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A Comparison of the Natural History of HPV Infection and Cervical Abnormalities among HIV-positive and HIV-negative Women in Senegal, Africa

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Abstract

Background—There is evidence of an interaction between human immunodeficiency virus (HIV) and human papillomavirus (HPV) resulting in increased HPV-associated morbidity and cancer mortality among HIV-positive women. This study aims to determine how the natural history of cervical HPV infection differs by HIV status.

Methods—A total of 1,320 women (47% were positive for HIV-1 and/or HIV-2) were followed for an average of two years in Senegal, West Africa between 1994 and 2010. Cytology (with a sub-sample of histology) and HPV DNA testing were performed at approximately 4-month intervals yielding data from over 7,900 clinic visits. Competing risk modeling was used to estimate rates for transitioning between three clinically relevant natural history stages: Normal, HPV, and HSIL (high-grade squamous intraepithelial lesions). Among HIV-positive women, exploratory univariate analyses were conducted examining the impact of HPV type, infection with multiple HPV types, HIV type, CD4+ count, and age.

Results—HIV-positive women had higher rates of progression and lower rates of regression compared to HIV-negative women (i.e. adverse transitions). HIV-positive women had a 2.55 (95% CI: 1.69–3.86; $P < 0.0001$) times higher rate of progression from HPV to HSIL than HIV-negative

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women (with 24-month absolute risks of 0.18 and 0.07, respectively). Among HIV-positive women, HPV-16/18 infection and CD4+ count <200/mm³ were associated with adverse transitions.

Conclusions—Adverse HIV effects persist throughout HPV natural history stages.

Impact—In the limited-resource setting of sub-Saharan Africa where cervical cancer screening is not widely available, the high-risk population of HIV-positive women may be ideal for targeted screening.

Keywords

Human papillomavirus; human immunodeficiency virus; cervical cancer natural history; transition probabilities; competing risk

Introduction

Human papillomavirus (HPV) is the necessary cause of squamous cervical cancer (1), is highly transmissible, and generally acquired closely after sexual debut (2, 3). Persistent infection can lead to the development of pre-cancerous lesions which, in the absence of treatment or an effective immune response, can progress to cancer. Approximately 40 genotypes infect the genital tract and are classified based on oncogenic potential with high-risk types 16 and 18 accounting for roughly 70% of cervical cancer (4). There is evidence of an interaction between human immunodeficiency virus (HIV) and HPV, with HIV-positive women at an increased risk for HPV detection, pre-cancerous lesions, and cervical cancer compared to HIV-negative women (5–13). Studies have shown that HIV-positive women are 1.5 to 8 times more likely to have cervical cancer than HIV-negative women (9, 11, 14–17).

Despite numerous studies describing the increased burden of HPV and cervical disease among HIV-positive women, few have longitudinally examined the natural history within a single study population and provided direct comparisons to HIV-negative women. Thus, estimates for the probability of transitioning between each natural history stage (i.e. progression and regression) are limited, as well as our understanding of the point at which the natural history of HPV diverges for HIV-positive and HIV-negative women. We estimate and compare the probability of transitioning between three clinically relevant natural history stages (Normal, HPV, and high grade pre-cancerous lesions (HSIL)) for HIV-positive and HIV-negative women using data from multiple cohort studies conducted in Senegal, West Africa.

Materials & Methods

Sample

Data from six studies conducted from 1994 to 2010 in Senegal were used for the present analysis (Table 1). Protocols have been described elsewhere (9, 10, 18–25). Women age 15 at outpatient clinics were screened for participation in longitudinal research with HIV testing at baseline, as well as cervical cytology and HPV DNA sampling roughly every four months for at least two years. Recruitment occurred at two infectious disease, two family planning,

and two sexually transmitted disease clinics in or around Dakar. Studies were approved by the University of Washington and Senegalese Human Subjects Review Boards with informed consent obtained from each subject. Data were de-identified and shared variables across the studies were pooled for analysis.

HIV Serology and Lymphocyte Testing

Blood samples were tested for HIV-1 and HIV-2 antibodies using an enzyme-linked immunosorbent assay (ELISA; Genetic Systems, Seattle, WA, USA), a microwell plate enzyme immunoassay (HIV 1/2 EIA; Sanofi Diagnostics Pasteur, Paris, France), or by rapid HIV testing (Determine Alere, Inc, Jena, Germany). HIV-1 and HIV-2 infections were distinguished by a peptide-based assay, although the assay varied by study (Genie II, Genetic Systems; Multispot, Sanofi Diagnostics Pasteur; or Immunocomb II, bispot, Organics, Yavne, Israel). For HIV-positive women, peripheral blood was analyzed with a FACSCount analyzer (Becton Dickinson Biosciences, San Jose, CA, USA) to determine number of CD4+ cells per microliter.

Cytologic and Histologic Testing

Conventional Pap smears were used and evaluated in Dakar until 1998; thereafter, the Thin Prep monolayer cell preparation system (Cytoc Corp., Boxborough, MA) was used and evaluated by a cytopathologist in Seattle, USA. Results were classified according to the Bethesda system (atypical squamous cells of undetermined significance - ASCUS, low-grade squamous intraepithelial lesion - LSIL, high-grade squamous intraepithelial lesions - HSIL) (26). All slides obtained prior to 1998, those classified as LSIL or worse, and a random subset of negative slides, were re-read in Seattle.

Protocols for all six parent studies called for colposcopically-directed biopsies and treatment for women with evidence of HSIL or invasive cervical cancer (ICC). Biopsy/treatment participation was low in earlier studies, in part, due to delays resulting from samples being sent to and re-read in Seattle. Representative hemotoxylin-eosin-stained slides were prepared from paraffin-embedded biopsy specimens and reviewed by a blinded cytopathologist. World Health Organization (WHO) pathology criteria were used to classify specimens (19).

HPV DNA Detection and Typing

Polymerase chain reactions (PCRs) assays for detection of HPV DNA were performed. Lab methods evolved over time with expansion of type specific probes (Table 1). Testing was initially performed by use of HPV L1 consensus primers, HPV type-specific oligonucleotide probes, and a generic probe (27). This method yielded type-specific identification of HPV 6/11, 16, 18, 31/33/35/39, 45/56, and 51/52. With new probes available, in 1998 HPV detection and typing analyses were performed via a PCR-based reverse-line strip test method (Roche Molecular Systems, Alameda, CA) with probes for types 6, 11, 16, 18, 26, 31, 33, 35, 39, 40, 42, 45, 51, 52, 53, 54, 55, 56, 57, 58, 59, 66, 68, 73, 82, 83, and 84 (28). In 2000, a Luminex-based testing approach was adopted with additional probes for 61, 62, 64, 67, 69, 70, 71, 72, 81, IS39, and CP6108 (29).

Natural History Stage Classification

In combination with HPV DNA results, cervical histology was used when available, with cytology in all other cases, to classify women at each visit as: *Normal* (HPV-negative), *HPV* (HPV-positive with one or more type in the absence of HSIL), or *HSIL* (HPV-positive with one or more type in the presence of HSIL). HSIL was defined based on the presence of HSIL, CIN2, CIN3, or CIS. Due to a demonstrated lack of reproducibility for cytologic and histologic interpretations of low-grade lesions (i.e. ASCUS, LSIL, and CIN1), these results were not used to inform stage classifications (30). Transitions to ICC were not examined due to small numbers and possible HSIL treatment effects.

Statistical Analyses

Competing risk modeling was used to estimate cumulative incidence functions and the probability of a specific transition over time (31). Follow-up time was set as the midpoint between visits (32). At each visit, natural history stage was classified as either incident (i.e. the stage differed from the preceding visit) or prevalent (i.e. the baseline visit or the stage was the same as the preceding visit). As HPV can be transient, restricting analyses to women with incident classification to eliminate left-censoring may produce bias. For example, eliminating women who are classified as *Normal* throughout follow-up (prevalent classification), leads to overestimation of the probability of transitioning from *Normal* to *HPV* as only women with prior HPV detection who become HPV-negative during follow-up contribute to incident *Normal*. Thus, to bound results, analyses were calculated separately for prevalent and incident natural history stages.

The following exploratory analyses were conducted on covariates of interest: age (< 25 vs. >25), CD4+ count (<200/mm³ vs. ≥ 200/mm³), HPV type (16/18 vs. all others), HIV type (1 vs. 2), and concurrent infection with multiple HPV types (>1 vs. 1 type). These analyses were restricted to transitions from *Normal* and *HPV* due to limited numbers for *HSIL*. For HPV-16/18 type specific analyses, women were classified based on the presence of either one or both of these types (33). Women dually infected with HIV-1 and HIV-2 were removed from HIV type specific analyses due to small numbers (n=52).

Gray's method was used to test for statistical differences in cumulative incidence functions (34). Robust variances were used when multiple observations from the same woman were included when estimating a transition. Schoenfeld residuals were examined to assess the proportional hazards assumption. Analyses were conducted using SAS 9.4 software (SAS Institute, Cary, NC).

Results

Demographics (Table 2)

Of 1,320 women, the majority were in their thirties, Muslim, with no more than a primary school education. Forty-percent of married women were in a polygynous relationship. Forty-seven percent of women were HIV-positive, of which roughly two-thirds were infected with HIV-1. Antiretroviral therapy (ART) was reported by 31% of HIV-positive women. Twenty-five percent were classified as having AIDS. Median CD4+ count at baseline was 406 cells/

μL. HIV-positive women had a lower level of education, and were less likely to be currently married or report using contraception than HIV-negative women. HIV-positive women were more likely to be commercial sex workers (CSWs). Most women were followed for over two years (median = 796 days, IQR: 377–1,224 days) with clinic visits approximately every four months. At baseline, roughly 36%, 59%, and 5% of women were classified as *Normal*, *HPV*, and *HSIL*, respectively.

Progression and Regression (Table 3)

Of 1,079 cases classified as *Normal*, 56.3% were incident. HIV-positive *Normal* women had a 1.58 times higher rate of *HPV* detection than HIV-negative women (95% CI: 1.32–1.89). The 24-month predicted probability of *HPV* detection for HIV-positive women was 0.66 (95% CI: 0.60–0.73) and 0.82 (95% CI: 0.78–0.87) for prevalent and incident classification of *Normal*, respectively, compared to 0.50 (95% CI: 0.44–0.56) and 0.66 (95% CI: 0.60–0.73) for HIV-negative women (Figure 1A). Women identified as incident *Normal* had a 1.58 times higher rate of *HPV* detection than prevalent cases (95% CI: 1.33–1.88). HIV-positive *Normal* women also had a 1.53 times higher rate of progression to *HSIL* than HIV-negative women, although this association was not significant (95% CI: 0.73–3.21). The 24-month predicted probability of *HSIL* for HIV-positive women was 0.04 (95% CI: 0.02–0.08) and 0.07 (95% CI: 0.04–0.12) for prevalent and incident classification of *Normal*, respectively, compared to 0.03 (95% CI: 0.01–0.06) and 0.04 (95% CI: 0.02–0.08) for HIV-negative women (Figure 1B). Women identified as incident *Normal* had an increased rate of progression to *HSIL* than prevalent cases, although this was not significant.

Of 1,265 cases of *HPV* detection, 39.4% were incident. HIV-positive women had a 0.46 times lower rate of regression from *HPV* to *Normal* than HIV-negative women (95% CI: 0.39–0.54). The 24-month predicted probability of regression to *Normal* for HIV-positive women was 0.56 (95% CI: 0.50–0.62) and 0.78 (95% CI: 0.74–0.83) for prevalent and incident *HPV* detection, respectively, compared to 0.83 (95% CI: 0.80–0.87) and 0.96 (95% CI: 0.94–0.99) for HIV-negative women (Figure 1C). Women with incident *HPV* detection had a 1.87 times higher rate of regression to *Normal* than women with prevalent detection (95% CI: 1.60–2.18). HIV-positive women with *HPV* had a 2.55 times higher rate of progression to *HSIL* than HIV-negative women (95% CI: 1.69–3.86). The 24-month predicted probability of progression to *HSIL* for HIV-positive women was 0.18 (95% CI: 0.13–0.23) and 0.20 (95% CI: 0.16–0.25) for incident and prevalent *HPV* detection, respectively, compared to 0.07 (95% CI: 0.05–0.11) and 0.08 (95% CI: 0.06–0.12) for HIV-negative women (Figure 1D). Those with prevalent *HPV* detection had a slightly increased rate of progression to *HSIL* than incident cases, although this was not significant.

Of 179 cases of *HSIL*, 60.9% were incident. HIV-positive women had a 0.57 times lower rate of regression from *HSIL* to *Normal* than HIV-negative women (95% CI: 0.26–1.29). The 12-month predicted probability of regression to *Normal* for HIV-positive women was 0.09 (95% CI: 0.04–0.19) and 0.14 (95% CI: 0.08–0.24) for prevalent and incident *HSIL*, respectively, compared to 0.16 (95% CI: 0.07–0.34) and 0.23 (95% CI: 0.14–0.39) for HIV-negative women (Figure 1E). Women with incident *HSIL* had a 1.57 times higher rate of regression to *Normal* than women with prevalent classification, although this association

was not significant. HIV-positive and HIV-negative women with *HSIL* had a similar rate of regression to *HPV* (relative rate = 1.06, 95% CI: 0.71–1.59). For HIV-positive women the 12-month predicted probability of regression to *HPV* ranged from 0.70 (95% CI: 0.60–0.79) and 0.68 (95% CI: 0.56–0.84) for incident and prevalent *HSIL*, respectively, compared to 0.67 (95% CI: 0.52–0.86) and 0.67 (95% CI: 0.54–0.81) for HIV-negative women (Figure 1F). The rate of regression to *HPV* was similar for both prevalent and incident cases of *HSIL*.

Potential Effect Modifiers among HIV-positive Women (Table 4)

HIV-positive women with CD4+ counts <200 had a 2.86 times higher rate of transitioning from *Normal* to *HSIL* than women with higher CD4+ counts (95% CI: 0.95–8.58). Low CD4+ count was also associated with a higher rate of progression from *HPV* to *HSIL* (1.83, 95% CI: 1.07–3.14), and a lower rate of regression from *HPV* to *Normal* (0.57, 95% CI: 0.39–0.83).

HIV-positive *Normal* women with HPV-16/18 had a 4.62 times higher rate of progression to *HSIL* than those infected with other HPV types (95% CI: 1.10–19.42). Similarly, those with HPV-16/18 infection had a higher rate of progressing from *HPV* to *HSIL* (2.20, 95% CI: 1.34–3.62) and a lower rate of regression to *Normal* (0.35, 95% CI: 0.23–0.54) compared to those infected with other HPV types. Rates for acquisition of HPV (*Normal* to *HPV*) were similar across HPV type sub-groupings.

HIV-positive women with multiple *HPV* types had a higher rate of progression to *HSIL* (2.33, 95% CI: 1.40–3.88) than those with a single HPV type. HIV-positive women with multiple HPV types also had a 0.34 times lower rate of regression from *HPV* to *Normal* than women with a single HPV type (95% CI: 0.25–0.47). In contrast, progression rates from *Normal* to *HPV* and *Normal* to *HSIL* were similar for those who acquired multiple HPV types and those who acquired a single type.

Among *Normal* HIV-positive women, those age ≤ 25 years had a 1.56 times higher rate of incident *HPV* detection than those >25 (95% CI: 1.14–2.14). Age associations for the other transitions examined were non-significant.

Among *Normal* HIV-positive women, those with HIV-1 had a 1.85 times higher rate of incident *HPV* detection than those with HIV-2 (95% CI: 1.38–2.49). Similarly, women with HIV-1 had a 1.69 times higher rate of progression from *HPV* to *HSIL* compared to women with HIV-2 (95% CI: 0.94–3.03). Women with HIV-1 also had a lower rate of regression from *HPV* to *Normal* than those with HIV-2 (0.83, 95% CI: 0.64–1.08), although this association was not statistically significant. In contrast, those with HIV-1 had a lower rate of progression from *Normal* to *HSIL* (0.48, 95% CI: 0.19–1.21). Mutual adjustment for baseline age and CD4+ count (as surrogate measures of duration and severity of HIV infection), did not affect these findings.

Discussion

As HIV-positive women are at an increased risk of cervical cancer, understanding the distinct natural history within this population is essential for informing targeted prevention efforts. We found that HIV-positive women had higher rates of incident HPV detection and progression to HSIL, as well as lower rates of regression from HSIL and HPV infection when compared to HIV-negative women. The most notable difference between HIV-positive and HIV-negative women was the more than doubled rate of progression from *HPV* to *HSIL* (although absolute risk remained low). True transition probabilities likely lie between the incident and prevalent curves as a function of age and sexual activity.

While analyses of potential effect modifiers should be interpreted with caution due to limited sample sizes, overall results suggest that HIV-positive women with baseline CD4+ counts <200 (cells/ μ L) or infection with HPV-16/18 had higher rates of progression and lower rates of regression. Consistent with other research (35, 36), this study found increased incident detection of multiple HPV types in HIV-positive women compared to HIV-negative women, but also demonstrated that infection with multiple types increased progression and decreased regression rates from *HPV*. This is particularly relevant as >1 type was present in 57% of visits in which HIV-positive women had HPV detection, in comparison to 28% among HIV-negative women.

Evidence regarding the role of HIV type on the natural history of HPV is conflicting (9, 10, 24, 37–41), with some research indicating that HIV-2 is more strongly associated with HPV-related disease than HIV-1. In contrast, the present analysis found HIV-1 to be more strongly associated with adverse transitions than HIV-2 (with the notable exception of a lower probability of transitioning from *Normal* to *HSIL*). HIV-1 has a shorter incubation period, higher transmissibility, and more rapid development of immunosuppression compared to HIV-2 (22, 42). Thus, it is biologically plausible that women with HIV-1 are at greater risk due to more severe immunosuppression. However, longer survival among women with HIV-2 (43) may result in higher lifetime risk due to extended time to develop cervical cancer.

This study has limitations. The present analysis focused on three stages of natural history: *Normal* (HPV-negative), *HPV* (HPV-positive with one or more type and the absence of HSIL), or *HSIL* (HPV-positive with one or more type and the presence of HSIL). If a woman went from HPV-positive with type 16 to HPV-positive with type 18 in consecutive visits this was classified as a continuation of the *HPV* stage. This situation was unusual as many women had multiple HPV types such that one type was persistent during consecutive *HPV* visits. Histology was available for 10% of all clinic visits. Misclassification resulting from cytology likely leads to undetected cervical abnormalities (e.g. false negatives), such that some women are classified as *HPV* when in fact they are *HSIL*. Thus, detection of *HSIL* may be underestimated. However, it is important to note that histology is an imperfect gold standard with demonstrated low reproducibility (30). Data on ART were limited for several studies included in the analysis; therefore, the impact of CD4+ count was examined to indirectly capture some of these treatment effects. Further, this sample is largely comprised of women in their thirties who may have been previously infected with HPV, such that a

positive HPV test represents incident DNA detection rather than incident infection (i.e. potential reactivation effects).

This study has a number of strengths, most notably a large longitudinal sample to directly compare HIV-positive and HIV-negative women. These data allowed for the examination of the impact of HIV type, as both HIV-1 and HIV-2 are endemic to West Africa. HPV-16 and HPV-18 were tested for in each study included in the analysis; therefore, we were able to examine the roles of these highly oncogenic types separate from other types. Further, this sample includes registered CSWs. As CSW are estimated to represent roughly 25% of new HIV cases in Africa, this population is particularly relevant for co-infection research (44). Finally, factors known to impact HPV natural history were minimal in this sample (e.g. oral contraception use, smoking, prior cervical cancer screening/treatment, and HPV vaccination).

Our findings suggest that targeted screening programs for the high-risk population of HIV-positive women are needed, in addition to HPV vaccination of youth prior to HIV infection. However, optimal screening modality, frequency, triage, and treatment threshold remain unknown, requiring both expert clinical review and cost-effectiveness examination.

Several studies and pilot projects have demonstrated the feasibility of implementing cervical cancer screening and treatment in low-resource settings (45), although significant economic and infrastructural challenges remain (46). While screening with cytology has been widely adopted in high-resource settings resulting in major declines in cervical cancer incidence over the past 60 years, significant laboratory, equipment, and clinical expertise requirements have limited the capacity to establish cytology in low-resource settings (46, 47). Further, cytology has low reproducibility and sensitivity in comparison to other strategies, necessitating frequent screening to achieve high effectiveness (30, 47). As such, a substantial effort has been made to identify and evaluate screening strategies that are more contextually relevant for the high-risk setting of sub-Saharan Africa. Hybrid Capture 2 (HC2) HPV testing is highly sensitive and reproducible with the potential for self-collection of samples (48), yet requires significant investment in laboratory equipment and technician expertise (46). Both cytology and HPV testing involve laboratory processing time leading to delays in obtaining results and an additional clinic visit if treatment is needed. Visual inspection with acetic acid (VIA) may be most suitable to low-resource settings as it involves naked eye inspection of the cervix (yielding immediate results), and requires little clinical expertise and no laboratory equipment for processing samples. However, inter-observer differences in subjectively determining positivity, concerns regarding over treatment, and the potential for small lesions to remain undetected may reduce the effectiveness of VIA (46). The more recently developed careHPV™ test offers the benefits of standard HPV testing, yields rapid results (roughly two hours of processing time), uses mobile battery operated processing equipment, and requires minimal technical training (46, 49). Both VIA and rapid HPV testing can be implemented as part of a same day ‘screen and treat’ approach, minimizing losses to follow-up and delayed treatment. Immediate treatment with cryotherapy has minimal infrastructure requirements and can be provided by nurses and midwives (46). In contrast, surgical treatment with conization or loop electrosurgical excision procedure (LEEP) require significant infrastructure and technical training, and are further limited by

the lack of pathology services in sub-Saharan settings (50). These various factors for testing and treatment must be considered in order to provide contextually relevant screening recommendations.

Importantly, both test and treatment effectiveness is reduced when applied to HIV-positive women adding uncertainty regarding optimal cervical cancer prevention strategies for this high-risk population (51–56). For instance, given the high prevalence of HPV among HIV-positive women, HPV testing as a stand-alone strategy may be inefficient for identifying women for targeted follow-up (due to a low positive predictive value such that a positive HPV test is not in itself actionable and other intermediate steps may be required). Furthermore, while HIV-positive women have significantly higher rates of adverse transitions than HIV-negative women, absolute risk remains low which could indicate that testing frequencies currently used among the general population may be extended to HIV-positive women. However, increased false negatives among the HIV-positive population may indicate the need for more frequent testing, and potential poor retention in care for certain settings (i.e. rural with limited access to follow-up care) may necessitate aggressive treatment of pre-cancer lesions.

In summary, there is a pressing need for targeted screening in the high-risk population of HIV-positive women. These efforts can subsequently be used to inform the eventual implementation of comprehensive population-based screening paradigms which are not yet widely available in the limited resource setting of sub-Saharan Africa (57). Given the distinct natural history of HPV among HIV-positive women (as demonstrated in our analyses), as well as varying test and treatment attributes, it is important to quantitatively compare the potential impact of screening strategies to understand how these factors collectively impact HPV-associated morbidity and mortality. In the absence of comprehensive cost-effectiveness analyses specific to low-resource settings of sub-Saharan Africa, frequent screening with VIA beginning at the time of HIV diagnosis with conservative triage of positive cases may be worth examining. Identifying optimal targeted prevention strategies is essential given modeling evidence suggesting that cervical cancer among HIV-positive women may increase in the future due to extended life expectancy resulting from ART (58, 59).

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Abbreviations

HIV	human immunodeficiency virus
AIDS	autoimmune deficiency syndrome

HPV	human papillomavirus
ART	antiretroviral therapy
CSW	commercial sex worker
ASCUS	atypical squamous cells of undetermined significance
LSIL	low-grade squamous intraepithelial lesion
HSIL	high-grade squamous intraepithelial lesion
ICC	invasive cervical cancer
PCR	polymerase chain reaction
WHO	World Health Organization

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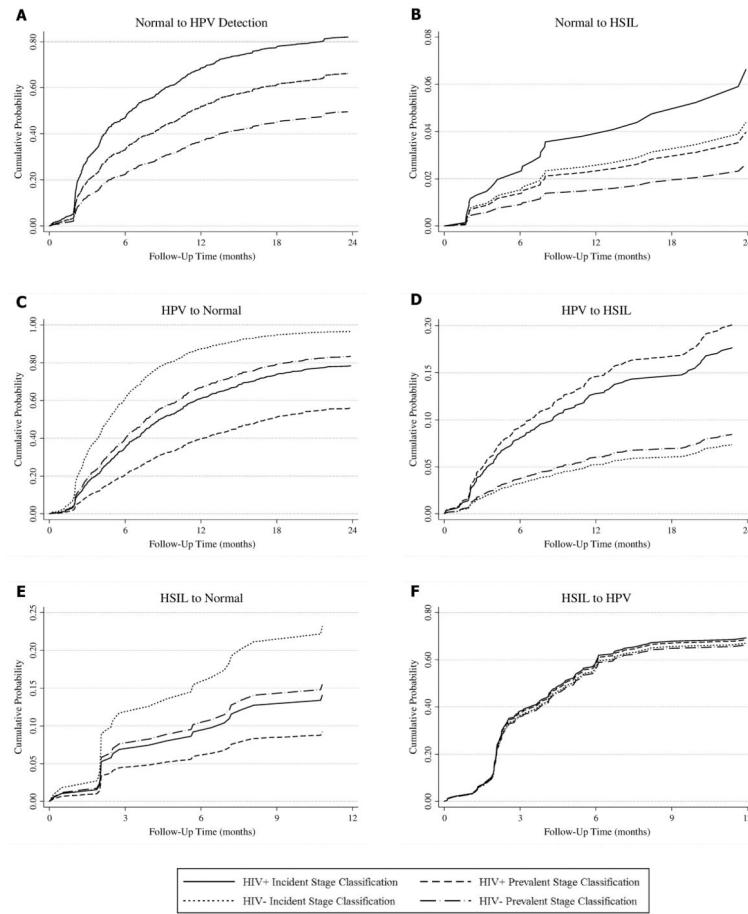


Figure 1. Predicted cumulative probabilities for HIV-positive and HIV-negative Senegalese women ^a
a: Note different x- and y- axis scaling between figures

Overview of included studies

Table 1

No.	Study Title	Study duration	Inclusion criteria	Study sample ^a	Type-specific HPV DNA detection	Relevant references
1	Natural history of cervical neoplasia in HIV-1 and HIV-2	1994 – 1999	Age 15	755 (291 HIV+)	12 types up to 1998 27 types thereafter	(9, 10, 18)
2	Control of HIV-1 by HIV-2 associated immune responses	2000 – 2006	Age 15 ART naive	121 (119 HIV+)	38 types	(22)
3	Epidemiology of HIV-1/HIV-2 dual infection	2001 – 2005	Age 15 HIV-1/2+ On ART	7 (7 HIV+)	38 types	(23)
4	Developing new approaches for cervical cancer control	2002 – 2007	Age 15	3 (3 HIV+)	38 types	(19)
5	Antiretroviral therapy for HIV-2 infection in Senegal	2005 – ongoing	Age 16 HIV-2+ On ART	48 (48 HIV+)	38 types	(20, 25)
6	HIV-associated DNA hyper-methylation in cervical cancer	2005 – 2010	Age 18	386 (150 HIV+)	38 types	(21, 24)

^aSample size for the present study differs from the total samples reported in prior publications due to missing HIV status and/or the lack of longitudinal cytology/histology and HPV DNA data.

Table 2

Baseline characteristics of pooled study sample

Characteristic	HIV- (n = 702)	HIV+ (n = 618)	Total N = 1,320
Age (years)			
Median	34	35	35
Interquartile range	27–44	29–42	28–43
Age at Sexual Debut (years)			
Median	17	17	17
Interquartile range	15–19	15–19	15–19
Lifetime sex partners (%)			
None	0.6	0.2	0.4
One	45.5	31.4	39.0
2–5	25.1	34.9	28.6
6–10	1.7	1.8	1.8
>10	27.1	31.7	30.2
Religion (%)			
Muslim	85.5	89.9	87.8
Christian	14.1	9.7	11.8
Other/None	0.4	0.4	0.4
Marital status (%)			
Married (monogamy)	34.4	26.7	30.9
Married (polygamy)	25.5	14.7	20.5
Never married	15.4	8.9	12.4
Separated/divorced	20.4	30.7	25.1
Widowed	4.3	19.0	11.1
Education (%)			
None	42.6	50.5	46.3
Primary School	30.1	33.3	31.6
Secondary School	24.3	15.2	20.0
University	3.0	1.0	2.1
Follow-up duration (days)			
Median	780	812	796
Interquartile range	368–1,155	384–1,344	377–1,224
Interval between clinic visits (days)			
Median	139	134	136
Interquartile range	122–232	119–238	121–235
Baseline cervical natural history stage (%)			
Normal	31.8	41.0	36.1
HPV	64.1	52.3	58.5
HSIL	4.1	6.7	5.4
Commercial sex worker (%)	24.8	30.4	27.4
Cigarette Use (% , ever)	15.4	18.9	17.0

Characteristic	HIV- (n = 702)	HIV+ (n = 618)	Total N = 1,320
Contraception Use (% , ever)	46.9	36.6	42.2
Alcohol Consumption (% , ever)	9.0	13.9	11.2
<i>HIV-positive subsample</i>			
HIV-type (%)			
HIV-1	-	63.3	-
HIV-2	-	28.3	-
Dual HIV-1/2 infection	-	8.4	-
CD4+ count (cells/ μ L)			
Median	-	406	-
Interquartile range	-	241–620	-
ART use (% , ever)	-	30.8	-
AIDS ^a (%)	-	25.1	-

^aClassified as either having CD4+ count <200 recorded at any point during follow-up or a WHO Stage 4 AIDS defining event.

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Table 3

Hazard ratios for HIV-positive and HIV-negative Senegalese women ^a

Initial Health Stage	Total	N	Transition Health Stage		
			Hazard ratio	P-value	n
			<i>HPV</i>		
<i>Normal</i>	1079	512			28
HIV+	509	298	1.58 (1.32–1.89)	<0.0001	17
HIV–	570	214	ref.	-	11
Incident classification	607	298	1.58 (1.33–1.88)	<0.0001	18
Prevalent classification	472	214	ref.	-	10
			<i>HSIL</i>		
			<i>Normal</i>		
Total	1265	764			
HIV+	634	308	0.46 (0.39–0.54)	<0.0001	87
HIV–	631	456	ref.	-	39
Incident classification	499	327	1.87 (1.60–2.18)	<0.0001	51
Prevalent classification	766	437	ref.	-	75
			<i>HPV</i>		
			<i>Normal</i>		
Total	179	25			
HIV+	117	14	0.57 (0.26–1.29)	0.18	81
HIV–	62	11	ref.	-	37
Incident classification	109	17	1.57 (0.70–3.49)	0.27	71
Prevalent classification	70	8	ref.	-	47

^aTransition numbers (by row) do not sum to the total provided as women remained in the same state for the duration of follow-up or transitioned to cervical cancer. Transitions to cancer were not examined due to extremely limited sample sizes.

Abbreviations: Normal, HPV-negative; HPV, HPV-positive with one or more type in the absence of HSIL; HSIL, HPV-positive with one or more type with the presence of HSIL; Incident, stage differed from the preceding visit; Prevalent, baseline visit or stage was the same as the preceding visit.

Table 4

Univariate analyses of potential effect modifiers, HIV-positive women ^a

Initial Health Stage	Total	n	Transition Health Stage				
			Hazard ratio	P-value	N	Hazard ratio	P-value
<i>Normal</i>							
CD4+ count < 200	57	29	1.09 (0.72–1.67)	0.68	4	2.86 (0.95–8.58)	0.06
CD4+ count ≥ 200	423	260	ref.	-	13	ref.	-
HPV - 16/18	27	24	0.97 (0.64–1.49)	0.90	3	4.62 (1.10–19.42)	0.04
HPV - Other	206	189	ref.	-	5	ref.	-
HPV - Multiple Types	71	64	1.07 (0.79–1.43)	0.67	3	1.37 (0.33–5.69)	0.66
HPV - Single Type	162	149	ref.	-	5	ref.	-
Age ≤ 25	69	47	1.56 (1.14–2.14)	0.0060	1	0.41 (0.05–3.10)	0.39
Age > 25	436	249	ref.	-	16	ref.	-
HIV-1	316	205	1.85 (1.38–2.49)	<0.0001	8	0.48 (0.19–1.21)	0.12
HIV-2	155	74	ref.	-	9	ref.	-
<i>HPV</i>							
<i>Normal</i>							
CD4+ count < 200	116	38	0.57 (0.39–0.83)	0.003	23	1.83 (1.07–3.14)	0.0282
CD4+ count ≥ 200	477	261	ref.	-	58	ref.	-
HPV - 16/18	176	29	0.35 (0.23–0.54)	<0.0001	32	2.20 (1.34–3.62)	0.0018
HPV - Other	473	186	ref.	-	42	ref.	-
HPV - Multiple Types	327	66	0.34 (0.25–0.47)	<0.0001	51	2.33 (1.40–3.88)	0.0011
HPV - Single Type	322	149	ref.	-	23	ref.	-
Age ≤ 25	87	46	1.29 (0.89–1.87)	0.17	13	1.14 (0.57–2.27)	0.72
Age > 25	545	261	ref.	-	70	ref.	-
HIV-1	425	192	0.83 (0.64–1.08)	0.17	63	1.69 (0.94–3.03)	0.08
HIV-2	163	94	ref.	-	16	ref.	-
<i>HSIL</i>							

^aNote that transition numbers (by row) do not sum to total as some women remained in the same state for the duration of follow-up. HPV-type specific analyses for transitions from *Normal* (the initial state) were conducted based on the type acquired, while analyses of transitions from *HPV* (the initial state) were conducted based on what type women had while HPV-positive.

Abbreviations: Normal, HPV-negative; HPV, HPV-positive with one or more type in the absence of HSIL; HSIL, HPV-positive with one or more type with the presence of HSIL; Incident, stage differed from the preceding visit; Prevalent, baseline visit or stage was the same as the preceding visit.

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