## **UCSF UC San Francisco Previously Published Works**

### **Title**

Risk Factors for Non–Human Papillomavirus (HPV) Type 16/18 Cervical Infections and Associated Lesions Among HPV DNA–Negative Women Vaccinated Against HPV-16/18 in the Costa Rica Vaccine Trial

### **Permalink**

<https://escholarship.org/uc/item/3hd390x6>

### **Journal**

The Journal of Infectious Diseases, 224(3)

### **ISSN**

0022-1899

### **Authors**

Sierra, Mónica S Tsang, Sabrina H Hu, Shangying [et al.](https://escholarship.org/uc/item/3hd390x6#author)

**Publication Date** 2021-08-02

### **DOI**

10.1093/infdis/jiaa768

Peer reviewed



# Risk Factors for Non–Human Papillomavirus (HPV) Type 16/18 Cervical Infections and Associated Lesions Among HPV DNA–Negative Women Vaccinated Against HPV-16/18 in the Costa Rica Vaccine Trial

Mónica S. Sierra,<sup>1,0</sup> Sabrina H. Tsang,<sup>1</sup> Shangying Hu,<sup>1</sup> Carolina Porras,<sup>2</sup> Rolando Herrero,<sup>2,3</sup> Aimée R. Kreimer,<sup>1</sup> John Schussler,<sup>4</sup> Joseph Boland,<sup>1,5</sup> Sarah Wagner,<sup>1,5,©</sup> Bernal Cortes,<sup>2</sup> Ana C. Rodríguez,<sup>6</sup> Wim Quint,<sup>7</sup> Leen-Jan van Doorn,<sup>7</sup> Mark Schiffman,<sup>1</sup> Joshua N. Sampson,<sup>1</sup> and Allan Hildesheim<sup>1,©</sup>; **for the Costa Rica Human Papillomavirus Vaccine Trial (CVT) Group**

<sup>1</sup>Division of Cancer Epidemiology and Genetics, National Cancer Institute, Bethesda, Maryland, USA, <sup>2</sup>Agencia Costarricense de Investigaciones Biomédicas, formerly Proyecto Epidemiológico Guanacaste, Fundación INCIENSA, San José, Costa Rica, <sup>3</sup>Prevention and Implementation Group, International Agency for Research on Cancer, Lyon, France, <sup>4</sup>Information Management Services, Silver Spring, Maryland, USA, <sup>5</sup>Cancer Genomics Research Laboratory, Frederick National Laboratory for Cancer Research, Leidos Biomedical Research Inc, Frederick, Maryland, USA, <sup>6</sup>Independent Consultant, San José, Costa Rica, and <sup>7</sup>DDL Diagnostic Laboratory, Rijswijk, The Netherlands

*Background.* Factors that lead human papillomavirus (HPV) infections to persist and progress to cancer are not fully understood. We evaluated co-factors for acquisition, persistence, and progression of non–HPV-16/18 infections among HPV-vaccinated women.

*Methods.* We analyzed 2153 women aged 18–25 years randomized to the HPV-vaccine arm of the Costa Rica HPV Vaccine Trial. Women were HPV DNA negative for all types at baseline and followed for approximately 11 years. Generalized estimating equation methods were used to account for correlated observations. Time-dependent factors evaluated were age, sexual behavior, marital status, hormonally related factors, number of full-term pregnancies (FTPs), smoking behavior, and baseline body mass index.

*Results.* A total of 1777 incident oncogenic non–HPV-16/18 infections were detected in 12 292 visits (average, 0.14 infections/ visit). Age and sexual behavior–related variables were associated with oncogenic non–HPV-16/18 acquisition. Twenty-six percent of incident infections persisted for ≥1 year. None of the factors evaluated were statistically associated with persistence of oncogenic non–HPV-16/18 infections. Risk of progression to Cervical Intraepithelial Neoplasia grade 2 or worst (CIN2+) increased with increasing age (*P* for trend = .001), injectable contraceptive use (relative risk, 2.61 [95% confidence interval, 1.19–5.73] ever vs never), and increasing FTPs (*P* for trend = .034).

*Conclusions.* In a cohort of HPV-16/18–vaccinated women, age and sexual behavior variables are associated with acquisition of oncogenic non–HPV-16/18 infections; no notable factors are associated with persistence of acquired infections; and age, parity, and hormonally related exposures are associated with progression to CIN2+.

**Keywords.** HPV infection; incidence; persistence; progression; CIN2+; HPV vaccine.

Human papillomavirus (HPV) infection is one of the most common sexually transmitted infections. Most women become infected at some point in their lives, but infections are transient and typically clear within a few months to 2 years. A small fraction of women with persistent, oncogenic HPV infections are at risk of development of high-grade precancerous lesions that may progress to cervical cancer if untreated [1, 2]. Approximately

#### **The Journal of Infectious Diseases® 2021;XX:1–14**

70%–75% of all cervical cancers worldwide are caused by oncogenic HPV types 16 and 18 and the remaining cancers are caused by HPV types 31/33/35/39/45/51/52/56/58/59 [2].

Several viral (eg, viral load, HPV genotypes) and nonviral (eg, smoking, oral contraceptive use, increased number of full-term pregnancies [FTPs]) co-factors have been associated with HPV persistence and progression [1]. However, our understanding of these co-factors and of the natural history of HPV infection is largely driven by studies of HPV-16/18 [2, 3].

Available HPV vaccines (bivalent, quadrivalent, and nonavalent) have the potential to eliminate oncogenic HPV-16/18 infection and associated cervical disease. Evidence from high-income countries indicates that after the introduction of the bivalent and quadrivalent vaccines, the incidence of HPV-16/18 infections is reduced [4]. In some populations, the incidence of the nonvaccine HPV types 31/33/45 has also

Received 8 October 2020; editorial decision 9 December 2020; accepted 11 December 2020; published online December 16, 2020.

Presented in part: 33rd International Papillomavirus Conference (virtual), Barcelona, Spain, July 2020.

Correspondence: Mónica S. Sierra, PhD, MSPH, Infections and Immunoepidemiology Branch, Division of Cancer Epidemiology and Genetics, National Cancer Institute, 9609 Medical Center Drive, RM 6-E206, Rockville, MD 20850 (monica.sierra@nih.gov).

Published by Oxford University Press for the Infectious Diseases Society of America 2020. This work is written by (a) US Government employee(s) and is in the public domain in the US. DOI: 10.1093/infdis/jiaa768

declined, suggesting cross-protection, specifically for the bivalent vaccine [4, 5]. We postulate that HPV infections have independent natural histories; thus, it is important to study whether the natural history of non–HPV-16/18 infections that occur in vaccinated populations differ from that previously observed for HPV-16/18 infections. In the present study, we evaluated risk factors for acquisition, persistence, and progression of non–HPV-16/18 infections in a population of HPV-negative women aged 18–25 years who received the bivalent HPV vaccine.

#### **METHODS**

We investigated this research question in women randomized to the HPV-16/18 vaccine arm of the Costa Rica Vaccine Trial (CVT). To be included in this analysis, women had to be HPV DNA negative for all types at baseline (to approximate an adolescent/naive population at vaccination [2153 of the 3727 women in the HPV arm]) and received at least 1 dose of the bivalent HPV vaccine. Women were actively followed for acquisition, persistence, and progression over a period of approximately 11 years.

#### **CVT Study Design**

CVT (NCT00128661) is a community-based, double-blind, randomized clinical trial aimed to investigate the safety and efficacy of the HPV-16/18 AS04-adjuvanted vaccine (Cervarix; GlaxoSmithKline Biologicals, Rixensart, Belgium) in the prevention of cervical precancers. The trial methodology has been published elsewhere [6]. In brief, 7466 consenting women aged 18–25 years were randomized 1:1 to receive 3 doses of Cervarix (treatment arm) or hepatitis A vaccine (Havrix, GlaxoSmithKline Biologicals, Rixensart, Belgium; control arm). At enrollment and annual follow-up visits, women provided a serum sample and sexually active women also had a pelvic examination to collect exfoliated cervical cells for cytology and HPV DNA infection determination. Women were followed for 4 years in the blinded phase, referred to as CVT. Women were provided cervical cancer screening via cytology with HPV triage of atypical squamous cells of unknown significance [7]. See Supplementary Methods for more information.

After the 4-year visit of CVT, women were invited to participate in an unblinded long-term follow-up (LTFU) study to evaluate the long-term risks and benefits of the prophylactic HPV vaccine over a decade. Detailed rationale and methods for LTFU are published elsewhere [7]. Approximately 80% (2919/3687) of the eligible HPV-vaccinated women were enrolled in the LTFU study. Clinician-collected cervical samples were obtained in years 7, 9, and 11. Once every 2 years, women underwent HPV DNA detection and cervical cancer screening, although more often if clinically indicated [7, 8] (see Supplementary Methods for more details).

During the 11 years of follow-up, women referred to colposcopy had a biopsy taken for histological evaluation. Women with histological Cervical Intraepithelial Neoplasia grade 2 or worst (CIN2+) lesions or with worrisome virologic patterns were treated by loop electrosurgical excisional procedure. Protocols were approved by the United States National Cancer Institute and the Costa Rica institutional review boards [6, 7].

#### **HPV DNA Detection and Genotyping**

Cervical specimens collected during the first 4 years of CVT and the early years of LTFU were tested at Delft Diagnostics Laboratory (Netherlands) using broad-spectrum polymerase chain reaction (PCR)–based HPV DNA testing using the SPF10 DNA enzyme immunoassay (DEIA) system and the LiPA25 line detection system [6, 9, 10]. If HPV DNA results from SPF10 DEIA were positive and LiPA25 was negative for HPV-16/18, specimens were tested for the presence of HPV-16 and HPV-18 using type-specific PCR primer sets [10]. Specimens collected during the later years of the LTFU study were tested at the National Cancer Institute Cancer Genomics Research Laboratory using a next-generation sequencing assay (TypeSeq). TypeSeq detects viral DNA for 51 HPV genotypes [11]. SPF10-LiPA25 and TypeSeq results had a 93.1% positive agreement for detecting any oncogenic HPV type; the agreement for non–HPV-16/18 oncogenic types ranged from 71.4% (HPV-59) to 100% (HPV-58), and no difference in vaccine efficacy was observed when using either test to define outcomes [12]. HPV determination was based on SPF10-LiPA25 for the initial 4 years of the study and on SPF10-LiPA25 or TypeSeq results thereafter. At some timepoints, a woman had HPV results available from both testing methods, SPF10-LiPA25 and TypeSeq. In these cases, we selected the most common method used at that timepoint and the less common method only when it was the sole option. Thus, the primary testing used was SPF10-LiPA25 for year 7 and TypeSeq for years 9 and 11 and intervening clinical management study visits. Sensitivity analyses that utilized alternative approaches to HPV classification yielded comparable results (data not shown).

#### **Outcomes**

An incident (acquisition) HPV infection was defined as a typespecific cervical HPV infection that was not present/detected at the previous scheduled visit (years 1, 2, 3, 4, 7, 9, and 11). Note, there could be multiple incident infections within a woman at a given visit. Persistence of an HPV infection was defined as the incident HPV infection persisting for at least 1 year, with the second detection at least 300 days after incidence without an intervening negative test. Progression was defined as the occurrence of CIN2+ histological diagnosis at a visit with a persistent infection.

#### **Exposures**

At each scheduled study visit, women responded to a structured questionnaire that included socioeconomic indicators, smoking, sexual and reproductive history, and contraceptive use [6, 7]. Factors for acquisition, persistence, and progression of non–HPV-16/18 infections included age, sexual behavior (age at first sexual intercourse [AFI], lifetime number of sexual partners [LNSP], monthly frequency of sex, marital status), contraceptive use (oral contraceptives [OC], injectable contraceptives, condoms, and other infrequent [<5%] contraceptive methods), number of FTPs, smoking behavior (status, intensity, age of initiation), and body mass index (BMI). BMI was categorized as underweight  $(<18.5 \text{ kg/m}^2)$ , normal weight (18.5–24.9  $\text{kg/m}^2$ ), overweight (25–29.9 kg/ m<sup>2</sup>), or obese ( $\geq$ 30 kg/m<sup>2</sup>). All of these exposures were considered time-dependent in the analyses, except for BMI (measured at enrollment).

#### **Statistical Analysis**

We first evaluated the relationship between risk factors and incidence of infection. We categorized each risk factor (eg, for age, we used 18–21, 22–25 years, etc) and then report the corresponding number of scheduled visits (eg, number of visits where the woman was 18-21 years old) and the number of incident infections occurring at those visits. Moreover, we reported the relative risk (RR) of acquiring an incident infection for each category. To calculate the RR, we created a dataset with 121 296 rows  $\approx$  number of women  $\times$  number of scheduled visits × number of infection types. Scheduled visits included follow-up visits at years 1, 2, 3, 4, 7, 9, and 11 (ie, not accelerated screening or colposcopy visits). We estimated the RRs using generalized estimating equations (GEEs) with incident infection as the dependent variable, the risk factor as the independent variable, subject as the cluster, an independent working correlation matrix, and a log-link function [13]. The risk factor was included as a continuous variable to obtain *P* values for trend ( $P_{trend}$ ). Our "base" models were adjusted for HPV type and our "adjusted" models also adjusted for assay type (SPF10-LiPA25/TypeSeq), visit protocol (CVT/LTFU), missed previous visit (yes/no), time since last HPV test, age, smoking, OC use, AFI, LNSP, frequency of sex, number of FTPs, and marital status.

We next evaluated the relationship between risk factors and persistence of an infection. We report the number of incident infections that occurred at scheduled visits (years 1, 2, 3, 4, 7, 9, and 11, excluding accelerated screening and colposcopy visits) within each category and then report the proportion of those infections that persisted for  $\geq$ 1 year. While a persistent infection needed to be identified in a scheduled visit, we used both scheduled visits and accelerated screening visits to confirm persistence. We used GEEs to estimate the RR of persistence.

Our dataset was restricted to incident infections that occurred at least 300 days prior to the final visit and was the first of its type. Our "base" models had no adjustments and our "adjusted" models adjusted for assay type at both incident (ie, visit with incident infection) and test visits (eg, first visit >300 days after incidence), visit protocol, missed visit prior to incident-visit, time between incident and test visits, number of visits between the incident visit and test visit, time between incident visit and its prior visit, age, smoking, OC use, AFI, LNSP, frequency of sex, number of FTPs, and marital status.

Finally, we evaluated the relationship between risk factors and progression to CIN2+. Following the above ideas, we report the number of visits where there was a persistent infection and the proportion of those visits where there was also a CIN2+ lesion (ie, progression). For this analysis, we included all visits with a persistent infection—that is, a persistent infection can be identified at a scheduled visit or at an accelerated screening visit. Women were truncated at their first CIN2+ lesion. We truncated follow-up at the time of treatment, as treatment of lesions interrupts their natural history. We used GEEs to estimate the RR but, because of the limited number of CIN2+ lesions, we only adjusted for age, number of oncogenic HPV persistent infections, and duration of the longest oncogenic HPV infection present.

We present acquisition, persistence, and progression models for oncogenic non–HPV-16/18 infections (types 31/33/35/39/45/51/52/56/58/59) in the main text and models for acquisition and persistence for "any" non–HPV-16/18 infections (oncogenic non–HPV-16/18 types and HPV-6/11/34/40/42/43/44/53/54/66/70/74/68/73) in Supplementary Tables 1 and 2. We tested numerous relationships, and therefore a significance threshold of  $P = .05$  can only be considered suggestive evidence of a relationship. All reported *P* values are 2-sided. Statistical analyses were conducted using PROC GENMOD in SAS version 9.4 software (SAS Institute, Cary, North Carolina).

#### RESULTS

#### **Determinants of Acquisition of Oncogenic Non–HPV-16/18 Infections**

We evaluated the relationship between risk factors and acquisition of HPV infection (Table 1). During the 11 years of follow-up in 2153 vaccinated women, we detected 1777 incident oncogenic non–HPV-16/18 infections during 12 292 visits (average, 0.14 HPV infections/visit). The risk of oncogenic non–HPV-16/18 acquisition decreased with increasing age  $(P_{\text{trend}} < .001)$  and risk of acquisition increased with increasing AFI ( $P_{\text{trend}} < .001$ ) and with increasing LNSP ( $P_{\text{trend}} < .001$ ). Unmarried women had twice the risk of oncogenic non–HPV-16/18 acquisition as married women (RR, 2.08 [95% confidence interval {CI}, 1.81–2.38] vs no).









"Adjusted model includes HPV type, assay type (includes all combinations of a sasay prior infection and assay of incident infection; SPF10–LiPA25/TypeSeq for each), visit protocol (Costa Rica Vaccine Trial/Iong-term follow "Adjusted model includes HPV type, assay type (includes all combinations of acident infection and assay of incident infection; SPF10-LiPA25/TypeSeq for each), visit protocol (Costa Rica Vaccine Tiial/long-term follow-up), visit (yes/nol, time since last HPV test (using a 4 degrees of freedom [*dff* cubic spline of log[time]), visit age (using a 4 df cubic spline of age), smoking, oral contraceptive use, AFI, LINSF, monthly frequency of sexu full-term pregnancies, and marital status.

Diaphragm, sponge, spermicide, intrauterine device, and others. cDiaphragm, sponge, spermicide, intrauterine device, and others.



Table 2. Determinants of Persistence of Oncogenic Non-Human Papillomavirus 16/18 Infections **Table 2. Determinants of Persistence of Oncogenic Non–Human Papillomavirus 16/18 Infections**





°Adjusted model includes assay (includes all combination of assay of prior incident infection, and first assay after 300 days—the assay used to declare persistence; SPF10–LIPA25/TypeSeq for each), number of tests between<br> Adjusted model includes all combination of assay of prior incident infection assay of incident infection, and first assay after 300 days—the assay used to declare persistence; SPF10-LIPA25/TypeSeq for each), number of test nodent and persistent infection (0, 21), time between test prior to incident infection and incident infection (using a 3 degrees of freedom [df] cubic spline of ling[]), time between test for incident infection and first e spline of log[time]], visit protocol (Costa Rica Vaccine Tiral/oncy term follow-up], missed previous visit age (using a 3 of cubic spline of age), smoking, oral contraceptive use, age at first sexual intercourse, LNSF, mon intercourse since last visit, number of full-term pregnancies, and marital status. *PDiaphragm, sponge, spermicide, intrauterine device, and others.* 



Table 3. Determinants of Progression to Cervical Intraepithelial Neoplasia grade 2 or worst (CIN2+) of Oncogenic Non-Human Papillomavirus 16/18 Persistent Infections **Table 3. Determinants of Progression to Cervical Intraepithelial Neoplasia grade 2 or worst (CIN2+) of Oncogenic Non–Human Papillomavirus 16/18 Persistent Infections**





\*Adjusted model includes visit age (using a 3 degrees of freedom cubic spline of diego), time of the longest oncogenic human papillomavirus (HPV) incident persistent infection (above or below the median infection length of Adjusted model includes visit age (using a 3 degrees of freedom cubic spline of age), time of fight one ost no orgenic human papillomavirus (HPV) incident persistent infection (above or below the median infection length of cogenic HPV persistent infections at that visit (1 vs >1).

<sup>b</sup>Diaphragm, sponge, spermicide, intrauterine device, and others. bDiaphragm, sponge, spermicide, intrauterine device, and others.

None of the contraception methods evaluated were statistically associated with oncogenic non–HPV-16/18 acquisition, except for suggestive evidence of a relationship with injectable contraceptive use (RR, 1.15 [95% CI, .99–1.33] ever vs never). Smoking status was not associated with oncogenic non–HPV-16/18 acquisition but among smokers, there was a slight reduction in acquisition with increasing smoking intensity ( $P_{\text{trend}}$  = .057) and a slight increase in risk with increasing age at smoking initiation ( $P_{\text{trend}}$  = .001). We found a statistically insignificant decrease in risk of infection with increasing BMI at enrollment ( $P_{\text{trend}} = .067$ ). Similar associations were obtained for the acquisition of "any" non–HPV-16/18 infection (Supplementary Table 1).

We further explored the combined relationship between AFI, LNSP, and acquisition of oncogenic non–HPV-16/18 infections. We observed positive associations for both AFI and LNSP within strata of the other, suggesting independent effects (data not shown).

#### **Determinants of Persistence of Oncogenic Non–HPV-16/18 Incident Infections**

We evaluated the relationship between risk factors and persistence in 1455 qualifying infections (Table 2); we did not have sufficient follow-up time to define persistence for 322 incident oncogenic non–HPV-16/18 infections. Twenty-six percent (375/1455) of oncogenic non–HPV-16/18 incident infections persisted for 1 year or longer. After adjusting for OC use, sexual behavior variables, marital status, and number of FTPs, none of the factors evaluated had statistically significant associations with persistence. We noted a suggestive evidence of a relationship between OC use (RR, 0.89 [95% CI, .69–1.14] for ever vs never), injectable contraceptive use (RR, 0.88 [95% CI, .70–1.12] for ever vs never), other contraceptive method use (RR, 0.78 [95% CI, .57–1.06] for ever vs never), and persistent oncogenic non–HPV-16/18 infections. Although there was not a linear trend with age at smoking initiation or increasing smoking intensity, the risk of persistence was approximately 3-fold higher for smoking initiation at ≤14 years (vs ≥19 years) and approximately 1.5-fold higher in underweight women (vs normal weight). Similar associations were obtained for the persistence of "any" non–HPV-16/18 infections (Supplementary Table 2).

#### **Determinants of Progression to CIN2+ of Oncogenic Non–HPV-16/18 Incident Persistent Infections**

We evaluated the relationship between risk factors and progression to CIN2+ at 878 follow-up visits with at least 1 persistent infection (Table 3). There were 36 visits with both a persistent infection and CIN2+ diagnosis. After adjusting for number of persistent oncogenic HPV infections and time of the longest oncogenic infection, the risk of progression to CIN2+ increased with increasing age ( $P_{trend}$  = .001). In the models that further

adjusted by age, the use of injectable contraceptives was associated with a 2.6-fold (95% CI, 1.19–5.73) increase in the risk of progression to CIN2+ and OC use with a nonsignificant 1.7 fold increase in risk of progression to CIN2+ (95% CI, .42–7.14) as compared to never users. There was a significant trend of increased risk of progression to CIN2+ with an increased number of FTPs ( $P_{trend} = .034$ ). No apparent associations with progression to CIN2+ were observed for smoking or sexual behavior variables, marital status, condom use, other contraceptive methods, or BMI.

Although limited by the number of CIN2+ cases, to explore whether observed effects for age, injectable contraceptives, and FTPs were independent of each other, we evaluated models in which we mutually adjusted for these variables. The effect of age remained after control for injectable contraceptives and FTPs  $(P_{trend} = .013)$ . The effects of injectable contraceptives and FTPs were evident after control for age but were no longer statistically significant (although patterns remained comparable to those shown in Table 3) upon further control for each other (*P* = .05 for injectable contraceptive use;  $P_{\text{trend}} = .41$  for FTPs).

Finally, we explored risk of progression of persistent HPV-31/35/52/58 to CIN2+ and obtained similar results (data not shown), as these were the most frequent persistent infections (30 of the 36 visits had both a persistent HPV-31/35/52/58 infection and CIN2+ diagnosis). We were unable to evaluate the risk of progression by other specific HPV types because of small sample sizes (Supplementary Table 3).

#### **DISCUSSION**

Much of what we know about the factors associated with acquisition, persistence, and progression of HPV infections is driven by our understanding of HPV-16/18 among unvaccinated women. We evaluated factors associated with acquisition, persistence, or progression of cervical oncogenic non–HPV-16/18 infections in the absence of HPV-16/18, among a cohort of >2000 HPV DNA–negative women vaccinated against HPV-16/18 and followed for 11 years. Our results are consistent with previous work showing that sexual behavior variables determine acquisition of oncogenic non– HPV-16/18 infections; identified no sociodemographic, behavioral, or exogenous factors to be associated with persistence of newly acquired infections; and importantly noted that hormonal/reproductive factors were significantly associated with risk of progression to precancer among women with persistent infections.

With respect to acquisition of non–HPV-16/18 oncogenic infections, our findings largely support previous work among unvaccinated populations that demonstrated a strong association between sexual behavior and HPV acquisition [14– 16], and declining acquisition with increasing age [14–20]. Specifically, we found that increasing LNSP and living as unmarried increased the risk of oncogenic non–HPV-16/18 acquisition. Surprisingly, we note that early AFI was associated with a reduced risk of acquisition, which contrasts with previous studies that consistently observed those with earlier AFI to have an increased risk of HPV acquisition. To fully explore this association, we stratified AFI by LNSP and confirmed that the reduced risk associated with early AFI was observed within each stratum of LNSP, thus confirming that the observed association with AFI was not explained by LNSP. Early AFI can be a marker of risky behaviors; thus, one could speculate that increased HPV exposure at young ages among those who initiate sexual activity earlier would lead to a better ability to control infections that occur in later years. Replication of this finding is needed before drawing strong conclusions.

Even though we evaluated nearly 1500 incident oncogenic non–HPV-16/18 infections and 26% of these infections persisted ≥1 year, none of the risk factors evaluated were consistently associated with persistence of incident infections. We observed significant increases in risk of persistence for women who initiated smoking early and women who were underweight. However, in the absence of an effect for current/ past smokers and a lack of a dose-response relationship for either smoking or BMI, we interpret these results with caution. Further corroborations from additional prospective studies are required before drawing any conclusions. Of note, we were unable to evaluate the host's immunological responses or viral characteristics as a determinant of oncogenic non–HPV-16/18 persistence in our study. It is likely that such responses/characteristics play a role on whether an HPV infection persists or clears [1].

Our findings with respect to factors associated with progression were notable. We found that, independent of age, the number of FTPs had the strongest and most consistent associations with progression to CIN2+, with a significant dose response and RRs increasing to 3 (for any oncogenic non–HPV-16/18) for women who reported >2 FTPs. We also observed that, independent of age, ever use of injectable contraceptives was associated with 2.6-fold increase in the risk of cervical precancer and, although not statistically significant, it is interesting to note that OC use was associated with a near 2-fold increase in risk of cervical precancer. Due to limited numbers, we were unable to fully evaluate whether effects of injectable contraceptives and number of FTPs were independent of each other. These findings, which suggest that endogenous and exogenous hormonal factors are important determinants of progression of persistent oncogenic incident non–HPV-16/18 infections, are largely consistent with studies in which the majority of CIN2+ cases were caused HPV-16/18 [21-30]. More specifically, multiple prevaccination era studies reported a dose-response relationship between pregnancies and cervical precancer/cancer [25, 28–31]. However, studies of unvaccinated women that evaluated the association between hormonal contraceptive use and cervical precancer/cancer yielded mixed results, with some studies showing positive associations [21–26] but others not [27, 32–34].

Possible biological explanations for these findings should be considered. Increased hormonal levels and impaired immune response during pregnancy might facilitate HPV exposure or enhance the role of HPV in cervical carcinogenesis; also, cervical trauma during delivery may explain the increased risk of precursor lesions and cervical cancer [29–31, 35–41]. It is unclear how hormonal contraceptive use might affect HPV acquisition or persistence, but published work points to a promoting effect of estrogens on the cervical carcinogenesis process initiated by HPV infection [21, 26, 33].

Smoking is an established HPV cofactor for the development of precursor lesions and cervical cancer [1]. We did not observe smoking to be associated with progression of persistent oncogenic non–HPV-16/18 infections, perhaps due to the small proportion (16%) of women who reported being ever smokers in our study.

Major strengths of this work are the sample size and follow-up, at ages when HPV acquisition rates are high. Another important strength is our ability to exclude women with prevalent infections at enrollment. This cohort allowed us to investigate the influence of risk factors on each stage in the natural history from HPV acquisition to persistence to progression to CIN2+. Despite the large sample size of our cohort, we were limited by the modest number of incident CIN2+ cases that developed during follow-up and were unable to evaluate individual or phylogenetically related groupings of HPV types. Nonetheless, we found that the most frequently persistent oncogenic HPV types in incident CIN2+ cases were HPV-31/35/52/58 (α-9 group) and HPV-39/59/68 (α-7 group), which have been frequently associated with cervical cancer [42]. Finally, we were unable to evaluate immunological responses to infection or viral characteristics that might be important determinants of persistence/progression.

In conclusion, our results revealed that age and sexual behavior variables are associated with the acquisition of oncogenic non–HPV-16/18 infections; there are no behavioral/ modifiable factors that strongly affect the risk of persistence of acquired infections; and age, parity and hormonally related exposures are associated with the progression of persistent infections to CIN2+. Results from our study may be generalizable to women without evidence of previous HPV exposure at time of vaccination with the bivalent HPV vaccine, a target group of adolescent/naive populations for whom vaccination could have the most impact. As more countries adopt HPV vaccination, understanding the co-factors for persistence and progression of oncogenic non–HPV-16/18 infections becomes more relevant, because current vaccines do not provide protection against all oncogenic HPV types that can cause cervical cancer.

#### Supplementary Data

Supplementary materials are available at *The Journal of Infectious Diseases* online. Consisting of data provided by the authors to benefit the reader, the posted materials are not copyedited and are the sole responsibility of the authors, so questions or comments should be addressed to the corresponding author.

#### **Notes**

*Investigators in the Costa Rica HPV Vaccine Trial (CVT) Group.* Bernal Cortés, Paula González, Rolando Herrero, Silvia E. Jiménez, Carolina Porras, and Ana Cecilia Rodríguez (Agencia Costarricense de Investigaciones Biomédicas, formerly Proyecto Epidemiológico Guanacaste, Fundación INCIENSA, San José, Costa Rica); Allan Hildesheim, Aimée R. Kreimer, Douglas R. Lowy, Mark Schiffman, John T. Schiller, Mark Sherman, and Sholom Wacholder (United States National Cancer Institute, Bethesda, Maryland); Ligia A. Pinto and Troy J. Kemp (Leidos Biomedical Research, Inc, Frederick National Laboratory for Cancer Research, Frederick, Maryland); Mary K. Sidawy (Georgetown University, Washington, D.C.); Wim Quint, Leen-Jan van Doorn and Linda Struijk (DDL Diagnostic Laboratory, the Netherlands); Joel M. Palefsky and Teresa M. Darragh (University of California, San Francisco); and Mark H. Stoler (University of Virginia, Charlottesville).

*Acknowledgments.* We extend a special thanks to the women of Guanacaste and Puntarenas, Costa Rica, who gave of themselves in participating in this effort. In Costa Rica, we acknowledge the tremendous effort and dedication of the staff involved in this project; we specifically acknowledge the meaningful contributions by Carlos Avila, Loretto Carvajal, Rebeca Ocampo, Cristian Montero, Diego Guillen, Jorge Morales, and Mario Alfaro. In the United States, we extend our appreciation to the team from Information Management Services responsible for the development and maintenance of the data system used in the trial and who serve as the data management center for this effort, especially Jean Cyr, Julie Buckland, John Schussler, and Brian Befano. We thank Dr Diane Solomon (CVT: medical monitor and quality control pathologist) for her invaluable contributions during the randomized blinded phase of the trial and the design of the long-term follow-up and Nora Macklin (CVT) and Kate Torres (long-term follow-up) for the expertise in coordinating the study. We thank the members of the Data and Safety Monitoring Board charged with protecting the safety and interest of participants during the randomized, blinded phase of our study (Steve Self, Chair, Adriana Benavides, Luis Diego Calzada, Ruth Karron, Ritu Nayar, and Nancy Roach) and members of the external Scientific HPV Working Group who have contributed to the success of our efforts over the years (Henriette Raventós, Chair, Joanna Cain, Diane Davey, Gypsyamber D'Souza, Elizabeth Fontham, Anne Gershon,

Elizabeth Holly, Silvia Lara, Wasima Rida, Richard Roden, Maria del Rocío Sáenz Madrigal, and Margaret Stanley).

*Disclaimer.* Where authors are identified as personnel of the International Agency for Research on Cancer (IARC)/World Health Organization (WHO), the authors alone are responsible for the views expressed in this article and they do not necessarily represent the decisions, policy, or views of the IARC/ WHO.

*Financial support.* The CVT is a long-standing collaboration between investigators in Costa Rica and the National Cancer Institute (NCI). The trial is sponsored and funded by the NCI (contract N01-CP-11005), with funding support from the National Institutes of Health (NIH) Office of Research on Women's Health. GlaxoSmithKline Biologicals provided vaccine and support for aspects of the trial associated with regulatory submission needs of the company under a Clinical Trials Agreement (FDA BB-IND 7920) during the 4-year, randomized blinded phase of our study. The long-term follow-up was funded by the NCI with support from the NIH Office of Research on Women's Health. The NCI and Costa Rica investigators are responsible for the design and conduct of the study; collection, management, analysis, and interpretation of the data; and preparation of the manuscript.

*Potential conflicts of interest.* S. H. T. is an employee of Merck Sharp & Dohme Corp, a subsidiary of Merck & Co, Inc (Kenilworth, New Jersey) but completed all work associated with this manuscript while employed at the NCI. R. H. reports the CVT was conducted under a clinical trials agreement between the NCI and GSK. The field work in Costa Rica and the work of R. H. at IARC were not funded by GSK and he has not received any funds or in kind contributions from this or any other company. A. C. R. discloses having received consulting fees from the NCI, outside the submitted work. M. S. reports having received HPV typing of specimens from Roche and BD at no cost for studies conducted by NCI. All other authors report no potential conflicts of interest.

All authors have submitted the ICMJE Form for Disclosure of Potential Conflicts of Interest. Conflicts that the editors consider relevant to the content of the manuscript have been disclosed.

#### References

- 1. International Agency for Research on Cancer Working Group on the Evaluation of Carcinogenic Risks to Humans. Biological agents. Volume 100 B. A review of human carcinogens. IARC Monogr Eval Carcinog Risks Hum **2012**; 100:1–441.
- 2. Bruni L, Albero G, Serrano B, et al; CO/IARC Information Centre on HPV and Cancer (HPV Information Centre). Human papillomavirus and related diseases in the world: summary report. 2019. https://www.hpvcentre.net/statistics/reports/XWX.pdf. Accessed 18 March 2019.
- 3. de Sanjosé S, Brotons M, Pavón MA. The natural history of human papillomavirus infection. Best Pract Res Clin Obstet Gynaecol **2018**; 47:2–13.
- 4. Harper DM, DeMars LR. HPV vaccines—a review of the first decade. Gynecol Oncol **2017**; 146:196–204.
- 5. Malagón T, Laurie C, Franco EL. Human papillomavirus vaccination and the role of herd effects in future cancer control planning: a review. Expert Rev Vaccines **2018**; 17:395–409.
- 6. Herrero R, Hildesheim A, Rodríguez AC, et al; Costa Rica Vaccine Trial (CVT) Group. Rationale and design of a community-based double-blind randomized clinical trial of an HPV 16 and 18 vaccine in Guanacaste, Costa Rica. Vaccine **2008**; 26:4795–808.
- 7. Gonzalez P, Hildesheim A, Herrero R, et al; Costa Rica HPV Vaccine Trial (CVT) Group. Rationale and design of a long term follow-up study of women who did and did not receive HPV 16/18 vaccination in Guanacaste, Costa Rica. Vaccine **2015**; 33:2141–51.
- 8. Porras C, Tsang SH, Herrero R, et al; Costa Rica Vaccine Trial Group. Efficacy of the bivalent HPV vaccine against HPV 16/18-associated precancer: long-term follow-up results from the Costa Rica Vaccine Trial. Lancet Oncol **2020**; 21:1643–52.
- 9. Kleter B, van Doorn LJ, Schrauwen L, et al. Development and clinical evaluation of a highly sensitive PCR-reverse hybridization line probe assay for detection and identification of anogenital human papillomavirus. J Clin Microbiol **1999**; 37:2508–17.
- 10. van Doorn LJ, Molijn A, Kleter B, Quint W, Colau B. Highly effective detection of human papillomavirus 16 and 18 DNA by a testing algorithm combining broad-spectrum and type-specific PCR. J Clin Microbiol **2006**; 44:3292–8.
- 11. Wagner S, Roberson D, Boland J, et al. Development of the TypeSeq assay for detection of 51 human papillomavirus genotypes by next-generation sequencing. J Clin Microbiol **2019**; 57:e01794-18.
- 12. Wagner S, Roberson D, Boland J, et al; CVT Group. Evaluation of TypeSeq, a novel high-throughput, low-cost, next-generation sequencing-based assay for detection of 51 human papillomavirus genotypes. J Infect Dis **2019**; 220:1609–19.
- 13. Xue X, Gange SJ, Zhong Y, et al. Marginal and mixed-effects models in the analysis of human papillomavirus natural history data. Cancer Epidemiol Biomarkers Prev **2010**; 19:159–69.
- 14. Ramanakumar AV, Naud P, Roteli-Martins CM, et al; HPV-007 Study Group. Incidence and duration of type-specific human papillomavirus infection in high-risk HPV-naïve women: results from the control arm of a phase II HPV-16/18 vaccine trial. BMJ Open **2016**; 6:e011371.
- 15. Kahn JA, Rosenthal SL, Succop PA, Ho GY, Burk RD. The interval between menarche and age of first sexual intercourse as a risk factor for subsequent HPV infection in adolescent and young adult women. J Pediatr **2002**; 141:718–23.
- 16. Fukuchi E, Sawaya GF, Chirenje M, et al. Cervical human papillomavirus incidence and persistence in a cohort of HIV-negative women in Zimbabwe. Sex Transm Dis **2009**; 36:305–11.
- 17. Sideri M, Igidbashian S, Boveri S, et al. Age distribution of HPV genotypes in cervical intraepithelial neoplasia. Gynecol Oncol **2011**; 121:510–3.
- 18. Baandrup L, Munk C, Andersen KK, Junge J, Iftner T, Kjær SK. HPV16 is associated with younger age in women with cervical intraepithelial neoplasia grade 2 and 3. Gynecol Oncol **2012**; 124:281–5.
- 19. Castle PE, Schiffman M, Wheeler CM, Wentzensen N, Gravitt PE. Human papillomavirus genotypes in cervical intraepithelial neoplasia grade 3. Cancer Epidemiol Biomarkers Prev **2010**; 19:1675–81.
- 20. Wheeler CM, Hunt WC, Joste NE, Key CR, Quint WG, Castle PE. Human papillomavirus genotype distributions: implications for vaccination and cancer screening in the United States. J Natl Cancer Inst **2009**; 101:475–87.
- 21. International Collaboration of Epidemiological Studies of Cervical Cancer. Cervical cancer and hormonal contraceptives: collaborative reanalysis of individual data for 16 573 women with cervical cancer and 35 509 women without cervical cancer from 24 epidemiological studies. Lancet **2007**; 370:1609–21.
- 22. Smith JS, Green J, Berrington de Gonzalez A, et al. Cervical cancer and use of hormonal contraceptives: a systematic review. Lancet **2003**; 361:1159–67.
- 23. Iversen L, Sivasubramaniam S, Lee AJ, Fielding S, Hannaford PC. Lifetime cancer risk and combined oral contraceptives: the Royal College of General Practitioners' Oral Contraception Study. Am J Obstet Gynecol **2017**; 216:580.e1–9.
- 24. Gadducci A, Barsotti C, Cosio S, Domenici L, Riccardo Genazzani A. Smoking habit, immune suppression, oral contraceptive use, and hormone replacement therapy use and cervical carcinogenesis: a review of the literature. Gynecol Endocrinol **2011**; 27: 597–604.
- 25. Syrjänen K, Shabalova I, Naud P, et al; NIS and the LAMS Study Research Groups; analysis of the combined prospective cohort of the NIS; LAMS Studies. Co-factors of high-risk human papillomavirus infections display unique profiles in incident CIN1, CIN2 and CIN3. Int J STD AIDS **2011**; 22:263–72.
- 26. Chung SH, Franceschi S, Lambert PF. Estrogen and ERalpha: culprits in cervical cancer? Trends Endocrinol Metab **2010**; 21:504–11.
- 27. Urban M, Banks E, Egger S, et al. Injectable and oral contraceptive use and cancers of the breast, cervix, ovary, and endometrium in black South African women: case-control study. PLoS Med **2012**; 9:e1001182.
- 28. Luhn P, Walker J, Schiffman M, et al. The role of co-factors in the progression from human papillomavirus infection to cervical cancer. Gynecol Oncol **2013**; 128:265–70.
- 29. Roura E, Travier N, Waterboer T, et al. The influence of hormonal factors on the risk of developing cervical cancer and pre-cancer: results from the EPIC cohort. PLoS One **2016**; 11:e0147029.
- 30. Jensen KE, Schmiedel S, Norrild B, Frederiksen K, Iftner T, Kjaer SK. Parity as a cofactor for high-grade cervical disease among women with persistent human papillomavirus infection: a 13-year follow-up. Br J Cancer **2013**; 108:234–9.
- 31. International Collaboration of Epidemiological Studies of Cervical Cancer. Cervical carcinoma and reproductive factors: collaborative reanalysis of individual data on 16 563 women with cervical carcinoma and 33 542 women without cervical carcinoma from 25 epidemiological studies. Int J Cancer **2006**; 119:1108–24.
- 32. Adhikari I, Eriksson T, Luostarinen T, Apter D, Lehtinen M. Is the risk of cervical atypia associated with the interval between menarche and the start of sexual activity? A population-based cohort study. BMJ Open **2019**; 9:e030091.
- 33. Longatto-Filho A, Hammes LS, Sarian LO, et al. Hormonal contraceptives and the length of their use are not independent risk factors for high-risk HPV infections or highgrade CIN. Gynecol Obstet Invest **2011**; 71:93–103.
- 34. Peng Y, Wang X, Feng H, Yan G. Is oral contraceptive use associated with an increased risk of cervical cancer? An evidence-based meta-analysis. J Obstet Gynaecol Res **2017**; 43:913–22.
- 35. Chen J, Gopala K, Akarsh PK, Struyf F, Rosillon D. Prevalence and incidence of human papillomavirus (HPV) infection before and after pregnancy: pooled analysis of

the control arms of efficacy trials of HPV-16/18 AS04 adjuvanted vaccine. Open Forum Infect Dis **2019**; 6:ofz486.

- 36. Sammarco ML, Del Riccio I, Tamburro M, Grasso GM, Ripabelli G. Type-specific persistence and associated risk factors of human papillomavirus infections in women living in central Italy. Eur J Obstet Gynecol Reprod Biol **2013**; 168:222–6.
- 37. Vaccarella S, Herrero R, Dai M, et al. Reproductive factors, oral contraceptive use, and human papillomavirus infection: pooled analysis of the IARC HPV prevalence surveys. Cancer Epidemiol Biomarkers Prev **2006**; 15:2148–53.
- 38. Stensen S, Kjaer SK, Jensen SM, et al. Factors associated with type-specific persistence of high-risk human papillomavirus infection: a population-based study. Int J Cancer **2016**; 138:361–8.
- 39. International Collaboration of Epidemiological Studies of Cervical Cancer. Cervical carcinoma and sexual behavior: collaborative reanalysis of individual data on 15 461 women with cervical carcinoma and 29 164 women without cervical carcinoma from 21 epidemiological studies. Cancer Epidemiol Biomarkers Prev **2009**; 18: 1060–9.
- 40. International Collaboration of Epidemiological Studies of Cervical Cancer. Comparison of risk factors for invasive squamous cell carcinoma and adenocarcinoma of the cervix: collaborative reanalysis of individual data on 8097 women with squamous cell carcinoma and 1374 women with adenocarcinoma from 12 epidemiological studies. Int J Cancer **2007**; 120:885–91.
- 41. Castellsagué X, Díaz M, Vaccarella S, et al. Intrauterine device use, cervical infection with human papillomavirus, and risk of cervical cancer: a pooled analysis of 26 epidemiological studies. Lancet Oncol **2011**; 12:1023–31.
- 42. Clifford G, Franceschi S. Members of the human papillomavirus type 18 family (alpha-7 species) share a common association with adenocarcinoma of the cervix. Int J Cancer **2008**; 122:1684–5.