ABSTRACT

Background: Near elimination of cervical cancer in the United States is possible in coming decades, yet inequities will delay this achievement for some populations. We sought to explore the effects of human papillomavirus (HPV) vaccination on disparities in cervical cancer incidence between high- and low-poverty U.S. counties.

Methods: We calibrated a dynamic simulation model of HPV infection to reflect average counties in the highest and lowest quartile of poverty (percent of population below federal poverty level), incorporating data on HPV prevalence, cervical cancer screening, and HPV vaccination. We projected cervical cancer incidence through 2070, estimating absolute and relative disparities in incident cervical cancer for high- versus low-poverty counties, and compared incidence with the near-elimination target (4 cases/100,000 women annually).

Results: We estimated that, on average, low-poverty counties will achieve near-elimination targets 14 years earlier than high-poverty counties (2029 vs. 2043). Absolute disparities by county poverty will decrease, but relative differences are estimated to increase. We estimate 21,694 cumulative excess cervical cancer cases in high-poverty counties over the next 50 years. Increasing HPV vaccine coverage nationally to the Healthy People 2020 goal (80%) would reduce excess cancer cases, but not alter estimated time to reach the near-elimination threshold.

Conclusions: High-poverty U.S. counties will likely be delayed in achieving near-elimination targets for cervical cancer and as a result will experience thousands of potentially preventable cancers.

Impact: Alongside vaccination efforts, it is important to address the role of social determinants and health care access in driving persistent inequities by area poverty.

Introduction

Geographic disparities in cancer are well documented in the United States, with those living in higher poverty areas experiencing higher morbidity and mortality from numerous preventable cancers (1–4). The largest disparities by area poverty are in cancers associated with human papillomavirus (HPV), including cervical, anal, and oropharyngeal cancers (1). Individuals in high-poverty areas are nearly twice as likely to be diagnosed with cervical cancer as those in low-poverty areas (1, 2). Geographic inequities in cervical cancer are complex and attributed to multiple, overlapping risk factors, including prevalence of high-risk HPV types and lower provision of cervical cancer screening to detect precancerous stages of disease (5–7).

In 2006, the introduction of HPV vaccine created an important new opportunity for cancer prevention. The most recently licensed HPV vaccine in the United States protects against seven oncogenic HPV types along with two types that cause approximately 90% of genital warts (8). The potential benefits of HPV vaccine are so promising that the near elimination of cervical cancer is considered an achievable goal in the United States (9–11). Previous simulation models of HPV vaccine coverage nation wide to the Healthy People 2020 goal (80%) would reduce excess cancer cases, but not alter estimated time to achieve near-elimination targets (14, 20). To understand the long-term implications of these patterns. Beyond reducing overall cancer burden, HPV vaccination has the potential to reduce geographic disparities in cancer outcomes by providing accessible prevention that has low out-of-pocket cost for patients (13). Studies examining HPV vaccine uptake by area poverty have suggested that HPV vaccination rates in high-poverty areas may be higher than in low-poverty areas (14–17), but these observations vary by how areas are defined and better longitudinal data are needed, along with models that can explore the long-term implications of these patterns.

To date, studies have modeled the potential impact of HPV vaccination in the United States, but the long-term effects of HPV vaccine have focused on the United States as a whole, potentially missing geographic heterogeneity in HPV prevalence, cervical cancer screening rates, and underlying cervical cancer risk (18–20). To understand the long-term implication of current HPV vaccination patterns on disparities in cervical cancer incidence between high- and low-poverty counties in the United States, we use a stratified dynamic HPV infection model, incorporating data on vaccination, screening, and HPV prevalence by county poverty.

Materials and Methods

HPV transmission model

We adapted a dynamic HPV transmission model to reflect composite high- and low-poverty U.S. counties (21). Briefly, the compartmental model simulates the transmission of HPV through sexual partnerships, allowing for both direct and indirect effects of...
vaccination (i.e., herd protection), as well as simulating progression of HPV to cervical cancer (Supplementary Fig. S1). Individuals are born, age in 1-year increments, and die at age-varying mortality rates. Individuals begin the model susceptible and may acquire an HPV infection based on age-specific sexual activity, prevalence of HPV among opposite-sex sexual partners (as a model simplification, we do not include same-sex sexual partnerships), and transmissibility of HPV. We collapse HPV types to separately describe high-risk types protected by HPV vaccine (HPV 16, 18, 31, 33, 45, 52, 58) and high-risk types not protected by current vaccines (HPV 35, 39, 51, 56, 59, 68). Women with a high-risk HPV infection can spontaneously clear infection, progress to precancerous cervical lesions, have lesions be identified and treated through routine cervical cancer screening, or have lesions progress to incident cervical cancer, at which point they exit the model. Our model was constructed using R (version 4.0.3).

Model inputs and target data by poverty quartile

Using the 2011–2015 American Community Survey, we classified U.S. counties into quartiles based on percent of residents living below 100% of the Federal Poverty Level (22). The lowest poverty quartile represents the approximately 84 million individuals living in counties where less than 11.9% of the population lives in poverty (Supplementary Table S1) while the highest poverty quartile represents approximately 45 million individuals living in counties with greater than 20.3% of the population living in poverty. We estimated age-specific all-cause mortality by poverty quartile using CDC Wonder (23).

We obtained data on characteristics by poverty quartiles from three large nationally representative surveys; the National Health Interview Survey (NHIS), the National Health and Nutrition Examination Survey (NHANES), and the National Immunization Survey–Teen (NIS-Teen) through the National Center for Health Statistics which allowed for use of county identifiers not available in the public data. For each survey, respondents’ county of residence was matched to poverty quartile from the American Community Survey. Quartile-matched data were analyzed at a secure Federal Research Data Center. As the study team did not have direct access to the individual county identifiers (only the matched quartile data), the study was determined to be exempt by the University of North Carolina Institutional Review Board.

Input: cervical cancer screening

We obtained data on cervical cancer screening from the NHIS (2013–2015), including approximately 3,600 women living in high-poverty counties each year and 6,600 women living in low-poverty counties in each year of data. We analyzed the proportion of women up to age 30 on cervical cancer screening, by poverty quartile, using complex survey weights to account for sampling design. We assessed differences using an independent sample t test with an alpha value of 0.05. All survey data were analyzed using Stata 16.

NHIS and other national U.S. data sources have not shown changes over time in the proportion of women receiving timely cervical cancer screening (24–26); however, U.S. recommendations for screening modality have shifted to recommend coadministering Pap smears with more sensitive HPV DNA testing (27, 28). Our base case assumed the adoption of HPV DNA screening would increase from 0% in 2005 to 70% of tests by 2020 in both high- and low-poverty counties (Supplementary Fig. S2). Because HPV DNA testing adoption varies geographically (29); sensitivity analyses modeled delayed adoption of HPV DNA testing in high-poverty counties (taking until 2030 for these counties to reach 70% use of HPV DNA testing). We estimate 26% of women in a high-poverty setting and 15% of women in a low-poverty setting fail to return for subsequent follow-up from an abnormal screening test within 1 year (30–32).

Input: HPV vaccination

HPV vaccination was obtained from the NIS-Teen (2008–2015), including provider-verified vaccination records from 5,000 adolescents living in high-poverty and 9,000 adolescents living in low-poverty counties each year. Data were available for girls starting in 2008 and boys starting in 2011. To facilitate comparison across settings, we present data on prevalent HPV vaccine coverage—percent of those age 11–17 with at least one dose and percent with all recommended doses by year, sex, and county poverty quartile. However, the inputs to our model are defined from these same data in a more specific way, as annual probability of incident HPV vaccine initiation (receiving first dose during the year) and completion (receiving final dose during the year) by sex, age group (11–12 vs. 13–17), and county poverty quartile. Model inputs for years before 2008 and after 2015 were projected, assuming stable uptake after 2020. (Supplementary Fig. S3).

Calibration and validation targets: HPV prevalence

We obtained HPV prevalence data from NHANES, a nationally sampled survey that combines personal interviews with physical examination and laboratory testing data. We used 2003–2006 data for model calibration and 2011–2014 data for validation of model output, we had approximately 500 high-poverty and 900 low-poverty participants in each 4-year combined cycle. Female NHANES participants ages 18–60 were tested for HPV DNA using a self-collected vaginal swab (33). We report prevalence of vaccine-protected high-risk types and nonprotected high-risk HPV types and prevalence of any high-risk type by age.

Model calibration and validation

The underlying causes of inequities in HPV prevalence and cervical cancer incidence are complex and likely mediated by differences in health care access, health behaviors, social determinants of health, and underlining health status that may affect HPV acquisition and the natural history pathway. We used model calibration to approximate differences in model parameters that could not be estimated directly from the literature, including sexual behavior and natural history of infection. We calibrated parameters that could plausibly vary by county poverty, using a single base model and varying parameter values by ±25% using a Latin hypercube sampling approach (34, 35). We compared 10,000 possible parameter sets using a log-likelihood estimate against setting-specific calibration targets of HPV prevalence, separately by HPV type (all women ages 18–60) and by age (any high-risk type). The 50 best-fitting parameter sets for each quartile were then used to estimate prostate cancer cervical incidence, with a directed search algorithm identifying for each set an HPV progression multiplier which best matched 2006 age-adjusted cancer incidence by poverty quartile [5.9 and 8.4 cases per 100,000 for low- and high-poverty counties, respectively (36)].

The result of this process was 50 paired combinations of parameters describing possible variation in unobservable characteristics leading to the observable differences in high-poverty and low-poverty counties. We assessed validity of these parameter sets by comparing the model estimate of HPV prevalence in 2014 to observed 2011–2014 NHANES prevalence estimates by HPV type and for all high-risk HPV types by age. As there is no single standard for evaluating validity to external data (37), we report the percent of model estimates that fall within two and within three SDs of their corresponding observed value.
Model projections

Fully calibrated models were run starting in 2006 incorporating HPV vaccination rates by county poverty quartile, repeating this process for each calibration set. We projected outcomes through 2070, as this covers the period of highest risk for cervical cancer among cohorts for which high-quality vaccination data are available. We compared a scenario assuming stable HPV vaccine rates after 2020 to a scenario with a 2020 increase to 80% HPV vaccine coverage. Our primary model outcome is projected annual age-adjusted cervical cancer incidence. We also report the year in which average high- and low-poverty counties would be projected to achieve targets for “near-elimination” (annual incidence below 4 per 100,000). We measure disparities through comparing paired combinations of our high- and low-poverty models, producing 50 estimates of (i) the absolute number of excess cervical cancer cases (risk difference) and (ii) the relative risk of cervical cancer in high versus low-poverty counties. We estimated total excess cervical cancer cases across the total female population of all high-poverty U.S. counties (~24 million women) if annual incidence were matched to that of low-poverty counties.

Results

Empirical data by county poverty quartile

Prevalence of high-risk HPV types covered by HPV vaccine did not significantly vary, at 12.9% in high-poverty and 10.2% in low-poverty counties in the prevaccine era (Fig. 1A). Prevalence of one or more high-risk HPV types not protected against by current HPV vaccines was higher in high-poverty counties than low-poverty counties (18.3% vs. 8.9%, \( P < 0.01 \)). Examining within age strata, prevalence of any high-risk HPV type was higher in high poverty, compared with low-poverty counties, for those 45–54 (Fig. 1B); other age groups showed the same direction of findings but were not statistically significant. Women ages 35–44 and 45–55 in high-poverty counties were less likely to be up to date on cervical cancer screening than for the same age groups in low-poverty counties (Supplementary Fig. S4).

Figure 1.
Prevalence of high-risk HPV by county poverty.
*Different by Wald test comparing high- versus low-poverty counties (\( P < 0.05 \)); Error bars represent 95% confidence interval; Data from National Health and Nutrition Examination Survey 2003–2006. A, Women ages 18–60 by HPV type. B, Prevalence of all high-risk (HR) HPV types by age group.
High- and low-poverty counties showed similar patterns of HPV vaccine uptake across all years (Fig. 2). By 2015, 69% of 11 to 17 years old girls and 51% of 11 to 17 years old boys in high-poverty counties had initiated HPV vaccination while 45% and 32% of girls and boys had completed the series. This was not different from low-poverty counties (initiation 63% and 53% for girls and boys, respectively; completion was 48% and 33%).

Model calibration and validation

The 50 best-fitting calibration sets for our high- and low-poverty models included values from across the search ranges (Supplementary Table S2), suggesting these sets represent a diversity of underlying drivers of disparities which each produce acceptable fits to prevaccine HPV prevalence data (Supplementary Fig. S5). Our validation against 2011–2014 NHANES data showed that all parameter sets resulted in reasonable fits to empirical data by both age and HPV type. 91% of estimates were within two SDs of the corresponding NHANES value and 97% of estimates were within three SDs (Supplementary Fig. S6).

Model projections

Projecting model estimates forward to 2070, both low- and high-poverty county models see substantial reductions in cervical cancer incidence (Fig. 3). In our low-poverty model, annual age-adjusted incidence rates were 5.9 per 100,000 in 2006 and were projected to fall to 0.7 per 100,000 by 2070, with parameter sets ranging from 0.5 to 1.2. In our high-poverty model, annual incidence declined from 8.4 per 100,000 in 2006 to 1.7 per 100,000 in 2070 (range: 1.1 to 2.5). We estimated low-poverty counties will, on average, achieve the near-elimination target by 2030 (range: 2027–2032) while, on average, high-poverty counties are not projected to reach this goal for another 14 years (2044 (2041–2048). Over the 50 years from 2020–2070, we estimate a total of 21,604 excess cervical cancer cases in high-poverty counties, relative to the burden if incidence were identical to that of low-poverty counties.

We estimate 2.5 excess incident cervical cancer cases per 100,000 women for average high-poverty counties relative to average low-poverty counties. We estimated this absolute disparity would shrink to

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**Figure 2.** HPV vaccine initiation and completion by poverty quartile. *Different by Wald test comparing high- versus low-poverty counties (P < 0.05); error bars show 95% confidence intervals. Data from provider-verified records of adolescents 11–17 years of age in 2008–2015 NIS-Teen. Initiation indicates receipt of at least one dose of HPV vaccine, and completion indicates receipt of all recommended doses.
1.0 cases per 100,000 by 2070 (Fig. 4A). The relative disparity is expected to increase (Fig. 4B), with women in our high-poverty county at 1.4 times the risk of incident cancer versus those in our low-poverty county in 2006 but 2.5 times the risk by 2070. These findings were consistent across paired calibration sets, with no comparisons showing a total disappearance of disparities by poverty quartile.

Assuming our high- and low-poverty county each achieved target thresholds of 80% vaccine completion in 2020, near-elimination timelines improved by less than a year (Table 1) and we found a small improvement in absolutely and relative disparities by 2070 compared with current practice. Assuming delayed adoption of HPV DNA testing in the high-poverty county, disparities initially widened in all analyses, but by 2070 absolute and relative disparities were similar to those in the base case. Alternative assumptions about protection from partial series completion had only small effects, with a small increase in both absolute and relative disparities if protection was lowered (50% of full series) and a slight improvement if fully protective (equivalent to completing the full series).

Discussion

Our dynamic HPV transmission model suggests current uptake of HPV vaccination is likely to dramatically reduce cervical cancer burden in both low- and high-poverty U.S. counties. Absolute disparities (i.e., number of excess cancers per 100,000) will decline as overall burden decreases in both high- and low-poverty counties. However, it is likely that relative differences will remain, with incident cervical cancer burden in high-poverty counties 1.5 to 3 times that of low-poverty counties. We found these conclusions robust to changes in model inputs and across calibrated parameter sets.

Differential access to advancements in cancer treatment and prevention often reduces overall cancer burden but widens inequity (38). Disparate access has not been generally seen in HPV vaccination, where uptake is similar or higher among multiple traditionally underserved groups (39, 40). While we did not find higher average uptake of HPV vaccine in high-poverty areas, as some studies have reported previously (14, 16), we found comparable uptake in high- and low-poverty counties, which is nonetheless promising. Differences in conclusions across studies are likely due to differences in the definition of geographies (e.g., states, counties, census tracts, and zip codes) as well as the socioeconomic variables selected for comparisons (17). A geospatial approach may help prioritize areas where improved HPV vaccination could have the highest benefits for equity, but as uptake remains below coverage goals in nearly all areas of the United States, broad approaches to improving HPV vaccine uptake are still urgently needed (41).

In addition, we found higher prevalence of high-risk HPV types that are not vaccine protected in high-poverty counties. Histological studies attribute only a small portion of invasive cervical cancers to these types, but differences in type distribution and co-infection may further disadvantage those already experiencing lower access to screening and preventive care (42). Better characterizing cancer attribution by HPV type, particularly among high-burden populations, is an important step for understanding disparities.

Our findings are comparable with previous modeling and empirical studies evaluating likely changes for the full U.S. population. Two microsimulation models projected near elimination in the United States by 2038 and 2046 (12). While studies on area poverty are limited, studies focusing on populations of color echo the pattern of our findings. For example, Black women have higher cervical cancer incidence (3) and higher HPV prevalence—particularly for high-risk types not protected by HPV vaccine (43), but higher HPV vaccine initiation (40). A study modeling the potential impact of HPV vaccine on racial disparities found that current HPV vaccination patterns were likely to decrease, but not eliminate, cervical cancer disparities by race (44).

As high-poverty counties in the United States have, on average, higher proportions of Black, American Indian, and Latinx populations than low-poverty counties and are more likely to be rural, the disparities we explore here reflect many of the same structural inequities that lead to racial disparities and rural/urban disparities (45). This includes lower access to cervical cancer screening, an important method of secondary prevention (46). Those living in high-poverty settings are also more likely to experience certain risk factors for HPV cancers, including higher smoking rates, higher parity, and co-occurring sexually transmitted infection (47–49).

Identifying multiple calibration sets allowed us to explore results across different combinations of uncertain variables. Our conclusions were largely robust to differences across calibrated parameter sets, but a better understanding of the underlying causes of observed disparities could be important for informing specific policy. We found that while increasing HPV vaccination to the target of 80%
U.S.-wide would reduce total number of incidence cervical cancers in the next 50 years, it will not improve the timeline for a high-poverty county to reach near elimination and is unlikely to completely eliminate disparities. This suggests additional consideration of social determinants and other prevention strategies are important for understanding and reducing cancer disparities by area poverty in the near term and long term.

We did not construct our model to evaluate cervical cancer screening in detail and thus did not incorporate fine-grained detail on screening or surveillance patterns. Future work should examine Figure 4.

**Table 1. Sensitivity analysis.**

<table>
<thead>
<tr>
<th>Partial series efficacy (relative to full series)</th>
<th>80% HPV vaccine coverage</th>
<th>Low HPV DNA uptake in high-poverty county*</th>
</tr>
</thead>
<tbody>
<tr>
<td>Median [min–max]</td>
<td>Base case</td>
<td>100%</td>
</tr>
<tr>
<td>Absolute disparity: 2070 (risk difference)</td>
<td>1.0 [0.2–1.9]</td>
<td>0.8 [0.2–1.8]</td>
</tr>
<tr>
<td>Relative disparity: 2070 (risk ratio)</td>
<td>2.5 [1.2–4.7]</td>
<td>1.9 [1.2–4.5]</td>
</tr>
</tbody>
</table>

*High-poverty county reaches 70% uptake of HPV DNA testing by 2030.
**Estimates scaled to total population of all high-poverty counties.
cervical cancer screening by county poverty in greater detail to understand whether higher rates of screening, diagnosis, and treatment could further reduce disparities. This is particularly important given evolving recommendations regarding screening frequency and modality. The shift to HPV testing, including new self-collection methods, may improve access to screening among traditionally underserved populations, including those living in poverty (50–52). However, disruptions in preventive care resulting from the COVID-19 pandemic have produced delays or omissions of both cervical cancer screening and HPV vaccination that are likely to result in thousands of excess cervical cancers nationwide (53–55). While no data are yet available on whether these delays differ by county poverty, it is likely that the disproportionate impact of COVID-19 along socioeconomic gradients may set back progress made by HPV vaccination and cervical cancer screening innovation. To continue meaningful gains in cancer prevention equity, it will be important to understand and address multilevel barriers to HPV vaccination, screening, follow-up, and treatment of precancerous diseases among traditionally underserved groups (56).

We note several limitations and strengths of our modeling approach. As the field’s understanding of cervical carcinogenesis continues to grow, granularity and accuracy of simulation models will continue to improve. Using a compartmental model limited our ability to incorporate concurrent infections with multiple genotypes or genotype replacement, potentially important sources of uncertainty in modeling vaccine impact (57, 58). We include only heterosexual partnerships, a simplification that reduces heterogeneity in mixing and excludes some populations with high burden of HPV disease (59, 60). Furthermore, recent work has proposed a new framework that moves away from histology-based classification (CIN1, CIN2,3) to a more parsimonious model that can be explicitly informed by time since HPV appearance, HPV genotype, and other disease biomarkers (61). As these approaches continue to evolve, we hope future work will be able to adapt them to better understand structural, behavioral, and biological causes of existing cervical cancer disparities.

While the field’s understanding of the natural history in cervical cancer is evolving, it is relatively well described among HPV cancers. Poverty disparities exist for all HPV cancer sites, yet the drivers of these differences are unclear (1). Future work should explore the implication of current vaccination patterns in other HPV cancer sites, including oropharyngeal cancer which has recently surpassed cervical cancer in annual incidence in the United States (62). Finally, we modeled a composite high- and low-poverty county using average data that may obscure heterogeneity between counties and we do not model migration between counties. As vaccination rates are similar in high- and low-poverty settings, migration would likely only obscure heterogeneity by county and we do not model migration between counties. As vaccination rates are similar in high- and low-poverty settings, migration would likely only influence our primary model conclusions if differential by vaccination status, but better data on migration between high- and low-poverty areas would also be valuable for improving our understanding of current disparities and assessing other strategies to improve equity.

Our study is strengthened by the use of quartile-matched data from multiple large national surveys to inform model inputs, as well as validation of our model estimates to empirical data and comparability with other cervical modeling studies (12, 19). Our study takes a nuanced approach to modeling cancer disparities through incorporating novel empirical data on HPV burden and HPV vaccination for both low- and high-poverty U.S. counties and we hope it will help to guide priority setting for national and regional cancer prevention efforts.

In addition to having potential for cancer prevention, HPV vaccine offers an unprecedented opportunity to reduce the large and persistent disparities in HPV cancer between high- and low-poverty counties in the United States. Current vaccination rates are projected to reduce, but not eliminate, the higher incidence of HPV cancers in high-poverty areas relative to low-poverty areas and may increase relative disparities. HPV vaccination alone is unlikely to achieve equity in HPV cancer in the near term; therefore, policymakers and advocates should continue broad efforts to increase HPV vaccination alongside more targeted efforts to improve social determinants of health potentially including improving access to preventive, screening, and diagnostic care for communities with disproportionate burden from HPV and HPV cancers.

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Authors’ Contributions
J.C. Spencer: Conceptualization, formal analysis, methodology, writing–original draft. N.T. Brewer: Conceptualization, writing–review and editing. T. Coyne-Beasley: Conceptualization, writing–review and editing. J.G. Trogdon: Conceptualization, writing–review and editing. M. Weinberger: Conceptualization, writing–review and editing. S.B. Wheeler: Conceptualization, methodology, writing–review and editing.

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