

Cervical Human Papillomavirus Infection in a Sample of Hispanic Women Living in Puerto Rico: Comparison with Cervical Cytology Reports

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Objective: Persistent infection with high-risk (HR) HPV is a necessary risk factor for the development of cervical cancer. Information on HPV infection is limited in Puerto Rico. This study determined the distribution of HPV types and the association of HR-HPV types with cervical pathology in a clinic-based sample of women in PR.

Methods: Data from 92 female participants aged 18 to 34 years and recruited from the University of Puerto Rico-Gynecology Clinic, were analyzed. Cervical cytology was performed. HPV testing was performed using L1 consensus primer PCR with MY09/MY11 primers and typed by dot-blot hybridization. Logistic regression modeling was used to determine the crude and covariate adjusted association between HR-HPV and cervical pathology.

Results: Twenty percent (n=18) of the patients had abnormal cytology, 45.7% (n=42) were HPV positive, and 30.4% (n=28) were HR HPV-positive. Women infected with HR-risk HPV types were 7.9 (95% CI = 2.5-25.5) times more likely to have abnormal cytology as compared to women without HR infection when adjusted by age and age at first sexual intercourse.

Conclusion: The burden of HPV infection was high, and, as expected, HR HPVs were strongly associated with dysplasia. A population-based study is needed to estimate HPV prevalence and its association with related malignancies in our population. This will be of great value in determining disease burden and will increase awareness of the HPV vaccination in our population. [*PR Health Sci J* 2013;1:3-7]

Key words: Cervical cancer, Prevention, Screening, Self-sampling

Cervical cancer is the fifth most commonly diagnosed cancer in Puerto Rican women (1). When compared to that of the United States (US), the age-adjusted incidence rate of cervical cancer (9.1 per 100,000) in Puerto Rico is close to double that of non-Hispanic Whites (NHW) (5.9 per 100,000), although it is lower than what is found in Hispanics (12.9 per 100,000) living on the US mainland (2).

Cervical cancer is almost entirely preventable and curable if detected early (3). Human papillomavirus (HPV), the most common sexually transmitted infection in the US, is linked etiologically to cervical cancer (4). Epidemiological and laboratory data have demonstrated that persistent infection with high-risk (HR) HPV types is a necessary risk factor for the development of invasive cervical cancer (5). Population-based type-specific HPV prevalence data provide baseline data for determining disease burden in a population and is essential for future assessment of the impact of HPV vaccination programs on cervical cytology (5). Dunne and colleagues reported that the most common HPV types detected in females aged 14 to 59 years from the National Health and

Nutrition Examination Survey in the years 2003-2004 were HPV 62 (3.3%), HPV 84 (3.3%), HPV 53 (2.8%), HPV 89 (2.4%), HPV 61(2.4%) and HPV 16 (1.5%). Overall, HPV types 6, 11, 16 and 18 were detected in 3.4% of the study participants, corresponding to 3.1 million females infected with HPV types included in the current quadrivalent HPV vaccine (4), which has been approved by FDA to prevent HPV infection in males and females from 9 to 26 years old.

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Information on HPV infection and its relation to HPV-related malignancies is very limited for PR; this information is essential for the further development and improvement of prevention and treatment strategies for cervical dysplasia and cervical cancer in the island, including vaccination programs. Thus, to our knowledge, this pilot study describes for the first time the frequency distribution of HPV types (high and low risk), by cervical pathology status, and assesses the association of HR-HPV types with cervical pathology in a clinic-based sample of women in PR.

Materials and Methods

Our study population consisted of 100 consecutive women who visited the Gynecology Clinic of the University of Puerto Rico (UPR) School of Medicine, who qualified for the study, and who consented to participate. This clinic is a general gynecology clinic at a tertiary institution at which gynecological care is provided to women, including those who are considered high-risk cases, those who are institutionalized, and those of lower incomes.

Complete information on HPV status was assessed for 92 women, and thus, data from these were included in the analysis. Inclusion criteria were that the woman be 18 to 34 years old, that she have an intact uterus, that she be sexually active, and that she be without a history of cervical carcinoma or any recent cervical procedures, such as cone biopsy or LEEP biopsy (loop electrical excisional procedure). An interviewer-administered questionnaire was used to collect data on potential demographic and lifestyle risk factors, including history of STI's, while information on sexual practices was collected through a self-administered questionnaire using an Audio Computer Assisted Self-Interview (ACASI) system implemented using the Questionnaire Development System (QDS) (Nova Research Co., Washington D.C.). Cervical HPV and cytology samples were collected from each participant. HPV samples were taken first by a study physician, and subsequently self-sampling was performed by the patients following written instructions (6). Self-collection followed similar procedures to those previously used in other studies (7).

HPV testing of samples was performed using L1 consensus primer PCR with MY09/MY11 primers at the University of California - San Francisco. Positive PCR products were typed by dot-blot hybridization using 39 individual type-specific probes, including oncogenic HPV types (16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58 and 59) and non-oncogenic types (6/11, 26/69, 30, 32/42, 34, 53, 54, 57/2/27, 61, 62, 66, 67, 68, 70, 71, 72, 73, 81, 82, 83, 84, 85, 86/87, 90/106, 97, 102/108, as well as 2 separate mixtures, mix1 contains 7/13/40/43/44/55/74/91, and mix2 contains 3/10/28/29/77/78/94 plus all those HPV types that hybridized only with the consensus probe) defined according to IARC (5) and were based on the results of both the physician- and the self-collected samples. Variables

studied included cervical cytology and HPV infection status. Cervical cytology results were determined to be either normal or abnormal; abnormal cytology included atypical squamous cells of undetermined significance (ASCUS), atypical squamous cells, cannot exclude high grade squamous intraepithelial lesion (ASC-H), low grade squamous intraepithelial lesion (LSIL), high grade squamous intraepithelial lesion (HSIL) and cancer. HPV infection was divided into HR (oncogenic) and low-risk (LR) (non-oncogenic) categories. Descriptive statistics were performed to summarize the demographic and clinical characteristics of the study group. The chi-square statistic was used to assess factors associated with cervical pathology. Logistic regression models were used to assess the crude and covariate-adjusted association between HR-HPV status and cervical cytology. Interaction terms in the model were evaluated with the likelihood ratio test.

Results

The median age in our sample was 26.4 years (percentiles 25 and 75: 23-30), with a mean age at first sexual intercourse of 18.1 \pm 3.0 years (Table 1). The median number of sexual partners in the last year and lifetime was 1.0 (percentiles 25 and 75: 1-2) and 4 (percentiles 25 and 75 : 2-6) partners, respectively. Only one woman (1.1%) aged 18 to 26 years (recommended vaccination ages) had been vaccinated with the HPV vaccine. Meanwhile, 20% of the women had an abnormal Pap test result (6.5% ASCUS, 9.8% LSIL, and 3.3% HSIL).

Regarding HPV results, 54.4% of women were HPV-negative and 45.7% were HPV-positive, with 30.4% (n=28) being positive for HR types only, 15.2% (n=14) positive for both LR and HR HPV types, and 30.4% (n=28) positive for LR HPV types only (Figure 1). The prevalence of HPV infection was higher in patients with abnormal cervical cytology results than among those with normal cervical cytology results (Figure 2). Of the patients with abnormal cytology, 33.3% (6/18) were negative for HR HPV and 66.7% (12/18) were positive for HR HPV types (p=0.002). Among those with HSIL (n=3), two were infected with HPV 16 and one with HPV 52. Of the patients with one or more HR types and an abnormal Pap smear, 11.1% (n=2/18) were positive for just one type, 44.4% (n=8/18) were positive for 2 types, and 11.1% (n=2/18) were positive for four (data not shown). Overall, the most prevalent HR HPV types were 16 (n=10, 10.9%), 52 (n=5, 5.4%), 53 (n=5, 5.4%), 56 (n=6, 6.5%) and 58 (n=5, 5.4%; only 2.2% of women (n=2) were infected with HPV 18. In patients with abnormal cytology, the most prevalent HR HPV types were 16 (22.2%), 18 (11.1%), 51 (11.1%), 52 (11.1%), and 58 (16.7%); in those with normal pathology, the prevalence of these HPV types was lower (Figure 3).

In bivariate analysis, the only factor statistically associated (p<0.05) with abnormal cytology results was infection with HR-HPV types (p=0.003). No significant associations with

abnormal cervical cytology results were observed for the following variables: age, marital status, education, age at first sexual intercourse, smoking status, history of STIs, number of sexual partners in the last year, or use of non-barrier contraceptives ($p>0.05$) (Table 1). In a logistic regression model, after adjusting for age and age at first sexual intercourse, the odds of having an abnormal cytology in patients positive for HR HPV types were 7.9 (95% CI = 2.5-25.5) times the odds of having an abnormal cytology in HR-HPV-negative patients ($p=0.001$) (Table 2). No significant ($p>0.05$) interaction terms were observed in this model.

Discussion

To our knowledge this is the first study to describe the frequency of HPV infection and its relation to cervical cytology in a clinic-based sample of women in PR. As expected, a strong

association of HR-HPV types with abnormal cervical cytology was found. Our study population was a high-risk population, as 20% of patients surveyed had an abnormal cytology result. This percentage is higher than what would be expected in the general US population (5%) (8), although just slightly higher than that reported in a similar clinic-based population of 9,657 women aged 14 to 65 years receiving routine cervical screening in the US (9). The high number of women with abnormal cytology in this study reflects the fact that this is a clinic-based study, and estimates of disease burden are expected to be higher than those in the general population. Furthermore, despite its being a general gynecology clinic, it also is a tertiary care center, one to which low socioeconomic status women—who could be at higher risk for sexually transmitted infections—are frequently referred (10). Nonetheless, despite the potential for selection bias in our study, patients with cervical infection with HR-HPV types had higher odds of having an abnormal cytology than did those without

Table 1. Characteristics of study participants by cytology status (n = 92)*

| Characteristics | Cytology Status n (%) | | Chi-square (p-value) |
|--|-----------------------|-----------------|----------------------|
| | Normal (n=74) | Abnormal (n=18) | |
| Age (years) | | | 0.661 |
| 18-25 | 33 (44.6%) | 7 (38.9%) | |
| 26-34 | 41 (55.4%) | 11 (61.1%) | |
| Education (years) | | | 0.963 |
| <12 | 12 (16.2%) | 3 (16.7%) | |
| ≥12 | 62 (83.8%) | 15 (83.3%) | |
| Marital status | | | 0.618 |
| Single | 44 (60.3%) | 12 (66.7%) | |
| Married/ cohabitating | 29 (39.7%) | 6 (33.3%) | |
| Age at first sexual intercourse (years) | | | 0.127 |
| <18 | 39 (52.7%) | 6 (33.3%) | |
| ≥18 | 35 (47.3%) | 12 (66.7%) | |
| Number of sexual partners (last 12 months) | | | 0.679 |
| 0-1 | 53 (71.6%) | 12 (66.7%) | |
| ≥2 | 21 (28.4%) | 6 (33.3%) | |
| Non-barrier contraceptive use | | | 0.837 |
| Yes | 35 (47.3%) | 9 (50.0%) | |
| No | 39 (52.7%) | 9 (50.0%) | |
| Ever smoking** | | | 0.526 |
| Yes | 16 (84%) | 3 (16%) | |
| No | 34 (76%) | 11 (24%) | |
| History of STI's*** | | | 0.545 |
| Yes | 20 (27.0%) | 3 (16.7%) | |
| No | 54 (73.0%) | 15 (83.3%) | |
| HR HPV-DNA | | | 0.002 |
| Negative | 58 (78.4%) | 6 (33.3%) | |
| Positive | 16 (21.6%) | 12 (66.7%) | |

*n<100 due to missing data; **n = 64 due to missing data on smoking status; ***self-reported history of the following STIs: HIV/AIDS, Gonorrhea, Syphilis, Genital warts, Genital Herpes, Chlamydia, Trichomonas, and HPV; no participant reported having HIV/AIDS.

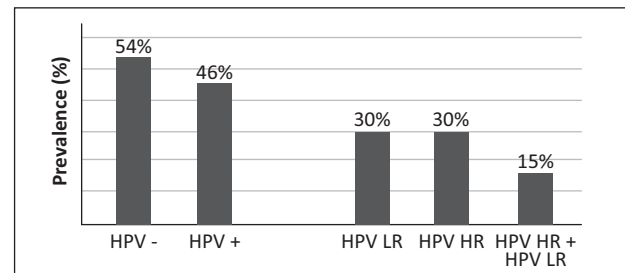


Figure 1. Overall HPV infection results (n=92) (HPV LR = low-risk human papillomavirus, HPV HR = high-risk human papillomavirus)

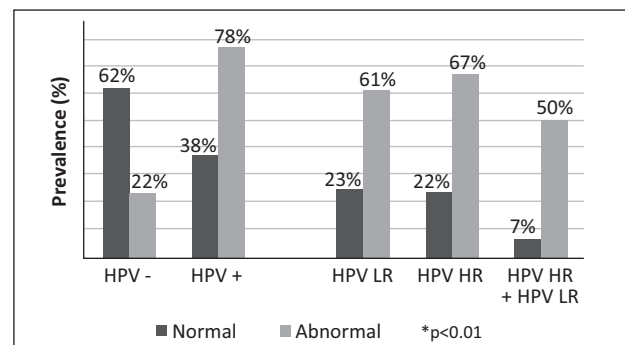


Figure 2. HPV infection in cervical samples, by cervical cytology results (n=92).

Table 2. Logistic regression models of the association of HR-HPV infection and cervical cytology

| Characteristics | Crude POR (95% CI) | Adjusted POR** (95% CI) |
|-----------------|---------------------------|--------------------------|
| HR-HPV-DNA | | |
| Negative* | 1.0 | 1.0 |
| Positive | 7.25 (2.4 - 22.3) p=0.001 | 7.9 (2.5 - 25.5) P=0.001 |

*no HPV or only LR HPV; **Adjusted for age and age at first sexual intercourse

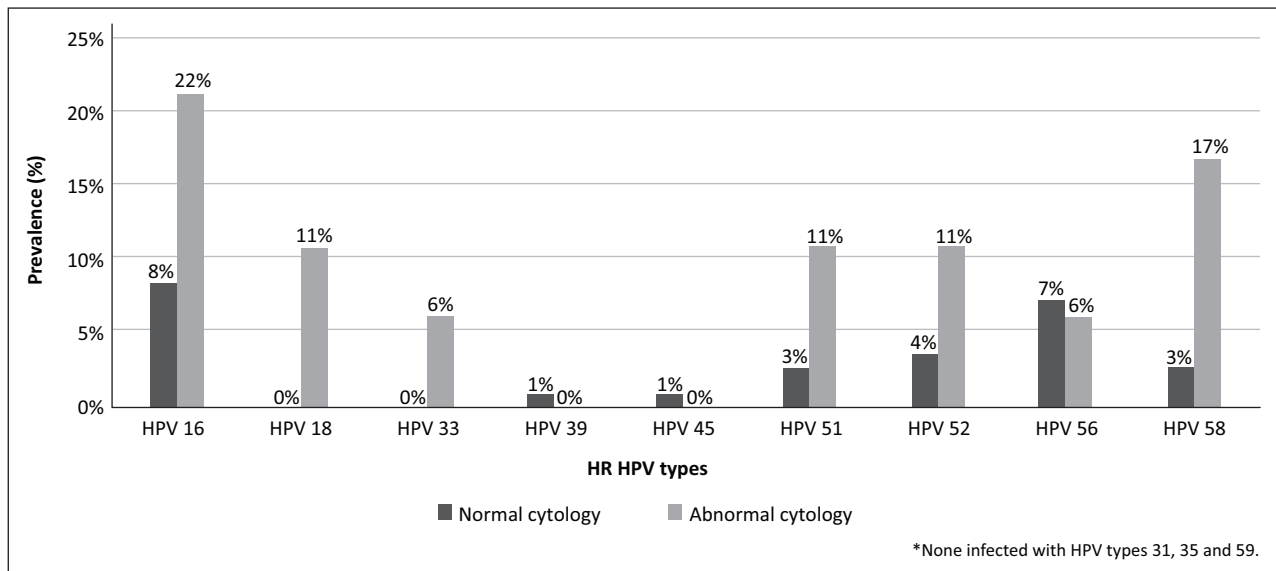


Figure 3. Prevalence of HR HPV types in cervical samples, by cervical cytology results.

HR-HPV infection, after adjusting for age and age at first sexual intercourse, which is consistent with other studies worldwide (11). In addition, this result is consistent with the well-established association between HPV and cervical dysplasia (5, 12).

Disparities in cervical cancer incidence and mortality rates exist among Hispanics and women of African ancestry, as well as in developing Latin American countries and the Caribbean (13). Ortiz and colleagues reported that the higher incidence of cervical cancer observed among US Hispanics, non-Hispanic Blacks, and Puerto Ricans, compared to that of non-Hispanic Whites, could reflect either a potential higher prevalence of HPV infection in these populations or lower screening rates in these groups (2). No population-based data on the prevalence of HPV exist for PR that might explain the burden of cervical cancer in this population. In our sample of 92 Puerto Rican women, 43.5% were positive for HPV infection, which is consistent with our hypothesis that women in PR have a high prevalence of HPV infection. Dunne et al. (4) reported an overall HPV prevalence of 26.8% among a representative sample of females aged 14 to 59 years in the US. Nonetheless, this has to be confirmed in population-based studies, as a higher prevalence of infection is expected in clinic-based samples (5). Despite the high disease burden, only 1.1% of the participants aged 26 years or younger had a history of HPV vaccine administration at the time of the study, which suggests a low uptake of the vaccine in this group, and this only two years after the approval of the first HPV vaccine in the US. As a result of this observation, all the patients in our study who were candidates for the vaccine were referred to the vaccination clinic so that they might receive information about the HPV vaccine and, when interested, receive the vaccine itself.

Among the study's limitations is the fact that subjects recruited for this study were women who underwent evaluation at a single

clinical referral center; thus, our results are not generalizable to the female population of PR. In addition, our small sample size limited the power of our study to assess the association between demographic and lifestyle factors (such as cigarette use, which is a known co-factor for a positive Pap test) and cytology status. Given the cross-sectional nature of our study, we cannot differentiate between incident and persistent HPV infections. Thus, even though many of the HPV infections detected among study participants will regress, some women might have incident infections, which have not yet progressed to premalignant lesions, underestimating the observed odds of the association between HPV infection and cervical malignancies.

In conclusion, our study showed that the burden of HPV infection was high in this clinic-based sample of women in PR, with a high prevalence of HPV types not included in currently available HPV vaccines. Also, as expected, HR HPVs were strongly associated with dysplasia and HPV 16 was the most common type among women with HSIL. A population-based study is needed to estimate HPV prevalence in PR and to assess, as well, its association with related malignancies. This will be of great value in the determination of disease burden and will increase awareness of HPV vaccination in our population.

Resumen

Objetivo: La infección persistente con el virus del papiloma humano (VPH) de alto riesgo es un factor de riesgo necesario para el desarrollo del cáncer de cuello uterino. Información sobre la infección con VPH en Puerto Rico (PR) es bien limitada. En este estudio se determinó la distribución de tipos de VPH y la asociación de estos con patología cervical, en una muestra clínica de mujeres en PR. **Métodos:** Se analizaron los

datos de 92 féminas, entre las edades 18-34 años, reclutadas en las Clínicas de Ginecología de la Universidad de PR. Se llevó a cabo citología cervical. Pruebas de VPH se llevaron a cabo usando la técnica de “PCR” con “primers MY09/MY11” e hibridación “dot-blot”. Modelos de regresión logística se usaron para determinar la asociación cruda y ajustada entre VPH de alto riesgo y la patología cervical. Resultados: Veinte por ciento (n=18) de las pacientes tuvieron una citología anormal, 45.7% (n=42) resultaron positivas para cualquier tipo de VPH y 30.4% (n=28) resultaron positivas para VPH de alto riesgo. Mujeres infectadas con VPH de alto riesgo tuvieron 7.9 (95% CI = 2.5-25.5) veces la posibilidad de tener una citología anormal comparado con mujeres sin la infección de alto riesgo cuando se ajusta por edad y edad al momento de la primera relación sexual. Conclusión: La magnitud de infección con VPH fue alta en esta población y como era esperado, tipos de alto riesgo resultaron estar fuertemente asociados con displasia cervical. Un estudio poblacional es necesario para estimar la prevalencia de VPH en la población de PR y su asociación con malignidades. Esto será de gran valor para determinar la magnitud de la enfermedad y aumentar el conocimiento en nuestra población sobre la necesidad de prevención con la vacuna de VPH.

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