Cross-sectional study of HPV-16 infection in a population-based subsample of Hispanic adults

A P Ortiz,1,2 E R Unger,3 C Muñoz,2 G Panicker,3 G Tortolero-Luna,1 M Soto-Salgado,4 Y Otero,4 E Suárez,2 C M Pérez2

ABSTRACT

Objective: This study aimed to estimate the prevalence and correlates of seropositivity to human papillomavirus (HPV)-16 in a subsample of adults who participated in the parent study Epidemiology of Hepatitis C in the adult population of Puerto Rico (PR).

Setting: The parent study was a population-based household survey aimed to estimate the seroprevalence of hepatitis C and other viral infections (hepatitis A, hepatitis B, HIV, and herpes simplex type 2) in PR (n=1654) between 2005 and 2008.

Participants: A subsample of the last 450 consecutive adults aged 21–64 years, recruited between February 2007 and January 2008, who participated in the parent study and agreed to participate in HPV testing.

Primary and secondary outcome measures: The samples were tested by ELISA for HPV-16 viral-like particle-specific immunoglobulin G. Information on sociodemographic, health, and lifestyle characteristics was collected. Logistic regression modelling was used to estimate the prevalence odds ratio (POR) to assess factors associated to HPV-16 seropositivity.

Results: Prevalence of seropositivity to HPV-16 was 11.3%. Seroprevalence was higher in women (15.8%) than men (5.6%; p=0.001). After adjusting for age and sex, ever smokers (POR 2.06, 95% CI 1.08 to 3.92) and participants with at least five lifetime sexual partners (POR 2.91, 95% CI 1.24 to 6.81) were more likely to be HPV-16 seropositive.

Conclusions: HPV-16 seropositivity is similar to that reported in the USA (10.4%) for NHANES 2003–2004 participants, although different assays were used in these studies. While future studies should evaluate HPV seroprevalence using a larger population-based sample, our results highlight the need to further understand the burden of HPV infection and HPV-related malignancies in PR, population with a low vaccine uptake.

INTRODUCTION

Human papillomavirus (HPV) is the most common sexually transmitted infection worldwide. Persistent infection with certain types of HPV has been established as a necessary cause for cervical cancer1,2 and has been associated with cancer of the anus, vulva, vagina, penis, and oropharynx.3,4 Approximately 5.2% of all cancers worldwide are attributable to HPV infection.3 Moreover, the economic burden of HPV infection is high, and second only to the cost of HIV infection.5 Currently, there are two HPV vaccines licensed for use worldwide that have proven to be effective in preventing HPV infection and progressive disease in people who were previously HPV naïve.6 With widespread use, the vaccines could provide a cost-effective prevention strategy.

Antibody response to HPV infection is considered a key determinant of protective immunity and may play a role as a predictor of HPV-associated cervical neoplasia.2,7 The protective antibody response is mainly typespecific and directed against conformational epitopes of the major capsid protein L1,8 while antibodies to E6 and E7 oncoproteins may be markers of invasive cervical cancer, with tumour stage and mass determining the magnitude of the response.9 Seroconversion

Strengths and limitations of this study

- This study is the first to provide initial insights of the epidemiology of human papillomavirus (HPV) infection by assessing HPV-16 seroprevalence and its correlates in Puerto Rico.
- Although different laboratory assays were used, our estimate of HPV-16 seropositivity for this subsample of adults in Puerto Rico (11.3%) from 2005 to 2008 is comparable with that reported in the USA (10.4%) for persons aged 14–59 years participating in the NHANES 2003–2004.
- Study limitations include the lack of HPV vaccination data of study participants and the modestly sized sample of HPV-16 seropositive individuals, limiting the power of our study to detect significant associations with HPV serology.
occurs several months after detection of HPV-DNA infection; approximately only 60% of women with an incident HPV-DNA infection seroconvert within 18 months after detection. No differences in median time of seroconversion are observed by HPV type, although antibody responses to high-risk HPV types have been found to persist longer.\(^1\)\(^1\)\(^1\) While HPV DNA testing detects current infection, serological testing serves as a useful epidemiological research tool to measure lifetime exposure to HPV infection because antibodies may persist even after the virus has cleared. Although HPV-16 is the most common HPV type detected in most regions of the world,\(^1\)\(^1\)\(^1\) no population-based seroepidemiological studies of HPV-16 infection have been conducted in Puerto Rico (PR). An estimate of natural HPV infection would not only help understand the burden of the disease, but also provide the necessary baseline data for HPV vaccine implementation and monitoring in PR. Hence, the purpose of this pilot study was to estimate prevalence of HPV-16 immunoglobulin G (IgG) responses and factors associated with HPV-16 seropositivity in a cross-section of adults in PR.

**MATERIALS AND METHODS**

**Study design and population**

An island-wide, population-based cross-sectional household survey aimed at estimating the seroprevalence of hepatitis C and other viral infections (hepatitis A, hepatitis B, HIV, and herpes simplex type 2) was performed in PR (n=1654) between 2005 and 2008. Detailed descriptions of the study sampling design and data collection procedures have been previously published.\(^1\)\(^1\)\(^1\) In brief, a cluster sampling design for household surveys using the census tracts of PR was employed, and one individual aged 21–64 years from each selected household was randomly selected to participate in the study. Participants underwent a personal interview and an audio computer-assisted self-interview (ACASI) using QDS (Nova Research Co., Washington DC, USA), and provided a sample of blood for serological testing. For the current analysis, we used residual serum (both reference and test samples) were serially diluted at 1:10, 1:31.6 and 1:100 in 1X TBST with 10% goat serum, 10% Super-Block and 10% insect cell lysate. An optimised concentration of goat antihuman IgG conjugated to alkaline phosphatase (EMD biosciences) diluted in 1X TBST with 10% goat serum and 10% Super-Block was used as the secondary antibody.

Sample antibody titres (IU/mL) were calculated using the parallel line method against a reference sample calibrated to the International Standard 16 (NIBSC 05/134) with known titre of 10 IU/mL.\(^1\)\(^1\)\(^1\)\(^1\)\(^1\)\(^1\)\(^1\)\(^1\)\(^1\)\(^1\)\(^1\)\(^1\) A pooled serum, negative for antibodies to HPV-16, 18, 6, and 11 as tested by an alternate, competitive Luminex assay,\(^1\)\(^1\)\(^1\)\(^1\)\(^1\)\(^1\) was used to establish cut-off value. The cut-off for positive results was set at values greater than or equal to the median antibody titre of the negative control plus two SDs (1.97 IU/mL).

**Study variables**

Demographic characteristics under study included age group (21–34, 35–50, 51–64 years), education (<12 vs ≥12 years), annual family income (<US$20,000 vs ≥US$20,000), health insurance coverage (private, government-sponsored, none) and marital status (single, married/living together, divorced/separated/widowed). Sexual practices (yes/no) included lifetime history of vaginal, anal, and oral sex as well as age of sexual initiation (≤18 vs >18 years) and number of lifetime sexual partners (0–1, 2–4, ≥5). In addition, we calculated the sexual exposure period by subtracting the participant’s age at the first sexual intercourse from the participant’s age at the time of interview. Then, the number of sexual partners was normalised to the sexual exposure period. Smoking status was assessed by a question asking participants whether they have ever smoked in their lifetime.

**Statistical analysis**

To characterise the demographic, clinical, and lifestyle characteristics of study participants, summary measures for continuous variables (mean±SD or median (25th and 75th centiles)) and frequency distributions for categorical variables were computed. Differences between HPV-16 seropositivity groups were assessed using Student’s t-test or Mann-Whitney test for continuous variables, and χ² test for independence or Fisher’s exact test, when appropriate, for categorical variables. Multivariable logistic regression models were fitted to estimate the prevalence odds ratio (POR) with 95% CIs for HPV-16 seropositivity.\(^1\)\(^1\)\(^1\)\(^1\)\(^1\)\(^1\)\(^1\)\(^1\)\(^1\)\(^1\)\(^1\)\(^1\)\(^1\) Variables statistically associated with HPV-16 seropositivity in age and sex-adjusted logistic regression models (p<0.05) were included in the multivariable model. All data were evaluated using Stata for Windows release 12.0 (Stata Corporation, College Station, Texas, USA). No significant interaction terms were found in the multivariable logistic regression models evaluated (p>0.05).

**RESULTS**

**Sociodemographic, clinical and lifestyle characteristics**

The characteristics of the subsample of 450 adults are shown in table 1. Three participants were HIV positive...
Although a comparison between the study subsample and the parent study population showed no significant differences in the clinical and lifestyle characteristics studied, a higher proportion of participants of the subsample reported less than 12 years of education (30.7% vs 23.2%), a government-
based health insurance (51.1% vs 41.9%) and an annual family income below US$20,000 (72.6% vs 63.8%; p<0.05; data not shown).

Seroprevalence of HPV-16

Overall, 11.3% of participants were seropositive to HPV-16. Seroprevalence among women (15.8%) was higher as compared with men (5.9%; p=0.001; figure 1). Median titres (IU/mL) for the whole sample were 6.07 (25th and 75th centiles: 2.78, 13.8); these titres were significantly higher in HPV-positive women (median: 8.39, 25th and 75th centiles: 2.68, 16.01) than in HPV-positive men (median: 3.32, 25th and 75th centiles: 3.04, 7.13; p<0.0001). Although no significant differences (p>0.05) in prevalence were observed across age groups in men or women, the seroprevalence was higher in younger women (figure 1). The mean age of HPV-16 seropositive individuals was lower (38.5±12.7) than among those HPV seronegative (41.9±12.2); this result was marginally significant (p=0.06). In bivariate analysis, no significant differences in seropositivity were observed by education level, household income, marital status, healthcare coverage, place of birth, sexual practices, and smoking status (p>0.05; table 1). After adjusting for age and sex, ever smokers (POR 2.06, 95% CI 1.08 to 3.92) and those reporting at least five lifetime sexual partners (POR 2.78, 95% CI 1.15 to 6.73) were more likely to be HPV-16 seropositive (table 2). Nonetheless, these associations attenuated to non-significance in the multivariable model. Only sex remained significantly associated with HPV-16 seropositivity in multivariable analysis, with women being more likely to be seropositive as compared with men (POR 4.16, 95% CI 1.91 to 9.03; table 2).

DISCUSSION

The first HPV vaccine, which includes HPV-16, was approved by the Food and Drug Administration (FDA) for use in the USA and PR in 2006 for women only (aged 9–26 years) and later in 2008 for men. Our estimate of HPV-16 seropositivity for this subsample of adults in PR (11.3%) from 2005 to 2008 is comparable to those reported in the USA (10.4%) for persons aged 14–59 years participating in the NHANES 2003–2004.17 Our findings are also comparable to those reported in studies worldwide. A study in the Netherlands during 2006–2007 among persons aged ≥14 years found an HPV-16 seroprevalence of 11.3%,11 whereas a study in England among persons aged 10–49 years reported a seroprevalence of 14.7%.18 Also, consistent with the study by Dunne et al,19 the prevalence of HPV seroreactivity in our study was higher among women (15.8%) than men.

<table>
<thead>
<tr>
<th>Characteristics</th>
<th>Crude POR</th>
<th>Age and sex-adjusted POR (95% CI)</th>
<th>Multivariable-adjusted POR* (95% CI)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Age group in years</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>51–64</td>
<td>1.00</td>
<td>–</td>
<td>1.00</td>
</tr>
<tr>
<td>35–50</td>
<td>0.96 (0.43 to 2.12)</td>
<td>–</td>
<td>0.99 (0.42 to 2.32)</td>
</tr>
<tr>
<td>21–34</td>
<td>1.81 (0.87 to 3.79)</td>
<td>–</td>
<td>1.52 (0.67 to 3.47)</td>
</tr>
<tr>
<td>Sex</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Male</td>
<td>1.00</td>
<td>–</td>
<td>1.00</td>
</tr>
<tr>
<td>Female</td>
<td>3.18 (1.58 to 6.37)</td>
<td>–</td>
<td>4.16 (1.91 to 9.03)</td>
</tr>
<tr>
<td>Number of lifetime sex partners</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>0–1</td>
<td>1.00</td>
<td>1.00</td>
<td>1.00</td>
</tr>
<tr>
<td>2–4</td>
<td>1.11 (0.46 to 2.64)</td>
<td>1.18 (0.48 to 2.87)</td>
<td>1.09 (0.44 to 2.68)</td>
</tr>
<tr>
<td>≥5</td>
<td>1.65 (0.73 to 3.71)</td>
<td>2.78 (1.15 to 6.73)</td>
<td>2.36 (0.94 to 5.90)</td>
</tr>
<tr>
<td>History of smoking</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>No</td>
<td>1.00</td>
<td>1.00</td>
<td>1.00</td>
</tr>
<tr>
<td>Yes</td>
<td>1.64 (0.89 to 3.04)</td>
<td>2.06 (1.08 to 3.92)</td>
<td>1.58 (0.79 to 3.16)</td>
</tr>
</tbody>
</table>

*Additionally adjusted by number of lifetime sexual partners and history of smoking; no significant interaction terms were detected in this logistic regression model (p=0.740).

HPV, human papillomavirus; POR, prevalence OR.

Figure 1 Seroprevalence of human papillomavirus-16 by age and sex (n=450).
(5.6%), with fourfold increased odds of infection. In a study performed by Stone et al., HPV-16 seroreactivity was over two times higher in women (17.9%) than in men (7.9%), and constant across all age and racial/ethnic groups evaluated in the study. Moreover, several studies have shown this sex difference in antibody response in all HPV vaccine types (6, 11, 16 and 18). Sex differences in HPV-16 seroreactivity are also observed among high-risk populations. Women attending sexually transmitted diseases clinics had higher prevalence of HPV-16 seropositivity (30.2%) than men (18.7%), supporting that there are biological reasons for men and women differing in serological responses. Potential explanations proposed by Thompson et al. include that men may be: (1) not as susceptible to HPV-16 infection, (2) more able to clear the infection spontaneously without developing a systemic antibody response and (3) less likely to get infected with HPV-16 given that their sexual exposure frequently involves keratinised epithelium (penis) rather than mucosal epithelium (cervix).

In concordance with the literature, HPV-16 seropositivity was also associated with lifetime number of sexual partners, a strong predictor of HPV seropositivity for men and women. In a study performed in Costa Rica, it was found that women with at least three lifetime sexual partners had a twofold increase in the detection of HPV-16 antibodies compared with women with one lifetime sexual partner. Similarly, results from the HPV Infection in Men study showed that men with multiple lifetime male partners (>11 partners) were more likely to be seropositive to HPV-16 (OR=7.74; 95% CI 3.96 to 15.12). We observed a strong association between smoking and HPV-16 seropositivity, with smokers being more likely to be positive for HPV-16 antibodies in this sample. Smoking habits have been identified as a risk factor for HPV infection and seropositivity, but these findings are inconsistent.

Unlike previous studies showing a decline in seropositivity among those aged 50+, we did not observe a significant association between age and HPV-16 seropositivity. This could be due to the small numbers of seropositive persons in this study.

Study limitations include the lack of HPV vaccination data of participants and the sample size; only 51 samples in this study.

REFERENCES


Cross-sectional study of HPV-16 infection in a population-based subsample of Hispanic adults

A P Ortiz, E R Unger, C Muñoz, G Panicker, G Tortolero-Luna, M Soto-Salgado, Y Otero, E Suárez and C M Pérez

BMJ Open 2014 4:
doi: 10.1136/bmjopen-2013-004203

Updated information and services can be found at:
http://bmjopen.bmj.com/content/4/2/e004203

These include:

References
This article cites 22 articles, 9 of which you can access for free at:
http://bmjopen.bmj.com/content/4/2/e004203#BIBL

Open Access
This is an Open Access article distributed in accordance with the Creative Commons Attribution Non Commercial (CC BY-NC 3.0) license, which permits others to distribute, remix, adapt, build upon this work non-commercially, and license their derivative works on different terms, provided the original work is properly cited and the use is non-commercial. See:
http://creativecommons.org/licenses/by-nc/3.0/

Email alerting service
Receive free email alerts when new articles cite this article. Sign up in the box at the top right corner of the online article.

Topic Collections
Articles on similar topics can be found in the following collections

Epidemiology (2038)
Infectious diseases (548)

Notes

To request permissions go to:
http://group.bmj.com/group/rights-licensing/permissions

To order reprints go to:
http://journals.bmj.com/cgi/reprintform

To subscribe to BMJ go to:
http://group.bmj.com/subscribe/