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# Effect of Prophylactic Human Papillomavirus (HPV) Vaccination on Oral HPV Infections Among Young Adults in the United States

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BSTRACT

Α

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# Purpose

The incidence of human papilloma virus (HPV)–positive oropharyngeal cancers has risen rapidly in recent decades among men in the United States. We investigated the US population–level effect of prophylactic HPV vaccination on the burden of oral HPV infection, the principal cause of HPV-positive oropharyngeal cancers.

#### Methods

We conducted a cross-sectional study of men and women 18 to 33 years of age (N = 2,627) within the National Health and Nutrition Examination Survey 2011 to 2014, a representative sample of the US population. Oral HPV infection with vaccine types 16, 18, 6, or 11 was compared by HPV vaccination status, as measured by self-reported receipt of at least one dose of the HPV vaccine. Analyses accounted for the complex sampling design and were adjusted for age, sex, and race. Statistical significance was assessed using a quasi-score test.

#### Results

Between 2011 and 2014, 18.3% of the US population 18 to 33 years of age reported receipt of at least one dose of the HPV vaccine before the age of 26 years (29.2% in women and 6.9% in men; P < .001). The prevalence of oral HPV16/18/6/11 infections was significantly reduced in vaccinated versus unvaccinated individuals (0.11% v1.61%;  $P_{adj}$  = .008), corresponding to an estimated 88.2% (95% CI, 5.7% to 98.5%) reduction in prevalence after model adjustment for age, sex, and race. Notably, the prevalence of oral HPV16/18/6/11 infections was significantly reduced in vaccinated versus unvaccinated men (0.0% v2.13%;  $P_{adj}$  = .007). Accounting for vaccine uptake, the population-level effect of HPV vaccination on the burden of oral HPV16/18/6/11 infections was 17.0% overall, 25.0% in women, and 6.9% in men.

#### Conclusion

HPV vaccination was associated with reduction in vaccine-type oral HPV prevalence among young US adults. However, because of low vaccine uptake, the population-level effect was modest overall and particularly low in men.

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# INTRODUCTION

The incidence of oropharyngeal cancer caused by human papillomavirus (HPV) infection has increased rapidly in recent decades in men in the United States as well as numerous other developed countries worldwide.<sup>1</sup> Furthermore, HPV-positive oropharyngeal cancer is projected to become the most common HPV-caused cancer in the United States by 2020, with the majority of the burden in men.<sup>1</sup> More than 70% of the approximately 12,000 oropharyngeal cancers diagnosed annually in the United States are caused by HPV, with approximately 90% of HPV-positive oropharyngeal cancers caused by HPV16 and the remainder caused by other oncogenic HPV types.<sup>1-3</sup> Given the absence of screening and secondary prevention strategies, prophylactic HPV vaccination has the greatest potential to prevent HPV-positive oropharyngeal cancers.<sup>4</sup>

Prophylactic HPV vaccination with the bivalent (HPV16/18), quadrivalent (HPV16/18/6/11), or nonavalent (HPV16/18/6/11/31/33/45/52/58) vaccine is currently recommended for US females and males (quadrivalent and nonavalent) ages 9 to

ASSOCIATED CONTENT



DOI: https://doi.org/10.1200/JCO.2017. 75.0141 26 years.<sup>5</sup> HPV vaccination was recommended for routine use in US females in 2006, permissive use in US males in 2009, and routine use in US males in 2011. These vaccines are indicated for the prevention of genital warts (caused by HPV types 6 and 11) and cervical, anal, vulvar, and vaginal precancers and cancers in females and genital warts and anal precancer and cancer in males.<sup>5</sup> Randomized clinical trials demonstrate > 90% vaccine efficacy in the prevention of anogenital HPV infections and precancerous lesions.<sup>5,6</sup> Surveillance studies also show significant population-level reductions in genital HPV prevalence among young US females in the postlicensure era.<sup>7,8</sup> In contrast, few studies have evaluated the population-level effect of HPV vaccination on oral HPV infections.<sup>9-11</sup>

Herein, we provide a surveillance report of the populationlevel effect of prophylactic HPV vaccination on the burden of oral HPV infections in young US women and men.

## METHODS

We compared oral HPV prevalence in vaccinated versus unvaccinated men and women 18 to 33 years of age within the National Health and Nutrition Examination Survey (NHANES) cycles 2011 to 2012 and 2013 to 2014.<sup>12</sup> The NHANES is a representative cross-sectional, stratified, multistage probability sample of the noninstitutionalized civilian US population.<sup>12</sup>

In the NHANES Mobile Exam Center (MEC),<sup>13</sup> all participants 14 to 69 years of age provided a 10-mL scope/saline oral rinse and gargle sample. DNA from oral rinses was evaluated for the presence of 37 HPV genotypes using PGMY09/11 polymerase chain reaction and Roche Linear Array genotyping.<sup>13</sup> Demographic and behavioral factors were collected through an audio computer-assisted self-interview during the MEC visit.<sup>12</sup> Information on HPV vaccination (receipt, age at vaccination, and doses) was collected from participants 9 to 59 years of age by interviewers during the household visit.<sup>12</sup> Public-use data on both oral HPV infection and vaccination were available in individuals 18 to 59 years of age (Appendix Fig A1, online only).

#### Statistical Analyses

Analyses were restricted to 2,627 individuals 18 to 33 years of age; 33 years was the oldest observed age for individuals who were vaccinated through 26 years of age, the oldest recommended age for vaccination in the United States. Individuals who received at least one dose were considered vaccinated.

All analyses were conducted in SAS (Cary, NC) and SAS-Callable SUDAAN (RTI International, Raleigh, NC). The primary outcome was oral HPV16/18/6/11 prevalence, given the predominant use of the quadrivalent vaccine in the United States through 2014.5,7 Analyses accounted for the complex sampling design and potential bias from exclusions (Fig A1) through use of MEC sample weights, repost-stratified to match the age-bysex-by-race US population distribution. Because of the inclusion of two NHANES cycles in the analyses, per NHANES analytic guidelines, we used the sample weights of each NHANES cycles divided by two. Given the nonrandomized comparisons, the vaccinated and unvaccinated groups differed by demographic and behavioral factors (Table 1), which substantially diminished upon adjustment for age (at NHANES participation), sex, and race (Appendix Tables A1 and A2, online only). Thus, oral HPV comparisons by vaccination status were conducted using binary logistic regression, with model adjustment for age, sex, and race. As alternative analyses, we also used propensity weighting (for vaccination by age, sex, and race), which incorporates fewer assumptions but may have lower statistical power than model adjustment.<sup>14</sup> Additional adjustment for socioeconomic status factors, such as education, did not materially change the results (data not shown).

Comparisons between vaccinated and unvaccinated individuals are presented as infection prevalence and percent reduction in prevalence ([unvaccinated minus vaccinated/unvaccinated]  $\times$  100). Prevaccine-era oral HPV prevalence data are not available in the US population; thus, we used the observed prevalence in unvaccinated individuals as the best-available surrogate for prevaccine-era oral HPV prevalence. We then estimated the total number of infections in the absence of HPV vaccination in the US population (US population size  $\times$  prevalence in unvaccinated), the number of preventable infections at 100% vaccination levels (US population size  $\times$  prevalence in vaccinated), and the number of potentially vaccine-prevented infections at current HPV vaccine–uptake levels (US population size of vaccinated individuals  $\times$  prevalence difference between unvaccinated and vaccinated). As a measure of the population-level effect of HPV vaccination on the burden of oral HPV infection, we estimated the proportion of potentially vaccine-prevented infections at the current HPV vaccine–uptake levels (number of potentially vaccine-prevented infections) total number of potentially vaccination).

Given the sparsity of oral HPV outcomes in the vaccinated group, statistical comparisons were conducted using a quasi-score test.<sup>15</sup> Specifically, the score test and the corresponding P value were computed under the null hypothesis of no difference in oral HPV prevalence between the vaccinated and unvaccinated groups, which allowed pooling of the two groups, thus ameliorating small event rates in either group. Prior simulation studies showed better performance of the score test compared with standard Wald tests for obtaining accurate P values under studies using logistic regression with small event rates, while also accounting for clustered sampling designs.<sup>15</sup> We used logistic regression modeling through either model adjustment or propensity adjustment to account for the imbalance in confounders between the vaccinated and unvaccinated individuals; we then computed the score test P value for the comparison of oral HPV prevalence between vaccinated and unvaccinated individuals. Using the model-adjustment approach, predicted margins were computed to obtain adjusted prevalence and prevalence ratios (vaccinated v unvaccinated), which are directly standardized to the distribution in the United States of the covariates used in the logistic modeling. Statistical significance was assessed at a two-sided P < .05.

# RESULTS

Between 2011 and 2014, 18.3% of the US population 18 to 33 years of age reported receipt of at least one dose of HPV vaccine through the age of 26 years. Vaccination rates were significantly higher in women than men (29.2%  $\nu$  6.9%; P < .001). Vaccinated and unvaccinated individuals significantly differed by demographic and behavioral characteristics in unadjusted analyses (Table 1). However, characteristics were similar upon adjustment for age, sex, and race through either model adjustment (Appendix Table A1) or propensity weighting (Appendix Table A2).

Oral HPV prevalence was assessed at an average of 4.1 years after vaccination. The prevalence of vaccine-type oral HPV infections (HPV16/18/6/11) was significantly reduced in vaccinated versus unvaccinated individuals 18 to 33 years of age (0.11%  $\nu$ 1.61%; model-adjusted P = .008; Table 2). This corresponded to an adjusted (model adjustment for age, sex, and race) prevalence ratio of 8.45 for unvaccinated versus vaccinated individuals (95% CI, 1.06 to 67.22) and an estimated 88.2% (95% CI, 5.7% to 98.5%) reduction in vaccine-type infections among vaccinated individuals. Prevalence of oral HPV16, which accounts for approximately 90% of HPV-positive oropharyngeal cancers, was also nonsignificantly reduced in vaccinated versus unvaccinated individuals (0.11% v 0.94%; model-adjusted P = .063). In contrast, prevalence of 33 nonvaccine HPV types was similar between vaccinated versus unvaccinated individuals (Table 2: 3.98% v 4.74%; model-adjusted P = .24). Event numbers for HPV18, HPV6, HPV11, and types with limited cross

Current classical         Maccinated         Procession         Vaccinated         Vaccinated         Procession         Vaccinated         Procession         Vaccinated         Vaccinated         Vaccinated         Vaccinated         Vaccinated         Vaccinated         Vaccinated         Vaccinated			Overall			Women			Men	
	Characteristic	Vaccinated	Unvaccinated	Ъ*	Vaccinated	Unvaccinated	Р*	Vaccinated	Unvaccinated	P*
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	No. of individuals	496†/11,310,006‡	2,131†/50,516,709‡		3941/9,225,357‡	905†/22,394,263‡		102†/2,084,649‡	1,226†/28,122,446‡	
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	No. of doses, %									
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	-	15.6			11.6			35.2		
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	2	17.6			18.9			10.9		
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	m	66.8			69.4			53.9		
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	Age at vaccination, years§	18.5 (0.29)			18.6 (0.32)			18.1 (0.37)		
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	Range	8-26			10-26			8-26		
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	Time since vaccination, years§	4.1 (0.18)			4.2 (0.18)			3.7 (0.44)		
$\begin{array}{c c c c c c c c c c c c c c c c c c c $	Age at NHANES participation, years	22.5 (0.25)	25.9 (0.16)	< .001	22.7 (0.29)	26.3 (0.28)	< .001	21.8 (0.39)	25.7 (0.16)	< .001
0.01 $163 (0.12)$ $17.1 (0.13)$ $< .001$ $163 (0.12)$ $168 (0.18)$ $0.02$ $0.02$ $0.44$ $1.21$ $2.7$ $0.02$ $63.6$ $54.2$ $65.5$ $57.2$ $0.27$ $13.6$ $14.6$ $13.9$ $12.0$ $0.27$ $13.6$ $14.6$ $13.9$ $12.0$ $0.27$ $13.6$ $14.6$ $13.9$ $12.0$ $0.27$ $13.6$ $14.6$ $13.9$ $12.0$ $0.27$ $13.6$ $14.6$ $12.0$ $13.9$ $12.0$ $0.27$ $7.4$ $11.0$ $2.1$ $13.9$ $12.0$ $0.27$ $7.4$ $11.0$ $14.8$ $13.9$ $12.0$ $0.27$ $10.1$ $6.4$ $90.0$ $91.0$ $91.3$ $0.11$ $0.11$ $0.12$ $14.0.12$ $14.0.12$ $91.0$ $91.3$ $0.0445$ $2.01$ $6.046$ $1.4.0.26$ $1.4.0.64$ $11.0.07$ $0.0109$ $5.6$ $1.4.0.29$ $2.01.029$ $1.14.0.29$	Range	18-33	18-33		18-33			18-33	18-33	
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	Age at sexual debut, years§	16.3 (0.13)	16.9 (0.14)	.001	16.3 (0.12)	17.1 (0.13)	< .001	16.0 (0.29)	16.8 (0.18)	.057
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	Race/ethnicity, %			.002			.044			.40
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	Hispanic	13.5	21.0		13.8	21.6		12.1	20.7	
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	White, non-Hispanic	63.9	55.9		63.6	54.2		65.5	57.2	
30 $9.6$ $8.5$ $10.1$ $37$ $74.5$ $70.2$ $21$ $61.7$ $7.4$ $11.0$ $21$ $55.6$ $61.7$ $7.4$ $11.0$ $21$ $55.6$ $61.7$ $7.4$ $11.0$ $26.6$ $24.4$ $18.1$ $18.8$ $0.81$ $26.6$ $24.4$ $7.7$ $10.1$ $6.4$ $9.0$ $8.7$ $23$ $89.9$ $33.6$ $91.0$ $91.0$ $91.3$ $23$ $6.011$ $6.4$ $7.2$ $91.0$ $91.3$ $23$ $6.011$ $6.4$ $7.2$ $91.0$ $91.3$ $23$ $6.01$ $7.3$ $91.0$ $91.0$ $91.3$ $23$ $6.01$ $1.4$ $6.12$ $1.4$ $6.6$ $6.5$ $6.6$ $7.009$ $5.6$ $1.4$ $0.026$ $1.9$ $1.14$ $0.04$ $7.001$ $6.5$ $0.026$ $1.14$ $0.12$ $0.14$ $0.14$ $0.14$ $0.14$ $0.14$	Black, non-Hispanic	13.7	13.2		13.6	14.6		13.9	12.0	
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	Other	8.9	9.9		9.0	9.6		8.5	10.1	
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	Cigarette use, %			.027			.21			<u> 6</u>
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	Never	72.1	63.7		74.5	70.2		55.6	61.7	
32 $18.1$ $18.8$ $26.6$ $24.4$ $22$ $.081$ $.081$ $.081$ $.081$ $.081$ $27$ $10.1$ $6.4$ $9.0$ $9.0$ $9.13$ $23.6$ $93.6$ $93.6$ $91.0$ $91.3$ $20.066$ $< .001$ $6.5 (0.46)$ $7.9 (1.05)$ $11.4 (0.64)$ $3.0 (0.45)$ $.31$ $4.4 (0.54)$ $7.9 (1.05)$ $11.6 (0.7)$ $5.6$ $1.1 (0.12)$ $.34$ $4.1 (0.56)$ $5.5 (0.55)$ $1.1 (0.07)$ $.42$ $1.2 (0.08)$ $1.1 (0.12)$ $.56$ $1.3 (0.16)$ $1.1 (0.07)$ $.42$ $1.2 (0.08)$ $1.1 (0.12)$ $.56$ $1.3 (0.16)$ $1.1 (0.07)$ $.12 (0.08)$ $1.1 (0.12)$ $.56$ $1.3 (0.16)$ $1.1 (0.07)$ $1.1 (0.07)$ $.12 (0.08)$ $1.1 (0.12)$ $.56$ $1.3 (0.16)$ $1.1 (0.07)$ $1.1 (0.07)$ $.12 (0.08)$ $1.1 (0.12)$ $.56$ $1.3 (0.16)$ $1.1 (0.07)$ $1.0 (0.17)$ $.18$ $1.5 (0.12)$ $1.5 (0.25)$ $.79$ $2.0 (0.28)$ $2.2 (0.24)$	Former	8.6	13.1		7.4	11.0		14.8	13.9	
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	Current	19.3	23.2		18.1	18.8		26.6	24.4	
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	Ever had sex (vaginal, anal, oral), %			.22			.081			<u> 06</u>
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	No	9.9	7.7		10.1	6.4		9.0	8.7	
3.9 (0.66)       < .001	Yes	90.1	92.3		89.9	93.6		91.0	91.3	
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	No. of lifetime partners, any type of sex§	6.6 (0.46)	9.9 (0.66)	< .001	6.5 (0.46)	7.9 (1.05)	.16	7.2 (0.97)	11.4 (0.64)	.002
50 (0.45) $.31$ $4.4 (0.54)$ $4.3 (0.65)$ $.94$ $4.1 (0.69)$ $5.5 (0.55)$ $1.1 (0.07)$ $.42$ $1.2 (0.08)$ $1.1 (0.12)$ $.56$ $1.3 (0.16)$ $1.1 (0.07)$ $3.1 (0.63)$ $< .001$ $6.0 (0.42)$ $7.6 (1.04)$ $.11$ $6.8 (1.01)$ $10.4 (0.59)$ $3.1 (0.63)$ $< .001$ $6.0 (0.42)$ $7.6 (1.04)$ $.11$ $6.8 (1.01)$ $10.4 (0.59)$ $1.9 (0.17)$ $.18$ $1.5 (0.12)$ $1.5 (0.25)$ $.79$ $2.0 (0.28)$ $2.2 (0.24)$	No. of recent partners, any type of sex§	1.5 (0.13)	1.7 (0.09)	.56	1.4 (0.12)	1.4 (0.12)	.80	2.0 (0.26)	1.9 (0.14)	.65
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	No. of lifetime oral sex partners§	4.3 (0.50)	5.0 (0.45)	.31	4.4 (0.54)	4.3 (0.65)	.94	4.1 (0.69)	5.5 (0.55)	.12
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	No. of recent oral sex partners§	1.2 (0.08)	1.1 (0.07)	.42	1.2 (0.08)	1.1 (0.12)	.56	1.3 (0.16)	1.1 (0.07)	30
1.9 (0.17)         .18         1.5 (0.12)         1.5 (0.25)         .79         2.0 (0.28)         2.2 (0.24)	No. of lifetime vaginal sex partners§	6.1 (0.45)	9.1 (0.63)	< .001	6.0 (0.42)	7.6 (1.04)	.11	6.8 (1.01)	10.4 (0.59)	.000
Abbreviation: NHANES, National Health and Nutrition Examination Survey. *Wald-F P values from univariable regression models. *Unveighted totals. *Weighted estimates. *Weighted mean (standard error). *Weinstret Column encentrares.	No. of recent vaginal sex partners§	1.6 (0.14)	1.9 (0.17)	.18	1.5 (0.12)	1.5 (0.25)	.79	2.0 (0.28)	2.2 (0.24)	.66
	Abbreviation: NHANES, National Health and *Vald-F <i>P</i> values from univariable regressiv †Unweighted totals. #Weighted estimates.	d Nutrition Examination ion models.	i Survey.							

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		Overall			Women			Men	
HPV Type	Vaccinated	Unvaccinated	P (model adjusted */ propensity adjusted†)	Vaccinated	Unvaccinated	P (model adjusted */ propensity adjusted†)	Vaccinated	Unvaccinated	P (model adjusted */ propensity adjusted†)
No. of individuals	496‡/11,310,006§	2,131‡/50,516,709§		394‡/9,225,357§	905‡/22,394,263§		102 ±/2,084,649§	1,226‡/28,122,446§	
Vaccine type HPV16/18/6/11						.087/.14	a		.007/.003
No. of infections‡	-	32	.008	-	0		0.0	23	
Prevalence, % (95% CI)§	0.11 (0.0 to 0.96)	1.61 (1.00 to 2.47)	.054	0.14 (0.00 to 1.21)	0.97 (0.40 to 1.95)		Ι	2.13 (1.12 to 3.65)	
HPV16			.063/.14			.19/.21	0		.081/.071
No. of infections <sup>‡</sup>	-	19		-	9		0.0	13	
Prevalence, % (95% CI)§	0.11 (0.0 to 0.96)	0.94 (0.47 to 1.67)		0.14 (0.0 to 1.21)	0.71 (0.23 to 1.64)		Ι	1.12 (0.40 to 2.46)	
Nonvaccine types			.24/.52			.42/.58			.93/.83
No. of infections <sup>‡</sup>	24	116		16	23		œ	93	
Prevalence, % (95% CI)§	3.98 (2.42 to 6.13)	4.74 (3.52 to 6.35)		3.70 (1.76 to 6.77)	2.29 (1.36 to 3.60)		5.19 (1.78 to 11.46)	6.69 (4.92 to 9.05)	
Abbreviation: HPV, human papilloma virus. *Binary logistic regression modeling was initially conducted with oral HPV infection as the outcome and age, sex, and race as predictors to account for the imbalance in confounders between vaccinated and unvaccinated individuals. The binary logistic regression models were adjusted for age (linear), sex, and race (Hispanic, non-Hispanic white, non-Hispanic white, non-Hispanic black, and other races). Models in women and men were adjusted for age (linear), sex, and race (Hispanic, non-Hispanic white, non-Hispanic black, and other races). Models for HPV for men were adjusted for age (linear), sex, and race (Hispanic, non-Hispanic white, non-Hispanic black, and other races). Models for HPV for menone and other races). Models for HPV for menonesity model had work conclusted for age (linear), sex, and race (Hispanic, non-Hispanic white, non-Hispanic black, and other races). Models for HPV for menone adjusted for age (linear) and race (Hispanic, non-Hispanic white, non-Hispanic white, non-Hispanic black, and other races). Models for HPV for mercensity and other races). Models for MPV for mercensity and other races). Models for MPV for mercensity and other races). Models for MPV for mercensity and other races) and race (Hispanic, non-Hispanic white, non-Hispanic black, and other races) are redictors. Predictor Area for age (linear) and race (Hispanic, non-Hispanic black, and other races) are redictors. Predictor Area for the unvaccinated and unvaccinated and unvaccinated individuals using a quasi-score test. Sex-specific propensity models included age (linear) and race (Hispanic, non-Hispanic black, and other races) are redictors. Predictor Area for the comparison of race in the vacinated and unvaccinated and unvaccinated and unvaccinated individuals using a quasi-score test. Sex-specific propensity models included age (linear) and race (Hispanic, non-Hispanic black, and other races) are redictors. Predicted odds, and other races). Are the unvaccinated and unvaccinated and unv	illoma virus. deling was initially co. which the score test <i>P</i> drace (Hispanic, non-H at drace (Hispanic, non-H had vaccination as the constitutified National nated and unvaccinate ass: HPV 26, 31, 34, 3	inducted with oral HPV	infection as thu r the comparis, mic black, and c ear) and race nination Surve, si-score test. S 5, 51, 52, 53, F	aral HPV infection as the outcome and age, sex, and race as predictors to account for the imbalance in confounders pouted for the comparison of oral HPV prevalence between vaccinated and unvaccinated individuals. The binary logistic non-Hispanic black, and other races). Models in women and men were adjusted for age (linear) and race (Hispanic, non-Hispanic black, and other races) to avoid zero cells. age (linear), sex, and race (non-Hispanic white, non-Hispanic black, and other races) to avoid zero cells. Bige (linear), sex, and race (non-Hispanic white, non-Hispanic black, and other races) as the predictors. Predicted tition Examination Survey Mobile Exam Center weights of the unvaccinated individuals, which were then used as weights ing a quasi-score test. Sex-specific propensity models included age (linear) and race (Hispanic, non-Hispanic white, non add a duasi-score test. Sex-specific propensity models included age (linear) and race (Hispanic, non-Hispanic white, non add a duasi-score test. Sex-specific propensity models included age (linear) and race (Hispanic, non-Hispanic white, non add a duasi-score test. Sex-specific propensity models included age (linear) and race (Hispanic, non-Hispanic white, non add a duasi-score test. 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Models in women and men were adjusted for age (linear) and race (Hispanic, non-Hispanic black, and other races) to avoid zero cells. ge (linear) and race (non-Hispanic white, non-Hispanic black, and other races) to avoid zero cells. titon Examination Survey Mobile Exam Center weights of the unvaccinated individuals, which were then used as weights for the comparison of oral ing a quasi-score test. Sex-specific propensity models included age (linear) and race (Hispanic, non-Hispanic black, and other races) as the predictors. Predicted odds from the propensity tition Examination Survey Mobile Exam Center weights of the unvaccinated individuals, which were then used as weights for the comparison of oral ing a quasi-score test. Sex-specific propensity models included age (linear) and race (Hispanic, non-Hispanic black, and other ing a quasi-score test. Sex-specific propensity models included age (linear) and race (Hispanic, non-Hispanic black, and other ing a quasi-score test. 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protection (HPV31/33/45) were too low for reliable estimation (Appendix Table A3, online only).

Results were similar in sex-stratified as well as propensityadjusted analyses, albeit at lower significance levels (Table 2). Notably, prevalence of vaccine-type oral HPV infections (HPV16/ 18/6/11) was significantly reduced in vaccinated men versus unvaccinated men (Table 2: 0.0% v 2.13%; model-adjusted P = .007).

We estimated the population-level effect of HPV vaccination on the burden of vaccine-type oral HPV infections between 2011 and 2014 in the US population of individuals 18 to 33 years of age. These estimates combined the reduction in prevalence among vaccinated individuals with current HPV vaccination rates (Fig 1). HPV vaccination potentially prevented an estimated 169,650 (95% CI, 90,668 to 248,862) oral HPV16/18/6/11 infections, including 76,570 (95% CI, 7,665 to 145,770) among women and 44,403 (95% CI, 19,919 to 68,750) among men. The corresponding estimate of population-level effect was 17.0% overall, 25.0% in women, and 6.9% in men. For HPV16, an estimated 93,873 (95% CI, 27,749 to 158,298) infections were potentially prevented (52,585 [95% CI, -10,185 to 115,241] among women and 23,348 [95% CI, 4,301 to 42,315] among men), corresponding to a population-level effect of 16.2% overall, 23.4% in women, and 6.9% in men.

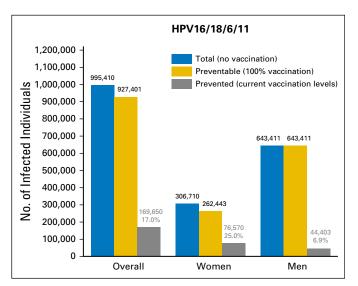


Fig 1. Effect of human papillomavirus (HPV) vaccination on the burden of vaccinetype oral HPV infections in the US population of individuals 18 to 33 years of age, National Health and Nutrition Examination Survey 2011 to 2014. The bar graph shows the estimated effect of prophylactic HPV vaccination on the burden of vaccine-type oral HPV infections (HPV16/18/6/11) among individuals 18 to 33 years of age at participation in the National Health and Nutrition Examination Survey 2011 to 2014. Results are shown overall as well as separately in men and women. Results are presented as the total number of infections in the absence of HPV vaccination in the US population: (US population size × prevalence in unvaccinated) = blue bars; the number of preventable infections at 100% vaccination levels (US population size × prevalence in vaccinated) = gold bars; and the number of potentially vaccineprevented infections at current HPV vaccine-uptake levels (US population size of vaccinated individuals × prevalence difference between unvaccinated and vaccinated) = gray bars. As a measure of the population-level effect of HPV vaccination on the burden of oral HPV infection, we estimated the proportion of potentially vaccine-prevented infections at the current HPV vaccine-uptake levels (number of potentially vaccine-prevented infections/total number of infections in the absence of HPV vaccination). Prevalence estimates used for these computations were not adjusted for any factors. The prevalence of individual vaccine-type HPV infections in the US population of individuals 18 to 33 years of age was as follows: HPV16 = 0.79%, HPV18 = 0.20%, HPV6 = 0.29%, and HPV11 = 0.14%

We evaluated the effect of HPV vaccination on the population-level burden of oral HPV infections in the United States. HPV vaccination was associated with an estimated 88% reduction in prevalence of vaccine-type oral HPV16/18/6/11 infections among vaccinated young adults in the United States. However, because of a vaccination rate of only 18.3% between 2011 and 2014 among individuals 18 to 33 years of age, the population-level effect of HPV vaccination on oral HPV16/18/6/11 infections was a modest 17.0%.

Our results are similar to a prior study in Costa Rica that reported a 93% reduction in HPV16/18 prevalence in women who received the bivalent HPV vaccine.<sup>9,11</sup> A recent study by Hirth et al<sup>10</sup> also used NHANES data (2009 to 2014, individuals ages 18 to 30 years) and concluded that vaccine-type oral HPV prevalence was lower in individuals who received the HPV vaccine compared with unvaccinated individuals.<sup>10</sup> Compared with the study by Hirth et al,<sup>10</sup> the unique aspects of our study include the use of statistical methods optimal for the sparse sample sizes of oral HPV infections, adjustment for important confounders (such as age, sex, and race), the conduct of sex-stratified analyses, and an evaluation of the population-level effect of HPV vaccination on the burden of oral HPV infections.

To our knowledge, our study is the first to report a significant reduction (100%) in prevalence of vaccine-type oral HPV infections in vaccinated men. This reduction in vaccine-type oral HPV infections in vaccinated men is of particular importance because HPV-positive oropharyngeal cancer incidence is three to five times higher in men than in women worldwide.<sup>4</sup> HPV vaccination is currently the most promising prevention option to stem the rising tide of HPV-positive oropharyngeal cancers in men. Unfortunately, low vaccine uptake in adult men between 2011 and 2014 translated to a low population-level effect of HPV vaccination (approximately 6.9%) in men. We note that vaccine uptake in US boys 14 to 17 years of age increased in 2015, reaching 49.8%.<sup>16</sup> Nonetheless, uptake in males has remained significantly lower than in females, which in turn has not achieved the high rates needed for herd immunity to males.<sup>16</sup>

We note the limitations of our study. The nonrandomized comparisons of infection prevalence between vaccinated and unvaccinated individuals preclude a causal interpretation of our results. Despite the inclusion of two NHANES cycles, our analyses were performed on the basis of sparse sample sizes for oral HPV infections, particularly in vaccinated individuals. These sparse sample sizes partly reflect the restriction of analyses to a narrow age range (predefined on the basis of HPV vaccination recommendations),<sup>5</sup> the rarity of oral HPV prevalence compared with anogenital HPV prevalence,<sup>7</sup> and importantly, the high efficacy of HPV vaccination in reducing HPV infections.<sup>6</sup> The low oral HPV prevalence also precluded key subgroup analyses, such as analyses stratified by age at vaccination, vaccine doses, and time since vaccination as well as analyses of cross protection against nonvaccine HPV types. Because of the sparsity of data, we could not calculate adjusted estimates of the number of oral HPV infections (in the absence of vaccination and at current vaccination levels) as well as of population-level effect. Also, our analyses were performed on the basis of self-reported receipt of vaccination, which may have resulted in misclassification.<sup>17</sup> Nevertheless, this misclassification would be expected to be nondifferential by oral HPV prevalence status, which may have biased the effect of vaccination

toward the null. Unlike cervical/vaginal HPV infection, for which prevaccine-era prevalence estimates are available in NHANES and other national surveys, there are no prevaccine-era prevalence data for oral HPV infection in the US population. Consequently, we used the observed prevalence of oral HPV infection in unvaccinated individuals as the best-available surrogate for the prevaccine-era prevalence of oral HPV infection, and then modeled the population-level effect of vaccination on oral HPV infections. Despite these limitations, our results have high global public health significance given the rapid increases in recent decades in the incidence of oropharynx cancers, particularly in men in the United States and numerous other developed countries, including Australia, Canada, Denmark, Japan, Sweden, the Netherlands, and the United Kingdom.<sup>1</sup>

Because of the absence of vaccine-efficacy trials, HPV vaccines are not currently indicated for the prevention of oral HPV infection and HPV-positive oropharyngeal cancers. Previous efforts to conduct efficacy trials were thwarted by regulatory policy that required clinically identifiable surrogate end points, such as precancers, which are difficult to detect in the oropharynx. However, a recent report from a US National Cancer Institute–International Agency for Research on Cancer joint workshop endorsed prevention of incident and persistent oral HPV16 as an acceptable end point for efficacy trials.<sup>6</sup> Such efficacy trials would support evidence-based recommendations for the prevention of HPV-positive oropharyngeal cancer and potentially enhance vaccine uptake in males. In conclusion, HPV vaccination was associated with reduction in vaccine-type oral HPV prevalence among young US adults. However, because of low vaccine uptake, the population-level effect was modest overall and particularly low in men.

## AUTHORS' DISCLOSURES OF POTENTIAL CONFLICTS OF INTEREST

Disclosures provided by the authors are available with this article at jco.org.

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Conception and design: Anil K. Chaturvedi, Barry I. Graubard, Tatevik Broutian, Maura L. Gillison

Financial support: Anil K. Chaturvedi, Maura L. Gillison

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Human Services, Centers for Disease Control and

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#### **AUTHORS' DISCLOSURES OF POTENTIAL CONFLICTS OF INTEREST**

## Effect of Prophylactic Human Papillomavirus (HPV) Vaccination on Oral HPV Infections Among Young Adults in the United States

The following represents disclosure information provided by authors of this manuscript. All relationships are considered compensated. Relationships are self-held unless noted. I = Immediate Family Member, Inst = My Institution. Relationships may not relate to the subject matter of this manuscript. For more information about ASCO's conflict of interest policy, please refer to www.asco.org/rwc or ascopubs.org/jco/site/ifc.

Anil K. Chaturvedi No relationship to disclose

Barry I. Graubard Stock or Other Ownership: Medtronic

**Tatevik Broutian** No relationship to disclose

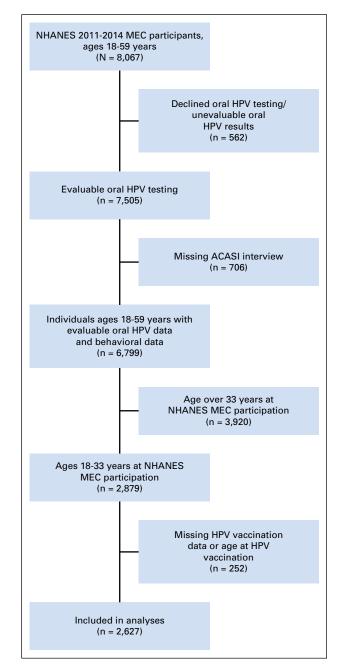
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# Appendix



**Fig A1.** Inclusions/exclusions for the analyses of the effect of human papilloma virus (HPV) vaccination on oral HPV infections, National Health and Nutrition Examination Survey (NHANES) 2011 to 2014. MEC, Mobile Exam Center. ACASI, Audio-computer assisted self-interview.

		Overall			Women			Men	
Characteristic	Vaccinated	Unvaccinated	*d	Vaccinated	Unvaccinated	Ρţ	Vaccinated	Unvaccinated	Ρţ
No. of individuals	496‡/11,310,006§	2,131‡/50,516,709§		394‡/9,225,357§	905‡/22,394,263§		102 #/2,084,649§	1,226‡/28,122,446§	
Age at sexual debut, years	16.6 (0.16)	16.9 (0.13)	.041	16.6 (0.14)	17.0 (0.13)	.017	16.5 (0.33)	16.7 (0.17)	.59
Cigarette use, %			.83			.68			.92
Never	66.7	64.9		73.1	70.7		58.6	58.8	
Former	11.8	12.4		9.6	10.1		16.3	14.6	
Current	21.5	22.7		17.3	19.2		25.1	26.6	
Ever had sex (vaginal, anal, oral), %			.35			.81			.10
No	7.1	8.4		7.2	7.6		5.5	9.1	
Yes	92.9	91.6		92.8	92.4		94.5	90.9	
No. of lifetime partners, any type of sex	8.8 (0.57)	9.4 (0.72)	.61	6.8 (0.54)	7.8 (1.21)	.52	9.7 (0.92)	11.2 (0.65)	.21
No. of recent partners, any type of sex	1.5 (0.15)	1.7 (0.10)	.56	1.3 (0.14)	1.4 (0.15)	.64	1.8 (0.31)	1.9 (0.15)	10
No. of lifetime oral sex partners	5.4 (0.55)	4.7 (0.46)	.33	4.7 (0.59)	4.2 (0.72)	.59	5.5 (0.62)	5.3 (0.55)	.85
No. of recent oral sex partners	1.2 (0.12)	1.1 (0.08)	77.	1.1 (0.14)	1.1 (0.16)	.87	1.4 (0.17)	1.1 (0.07)	.29
No. of lifetime vaginal sex partners	8.1 (0.56)	8.7 (0.69)	.57	6.2 (0.50)	7.5 (1.20)	.40	9.2 (0.95)	10.2 (0.60)	.43
No. of recent vaginal sex partners	1.6 (0.23)	1.9 (0.18)	.44	1.3 (0.20)	1.6 (0.31)	.48	1.9 (0.35)	2.2 (0.24)	.55
NOTE. Values in table are predicted marginals, with standard errors where appropriate. *Wald-F <i>P</i> values from regression models, adjusted for age (linear term), sex, and race tWald-F <i>P</i> values from regression models, adjusted for age (linear term) and race. #Unweighted totals. \$Weighted estimates.	als, with standard error adjusted for age (linear adjusted for age (linear	s where appropriate. term), sex, and race. term) and race.							

#### Chaturvedi et al

		Overall			Women			Men	
Characteristic	Vaccinated	Unvaccinated	*ď	Vaccinated	Unvaccinated	*ď	Vaccinated	Unvaccinated	*ď
No. of individuals	496†/11,310,006‡	2,131†/50,516,709‡		394†/9,225,357‡	905†/22,394,263‡		102†/2,084,649‡	1,226†/28,122,446‡	Ĩ
Age at NHANES participation, years	22.5 (0.25)	22.3 (0.22)	.46	22.7 (0.29)	22.5 (0.27)	.50	21.8 (0.39)	21.7 (0.12)	.92
Age at sexual debut, years	16.3 (0.13)	16.6 (0.10)	.082	16.3 (0.12)	16.7 (0.11)	.052	16.0 (0.29)	16.1 (0.15)	.76
Race/ethnicity, %*			1.0			1.0			1.0
Hispanic	13.5	13.3		13.8	13.5		12.1	11.9	
White, non-Hispanic	63.9	63.7		63.6	63.7		65.5	65.8	
Black, non-Hispanic	13.7	13.8		13.6	13.4		13.9	13.8	
Other	8.9	9.3		9.0	9.4		8.5	8.5	
Cigarette use, %			.64			.53			.97
Never	72.1	69.6		74.5	71.3		61.7	61.2	
Former	8.6	9.0		7.4	8.0		13.9	13.3	
Current	19.3	21.4		18.1	20.7		24.4	25.5	
Ever had sex (vaginal, anal, oral), %			39			.67			.085
No	9.9	12.0		10.1	11.0		9.0	15.0	
Yes	90.1	88.0		89.9	89.0		91.0	85.0	
No. of lifetime partners, any type of sex	6.6 (0.46)	8.1 (1.71)	.33	6.5 (0.46)	8.1 (2.1)	.43	7.2 (0.97)	8.7 (0.97)	.26
No. of recent partners, any type of sex	1.5 (0.13)	1.7 (0.26)	.54	1.4 (0.12)	1.6 (0.31)	.54	2.0 (0.26)	2.1 (0.30)	.81
No. of lifetime oral sex partners	4.3 (0.50)	4.3 (0.92)	.97	4.4 (0.54)	4.4 (1.12)	96.	4.1 (0.69)	3.7 (0.43)	.65
No. of recent oral sex partners	1.2 (0.08)	1.3 (0.28)	.76	1.2 (0.08)	1.4 (0.34)	.64	1.3 (0.16)	1.1 (0.10)	.24
No. of lifetime vaginal sex partners	6.1 (0.45)	7.7 (1.71)	.33	6.0 (0.42)	7.8 (2.1)	.36	6.8 (1.01)	7.8 (0.90)	.43
No. of recent vaginal sex partners	1.6 (0.14)	2.0 (0.48)	.42	1.5 (0.12)	1.9 (0.58)	.45	2.0 (0.28)	2.2 (0.28)	.61
Abbreviation: NHANES, National Health and Nutrition Examination Survey. NOTE. Values in table are weighted means and proportions, with standard errors where appropriate. *Vald-F Pvalues from regression models using propensity weights. The logistic propensity model had vaccination as the outcome and age (linear), sex, and race (Hispanic, non-Hispanic white, non-Hispanic black, and other races) as the predictors. Predicted odds from the propensity model were used to modify the poststratified NHANES Mobile Exam Center weights of the unvaccinated individuals, which were then used as weights for the logistic regression models with human papilloma virus infection as the outcome and vaccination as the predictor. Sex-specific propensity models included age (linear) and race (Hispanic, non-Hispanic, non-Hispanic white, non- Hispanic black, and other races). Huweighted totals.	nd Nutrition Examination ns and proportions, with using propensity weights ds from the propensity m an papilloma virus infecti	n Survey. In standard errors where <i>i</i> s. The logistic propensity u lodel were used to modify on as the outcome and va	appropriate model had , the postst accination <i>i</i>	vaccination as the outr ratified NHANES Mob is the predictor. Sex-sr	come and age (linear), si lie Exam Center weight secific propensity mode	ex, and rac s of the un ls includec	se (Hispanic, non-Hisp vaccinated individuals d age (linear) and race (	urvey. andard errors where appropriate. he logistic propensity model had vaccination as the outcome and age (linear), sex, and race (Hispanic, non-Hispanic white, non-Hispanic black, and he logistic propensity model had vaccination as the outcome and age (linear), sex, and race (Hispanic, non-Hispanic white, non-Hispanic black, and el were used to modify the poststratified NHANES Mobile Exam Center weights of the unvaccinated individuals, which were then used as weights as the outcome and vaccination as the predictor. Sex-specific propensity models included age (linear) and race (Hispanic, non-Hispanic white, non- as the outcome and vaccination as the predictor. Sex-specific propensity models included age (linear) and race (Hispanic, non-Hispanic white, non-	olack, and s weights hite, non-

Primodel adjusted / buncentaried         Primodel adjusted / bun	Overall			Women			Men	
No. of individuals         496±/11,310,0065         2,13           HPV18         0	P ( adji pro Unvaccinated adji	P (model adjusted */ propensity adjusted†)	Vaccinated	Unvaccinated	P (model adjusted */ propensity adjusted†)	Vaccinated	Unvaccinated	P (model adjusted */ propensity adjusted†)
HPV18     0       No. of infections‡     0.0       Prevalence, % (95% Cl)\$     0.0       HPV6     0.0       No. of infections‡     0.0       No. of infections‡     0.0       No. of infections‡     0.0       HPV11     0       No. of infections‡     0.0       HPV13     0       HPV14     0       No. of infections‡     0.0       Prevalence, % (95% Cl)\$     0.0       Prevalence, % (95% Cl)\$     0.0       No. of infections‡     0.0       Prevalence, % (95% Cl)\$     0.0       Prevalence, % (95% Cl)\$     0.0       No. of infections‡     1       Prevalence, % (95% Cl)\$     0.05 (0.0 to 0.85)       Abbreviations: HPV, human papilloma virus; NE, not estima       *Binary logistic regression modeling was initially conducte       urvaccinated individuals, after which he score test P value w       score test P value computation, we did not derive adjusted pn       urvaccinated individuals, after which he score test P value w       score test P value computation, we did not derive adjusted pn       urvaccinated individuals, after which he score test P value w       score test P value computation, we did not derive adjusted pn       urvaccinated individuals, after which he score test P value w	2,131‡/50,516,709§		394‡/9,225,357§	905‡/22,394,263§		102‡/2,084,649§	1,226‡/28,122,446§	
No. of infections‡ 0.0 Prevalence, % (95% CI)\$ - 0.0 HPV6 0 No. of infections‡ 0.0 HPV11 0 HPV11 0 HPV11 0 No. of infections‡ 0.0 HPV11 0 No. of infections‡ 0.0 Prevalence, % (95% CI)\$ 0.0 Prevalence, % (95%		073  /.072	. 0	0	NE/NE	. 0		.073  /.059
Prevalence, % (95% Cl)\$     0     0.2       HPV6     0     0.0       No. of infections‡     0.0       Prevalence, % (95% Cl)\$     0.0       HPV11     0       No. of infections‡     0.0       Prevalence, % (95% Cl)\$     0.0       Prevalence, % (95% Cl)\$     0.0       Prevalence, % (95% Cl)\$     0.0       No. of infections‡     0       Prevalence, % (95% Cl)\$     0.0       Prevalence, % (95% Cl)\$     0.0       Prevalence, % (95% Cl)\$     0.0       Abbreviations: HPV, human papilloma virus; NE, not estima virus; NE, not estima virus initially conducte urvaccinated individuals, after which the score test P value w sco	4		0.0	0.0		0.0	4	
HPV6 0 No. of infections‡ 0.0 Prevalence, % (95% Cl)\$ - 0.3 HPV11 0 No. of infections‡ 0.0 HPV11 0 No. of infections‡ 0.0 Prevalence, % (95% Cl)\$ - 0.1 Cross-protection types HPV31/33/45 1 No. of infections‡ 1 No. of infections‡ 1 Abbreviations: HPV, human papilloma virus; NE, not estima *Binary logistic regression modeling was initially conducte unvaccinated individuals, after which the score test <i>P</i> value w score test <i>P</i> value computation, we did not derive adjusted pri the logistic propensity model had vaccination as the outcore	0.25 (0.05 to 0.71)		Ι	Ι		Ι	0.44 (0.09 to 1.27)	
No. of infections‡ 0.0 Prevalence, % (95% CI)\$ 0.3 HPV11 0 0 No. of infections‡ 0.0 Prevalence, % (95% CI)\$ -0.1 Cross-protection types HPV31/33/45 1 No. of infections‡ 1 Prevalence, % (95% CI)\$ 0.05 (0.0 to 0.85) 0.2 Abbreviations: HPV, human papilloma virus; NE, not estime *Binary logistic regression modeling was initially conducte unvaccinated individuals, after which the score test <i>P</i> value w score test <i>P</i> value white, non-Hispanic black, and othe #The logistic propensity model had vaccination as the outcor	.03	037¶/.049	0		NE/.16	0		.055¶/.052
Prevalence, % (95% Cl)\$     0       HPV11     0       No. of infections‡     0.0       Prevalence, % (95% Cl)\$     0.05 (0.0 to 0.85)       No. of infections‡     1       Prevalence, % (95% Cl)\$     0.05 (0.0 to 0.85)       Abbreviations: HPV, human papilloma virus; NE, not estime unvaccinated individuals, after which the score test <i>P</i> value w concertest <i>P</i> value w three unvaccinated individuals, after white, non-Hispanic black, and other three adjustic propensity model had vaccination as the outcor	00		0.0	2		0.0	9	
HPV11 0 No. of infections‡ 0.0 Prevalence, % (95% CI)\$ 0.0 Cross-protection types HPV31/33/45 1 No. of infections‡ 1 No. of infections‡ 1 Prevalence, % (95% CI)\$ 0.05 (0.0 to 0.85) 0.2 Abbreviations: HPV, human papilloma virus; NE, not estime *Binary logistic regression modeling was initially conducte unvaccinated individuals, after which the score test <i>P</i> value w score test <i>P</i> value withe, non-Hispanic black, and othe #The logistic propensity model had vaccination as the outcor	0.36 (0.10 to 0.91)		Ι	0.14 (0.01 to 0.66)		Ι	0.54 (0.11 to 1.56)	
No. of infections‡ 0.0 Prevalence, % (95% CI)\$0.1 Cross-protection types HPV31/3345 1 No. of infections‡ 1 No. of infections‡ 1 Prevalence, % (95% CI)\$ 0.05 (0.0 to 0.85) 0.2 Abbreviations: HPV, human papilloma virus; NE, not estime *Binary logistic regression modeling was initially conducte unvaccinated individuals, after which the score test <i>P</i> value w score test <i>P</i> value on the interval of the territe adjusted pri- tripanic, non-Hispanic, white, non-Hispanic black, and othe #The logistic propensity model had vaccination as the outcor			0		NE/.33	0		NE/.33
Prevalence, % (95% Cl)§01. Cross-protection types HPV31/3345 No. of infections‡ No. of infections‡ No. of infections Prevalence, % (95% Cl)§ 0.05 (0.0 to 0.85) 0.2 Abbreviations: HPV, human papilloma virus; NE, not estime *Binary logistic regression modeling was initially conducte unvaccinated individuals, after which the score test <i>P</i> value w score test <i>P</i> value computation, we defined do not derive adjusted pri thilpsanic, non-Hispanic which the vaccination as the outcor if The logistic propensity model had vaccination as the outcor	2 .4	49  /.24	0.0	-		0.0	-	
Cross-protection types HPV31/33/45 No. of infections‡ Prevalence, % (95% Cl)§ 0.05 (0.0 to 0.85) 0.2 Abbreviations: HPV, human papilloma virus; NE, not estima *Binary logistic regression modeling was initially conducte unvaccinated individuals, after which the score test <i>P</i> value v score test <i>P</i> value computation, we did not derive adjusted pri Hispanic, non-Hispanic white, non-Hispanic black, and other iThe logistic propensity model had vaccination as the outcor	0.17 (0.01 to 0.70)		Ι	0.13 (0.0 to 0.74)		Ι	0.21 (0.0 to 1.23)	
HPV31/33/45 1 No. of infections‡ 1 No. of infections‡ 0.05 (0.0 to 0.85) 0.2 Abbreviations: HPV, human papilloma virus; NE, not estime *Binary logistic regression modeling was initially conducte unvaccinated individuals, after which the score test <i>P</i> value v score test <i>P</i> value computation, we did not derive adjusted pri Hispanic, non-Hispanic white, non-Hispanic black, and other †The logistic propensity model had vaccination as the outcor	ġ.	.91¶/.85			NE/.33			.15¶/.085
No. of infections <sup>‡</sup> 1 Prevalence, % (95% CI)\$ 0.05 (0.0 to 0.85) 0.2 Abbreviations: HPV, human papilloma virus; NE, not estime *Binary logistic regression modeling was initially conducte- unvaccinated individuals, after which the score test <i>P</i> value w unvaccinated individuals, after which the score test <i>P</i> value w the score test <i>P</i> value computation, we did not derive adjusted print (Hispanic, non-Hispanic white, non-Hispanic black, and other †The logistic propensity model had vaccination as the outcor				0		0		
Prevalence, % (95% CI)\$ 0.05 (0.0 to 0.85) 0.2: Abbreviations: HPV, human papilloma virus; NE, not estima *Binary logistic regression modeling was initially conducte- unvaccinated individuals, after which the score test <i>P</i> value w invection the computation, we did not derive adjusted print. [Hispanic, non-Hispanic white, non-Hispanic black, and other †The logistic propensity model had vaccination as the outcor	Ð		1	0.0		0.0	5	
Abbreviations: HPV, human papilloma virus; NE, not estima *Binary logistic regression modeling was initially conducter unvaccinated individuals, after which the score test <i>P</i> value w core test <i>P</i> value computation, we did not derive adjusted prin [Hispanic, non-Hispanic white, non-Hispanic black, and other †The logistic propensity model had vaccination as the outcor	0.27 (0.07 to 0.73)		0.07 (0.0 to 1.07)	I		I	0.49 (0.13 to 1.30)	
model were used to modify the poststratified National Health and Nutrition Examination Survey Mobile Exam Center weights of the unvaccinated individuals, which were then used as weights for the comparison of oral HPV prevalence between vaccinated and unvaccinated individuals using a quasi-score test. Sex-specific propensity models included age (linear) and race (Hispanic, non-Hispanic black, and other a Lonweighted vaccination status was pooled in the score test. Pvalue computation, we did not derive adjusted prevalence estimates or measures of association, such as adjusted odds ratios. SVeighted estimates or measures of association, such as adjusted odds ratios. SVeighted estimates are adjusted for accounted as adjusted odds ratios.	nable. ted with oral HPV infectic was computed for the co revalence estimates or m er races), unless noted o ome and gge (linear), sex, h and Nutrition Examinatic widuals using a quasi-scor e test <i>P</i> value computati	on as the ou mparison of a neasures of a neasures of a and race (H and suck y n Survey M re test. Sex- on, we did i	troome and age, se oral HPV prevalence issociation, such as spanic, non-Hispani bbile Exam Center w specific propensity r iot derive adjusted	x, and race as predict between vaccinated adjusted odds ratios. E white, non-Hispanic eights of the unvaccin nodels included age (I orevalence estimates	ors to accoun and unvaccina Binary logistic r black, and othe ated individual inear) and race or measures o	t for the imbalance ir ted individuals. Becat egression models we graces) as the predic s, which were then us (Hispanic, non-Hispa of association, such a	al HPV infection as the outcome and age, sex, and race as predictors to account for the imbalance in confounders between vaccinated and uted for the comparison of oral HPV prevalence between vaccinated and unvaccinated individuals. Because vaccination status was pooled in the settmates or measures of association, such as adjusted odds ratios. Binary logistic regression models were adjusted for age (linear), sex, and race nless noted otherwise. The propersity models into the invaccinated individuals. Because vaccination status was pooled in the nless noted otherwise. The propersity is such as adjusted odds ratios. Binary logistic regression models were adjusted for age (linear), sex, and race le (linear), sex, and race (Hispanic, non-Hispanic black, and other races) as the predictors. Predicted odds from the propensity ion Examination Survey Mobile Exam Center weights of the unvaccinated individuals, which were then used as weights for the comparison of oral g a quasi-score test. Sex-specific propensity models included age (linear) and race (Hispanic, non-Hispanic white, non-Hispanic black, and other lue computation, we did not derive adjusted prevalence estimates or measures of association, such as adjusted odds ratios.	vaccinated and as pooled in thu n), sex, and raco the propensit mparison of ore black, and othe

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