**condyloma acuminata**) (*Insinga et al., 2007*).