Association between male circumcision and human papillomavirus infection in males and

females: a systematic review, meta-analysis, and meta-regression

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ABSTRACT

Background: Human papillomavirus (HPV) infection is a necessary cause of cervical cancer and is associated with anal, penile, vaginal, and vulvar cancers. Previous studies have suggested a protective effect of male circumcision (MC) on HPV infections in males, and that this protection may be conferred to their female sexual partners. We synthesized the available evidence on the association between MC and HPV infections in males and females.

Methods: We performed a systematic review and meta-analysis of the effect of MC on the prevalence, incidence, and clearance of genital HPV infections in heterosexual males and their female sexual partners. We searched multiple databases for studies that assessed MC status and tested for the presence of genital HPV DNA. We used random-effects meta-analysis models to estimate summary measures of effect and 95% confidence intervals (CI) for the prevalence, incidence, and clearance of HPV infections in males and females. We assessed effect modification for prevalence in males using random-effects meta-regression.

Findings: We included 32 publications encompassing 25 unique study populations. MC was associated with decreased odds of prevalent HPV infections (odds ratio 0.45, CI 0.34-0.61), a reduced rate of incident HPV infections (incidence rate ratio 0.69, CI 0.57-0.83), and an increased risk of clearing HPV infections (risk ratio 1.44, CI 1.28-1.61) at the glans penis. Effect modification by sampling site was observed for HPV prevalence in males, with greater protection conferred by MC at the glans than the shaft (OR 0.68, 95% CI 0.48-0.98). Females with circumcised sexual partners were at reduced risk for all outcomes.

Interpretation: MC protects against various HPV infection outcomes, especially at the glans, and may be a viable prophylactic strategy in regions with a high burden of HPV-associated disease where the HPV vaccine is not commercially available. That the protective effect of MC on HPV

infection prevalence varies by penile site has important implications for epidemiologic studies of

HPV transmission.

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RESEARCH IN CONTEXT

Evidence before this study: Previous meta-analyses published in 2011, 2012, and 2017 have assessed the impact of MC on genital HPV infections in males, while systematic reviews published in 2017 and 2019 have described the impact of MC on women's sexual health outcomes. All meta-analyses of males found a protective effect of MC on HPV prevalence, with inconsistent evidence for the association between MC and HPV incidence and clearance. Systematic reviews in females found a protective effect of MC on HPV prevalence.

Added value of this study: We identified an additional 12 publications (including one randomized controlled trial) that were not included in the most recently published systematic review and metaanalysis. We found that in males, MC conferred protection against prevalent HPV infections at the glans and shaft of the penis, protected against the acquisition of HPV infections at the glans, and resulted in increased clearance of HPV infections at the glans and shaft. We also found that MC protected females against various HPV infection outcomes. We considered anatomical site in all analyses and explored effect modification using a meta-regression approach. Our meta-analysis also examined the impact of MC on various HPV infection outcomes in females. To our knowledge, the latter two types of analyses had not been done before.

Implications of all the available evidence: Countries with a high burden of HPV-associated diseases, or where the HPV vaccine is not commercially available, may wish to consider male circumcision as a preventive strategy. Both males and their female sexual partners may benefit from MC for protection from HPV infections.

1 INTRODUCTION

Human papillomavirus (HPV) is the most common sexually transmitted infection worldwide.¹
Persistent infection with high-risk HPV types (hrHPV) is a necessary cause of cervical cancer and
is associated with penile, anal, vaginal, vulvar, and head and neck cancers,²⁻⁴ while infection with
some low-risk HPV types (lrHPV) is associated with genital warts.¹

6 Male circumcision (MC) protects against a variety of sexually transmitted infections, including 7 human immunodeficiency virus (HIV), herpes simplex type 2, trichomoniasis, chancroid, and syphilis.⁵⁻⁷ Several randomized controlled trials (RCT) evaluating the association between MC and 8 HIV acquisition have also included analyses of HPV as secondary endpoints.^{8,9} Most observational 9 10 studies of the relationship between MC and HPV infections in males have been cross-sectional in 11 nature, and few have evaluated the risk of HPV infection in female partners of circumcised and uncircumcised males. Previous systematic reviews^{10,11} and meta-analyses¹²⁻¹⁴ found that MC 12 13 protects against a variety of HPV infection outcomes in males and their female sexual partners. 14 However, gaps in knowledge remain and multiple studies on the topic have been recently 15 published, necessitating an update to the existing literature. In this systematic review, we synthesize the growing evidence suggestive of a protective relationship between MC and HPV 16 17 infections in males, and the conferred protection to female sexual partners.

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19 METHODS

20 Search strategy and selection criteria

We searched for studies that 1) included participants with no HPV-associated genital lesions, 2) tested for the presence of HPV DNA in genital epithelial cells, 3) assessed the male circumcision status, and 4) assessed the prevalence, incidence, and/or clearance of HPV infections. We included

24 both observational and experimental study designs but excluded case reports and case series. We 25 included studies of both males and females of any age but excluded studies that focused solely on men who have sex with men and people living with HIV from the sample due to HIV's direct 26 27 effect on HPV infection risk due to immunosuppression and shared sexual transmission 28 characteristics.^{15,16} Multiple publications from the same study population were eligible for 29 inclusion if they assessed distinct outcomes. We applied no country, date, or language restrictions. 30 We searched the MEDLINE, Embase, Scopus, Cochrane, LILACS, and ProQuest Dissertations & 31 Theses Global databases to identify relevant records published up to 22 June 2022. We also 32 manually searched for potentially eligible studies from previous knowledge syntheses and 33 conference abstracts. The search strategy for each database, developed with input from a university 34 librarian, is included in Supplementary Table 1.

After de-duplicating search results in EndNote, S.S. and C.L. independently screened the abstract of each record to determine relevancy. For papers deemed potentially relevant, we obtained and independently screened the record's full text. Disagreements at both stages were resolved by consensus.

39 Data analysis

40 S.S. and C.L. performed data extraction using a standardized spreadsheet. Each author extracted 41 data from half of the included records, which was subsequently verified by the alternate author. 42 Extracted data included study characteristics (design, year(s), country(s) and their economic 43 development as defined by the World Bank,¹⁷ population description, number of visits if 44 longitudinal), exposure and outcome methods (MC assessment method, genital sites sampled, 45 frequency of genital sampling, sampling method, HPV DNA detection and genotyping method, 46 HPV types detected and genotyped), study population results (sample size, sex, age at baseline,

47 HPV prevalence at baseline), and outcome-related data (outcome type, i.e., prevalence, incidence, 48 clearance; HPV risk grouping; number of samples analyzed; number circumcised and 49 uncircumcised; number of prevalent or incident or cleared infections; person-time at risk; effect 50 estimate and 95% confidence interval (95% CI); and covariates adjusted for). Whenever possible, 51 we extracted separate estimates for infection with any HPV type, hrHPV, and lrHPV, as well as 52 separate estimates from samples of different sites of the penis: shaft and/or scrotum only (hereafter 53 referred to as shaft), glans and/or urethra and/or foreskin only (hereafter referred to as glans), and 54 from combinations of shaft sites and glans sites (hereafter referred to as combined site). We 55 extracted the adjusted estimate when available and the crude estimate otherwise. If raw data were 56 presented without effect estimates, we calculated the odds ratio and 95% CI using OpenEpi's two by two table function.¹⁸ If effect estimates used circumcised males as the reference category, we 57 58 took the reciprocal of the estimate and its 95% CI. If relevant data or analyses were mentioned but 59 not quantitatively reported, we contacted the study authors.

We assessed the risk of bias in each study using customized versions of the Newcastle-Ottawa scale for cross-sectional and cohort studies and the Cochrane risk-of-bias tool for randomized trials.¹⁹ Studies were deemed to have a low risk of bias if they were assigned a score of 7 or greater on the Newcastle-Ottawa scale or a score of low across at least 4 domains using the Cochrane riskof-bias tool.

We extracted effect estimates for the relationship between MC and HPV infection prevalence, incidence, and clearance for multiple sexes (male and female), HPV risk groupings (any HPV, hrHPV, and lrHPV), and sampling sites (glans, shaft, and combined site). We used the *meta* command in Stata (version 17.0, StataCorp, College Station, Texas) to calculate pooled odds ratios, risk ratios, incidence rate ratios, hazard ratios, and their corresponding 95% CIs using a

restricted maximum likelihood model. We assessed study heterogeneity using the I² statistic. We 70 71 used random effects models for analyses with an I² of greater than or equal to 25% and fixed effects 72 models otherwise. For all analyses, we performed subgroup analyses by sampling site (glans-only 73 vs. shaft-only or combined site) and HPV oncogenicity (hrHPV vs. lrHPV) to assess potential 74 effect modification. We conducted univariate random-effects meta-regression of prevalence studies in males with clustering by study using the *metafor* package²⁰ in R (version 4.2.0, R Core 75 76 Group, Vienna) to explore potential effect modification by study characteristics: year of 77 publication, sites sampled, study country's economic development, and whether the study 78 controlled for confounding. For studies of prevalence that reported risk or prevalence ratios, we 79 used the raw data to calculate an odds ratio so that the study could be included in the meta-80 regression.

We performed several sensitivity analyses to assess the robustness of our findings. We repeated our primary analyses of prevalence, incidence, and clearance in males including only studies judged to have a low risk of bias. We additionally repeated our analysis of prevalence in males stratifying by whether studies controlled for confounding, conducted leave-one-out analyses to assess the impact of any one study on the pooled estimate,²¹ and assessed publication bias using a funnel plot and the Egger test.²²

87 This study protocol was registered in PROSPERO (registration number CRD42020211591).

88 **Role of the funding source**

89 The funder of the study had no role in study design, data collection, data analysis, data90 interpretation, or writing of the report.

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92 **RESULTS**

93 We identified 1,409 potentially eligible records through systematic database searches and 10 94 through manual searches, of which 624 remained after de-duplication (Figure 1). We excluded 520 95 records after title and abstract screening, leaving 104 full-text records for assessment. We excluded 96 18 records for reporting on the same study population as an included record, 25 records that studied 97 participants with HPV-associated lesions, seven records for not sampling a genital site, seven 98 records for missing exposure or outcome data, seven records for not having an outcome of interest, 99 and eight records for failure to obtain needed data that were missing from the authors of the original 100 studies (Supplementary Table 2). In total, we included 32 records in our systematic review and 101 meta-analysis.

102 Characteristics of these publications,^{6,8,23-52} which were published between 2002 and 2022, are 103 presented in Table 1. The 32 studies encompassed 25 unique study populations.

104 Of these publications, 17 were cross-sectional studies, ten were cohort studies (of which two were 105 analyzed cross-sectionally), and five were RCTs. Studies were conducted in North America 106 (n=12), South America (n=3), Europe (n=4), Asia (n=1), Africa (n=8), and intercontinentally 107 (n=4). MC status was either self-reported or reported by a partner (n=11), reported by a clinician 108 (n=16), or randomized and verified by a clinician (n=5). All studies assessed the presence of HPV 109 DNA by PCR, 23 of which genotyped for 20 or more HPV types. Samples were taken via swab 110 (n=18), textured paper and swab (n=6), brush (n=5), and brush and swab (n=2). Samples in males 111 were taken from multiple sites, including the urethra (n=5), foreskin (n=15), glans and/or corona 112 (n=27), shaft (n=19), scrotum (n=15), and perianal area (n=4) whereas samples in females were 113 taken from the cervix and vagina (n=5). The PCR primer sets used for HPV DNA typing were 114 PGMY09/11 (n=13), MY09/11 (n=8), GP5+/6+ (n=4), SPF10 (n=3), CpI/CpIIG (n=1), and type-

115 specific and assay-specific primers (n=3). HPV prevalence among all participants at baseline 116 ranged from 8.7% to 69.8%.

117 A total of 21 studies reported estimates for the association between MC and prevalent HPV 118 infections in males (Supplementary Table 3). Sample sizes ranged from 37 to 3,969. MC was associated with significantly decreased odds of prevalent HPV infections at both the glans (OR 119 120 0.45, 95% CI $0.34-0.61, I^2=0.0\%$) and the shaft or combined sites (OR 0.66, 95% CI 0.50-0.87, 121 I^2 =67.1%), with a stronger effect observed at the glans (Figure 2). MC was associated with a 122 significantly decreased risk of prevalent HPV infections at the glans (RR 0.57, 95% CI 0.39-0.82, 123 $I^2=82\cdot 2\%$), but not at the shaft or combined sites (RR 0.96, 95% CI 0.92–1.01, $I^2=0.0\%$). Findings 124 were similar when stratifying by hrHPV and lrHPV types (Supplementary Figures 1 and 2). 125 Nine studies examined the association between MC and HPV incidence in males (Supplementary 126 Table 4), with sample sizes ranging from 210 to 4,033. A significant protective effect of MC was 127 observed for the incidence rate (IRR 0.69, 95% CI 0.57–0.83, $I^2=0.0\%$) at the glans, but not for 128 the hazard rate at the shaft or combined sites (HR 1.04, 95% CI 0.94-1.16, $I^2=0.0\%$) (Figure 3). 129 Results were similar when stratifying by HPV oncogenicity (Supplementary Figures 3 and 4). 130 Seven publications, with sample sizes of 285 to 4,033, examined the association between MC and 131 HPV clearance in males (Supplementary Table 5). Both the risk and hazard rate of HPV infection 132 clearance were significantly increased at the glans of circumcised males (HR 1.86, 95% CI 1.49– 133 2.31, $I^2=0.0\%$, RR 1.44, 95% CI 1.28-1.61, $I^2=0.0\%$) (Figure 5), while the hazard rate was 134 increased at the shaft and combined sites, albeit non-significantly (HR 1.41, 95% CI 0.81-2.42, 135 I^2 =86.9%). Results remained similar when separately examining hrHPV and lrHPV.

136 Six studies examined the association between MC and various HPV outcomes in females137 (Supplementary Table 6). Sample sizes ranged from 61 to 2,735. All studies assessed the

prevalence of HPV infections, while two additionally assessed the acquisition of HPV infections and one assessed the clearance of HPV infections. The risk of prevalent hrHPV infections and the incident rate of hrHPV infections were significantly reduced in female partners of circumcised males (RR 0.66, 95% CI 0.49–0.89, $I^2=35.0\%$, IRR 0.77, 95% CI 0.63–0.93, $I^2=0.0\%$) (Figure 5). For all other outcomes, point estimates were protective, but did not reach statistical significance.

We found evidence of an effect modification of the association between MC and HPV prevalence in males by sampling site, with a 32% increase in the protective effect of MC at the glans than at the shaft or combined sites (OR 0.68, 95% CI 0.48-0.98) (Table 2). No effect modification was observed for year of publication, primary study country's economic development, or whether the study accounted for confounding.

149 Most studies were judged to have a low risk of bias (Supplementary Tables 7–9). Restricting to 150 studies with a low risk of bias did not change any findings for studies of prevalence 151 (Supplementary Figure 7), incidence (Supplementary Figure 8), and clearance (Supplementary 152 Figure 9) of HPV infections in males. When stratifying studies of prevalence in males by whether 153 or not they controlled for confounding, we observed that studies that did control for confounding 154 found significantly protective effects of MC at the glans on both the odds and risk scales 155 (Supplementary Figure 10). Studies that did not control for confounding only evaluated the effects 156 of MC at the glans on the odds scale and did not find a protective effect. Studies that controlled 157 for confounding also found significantly protective effects of MC at the shaft and combined sites 158 on the odds scale, but not the risk scale, and studies that did not control for confounding did not 159 find any protective effect of MC at the shaft or combined sites. Excluding any given study of 160 prevalence in males reporting an OR did not significantly change the pooled estimate

(Supplementary Figure 11), and we did not observe evidence of publication bias for these studies
(Supplementary Figure 12, p value for Egger test 0.95). Publication bias could not be assessed for
other outcomes due to the limited number of studies that used the same effect measure.

164

165 **DISCUSSION**

166 Findings of this meta-analysis suggest that MC results in reduced prevalent and incident HPV 167 infections and increased clearance of HPV infections at the glans penis, as well as reduced 168 prevalent infections at the shaft. Protection may also be conferred to the female sexual partners of 169 circumcised males. Our findings of the protective effect of MC against various HPV infection outcomes are consistent with those of previous reviews.¹⁰⁻¹⁴ However, our analysis of the varying 170 171 effect of MC at different anatomical sites of the penis and the use of a meta-regression approach 172 to assess for effect modification have not been done before. To the best of our knowledge, our 173 analysis seems to be the first to include both males and females in the same review.

174 Infections with hrHPV are of most clinical relevance, as persistent infection with hrHPV is a 175 necessary cause of cervical cancer and is associated with various anogenital cancers.²⁻⁴ All 176 estimates for the association between MC and hrHPV infection prevalence, incidence, and 177 clearance found that MC had a significantly protective effect at the glans, and either a protective 178 effect or no effect at the shaft. MC was not found to be a risk factor for HPV infections in any of 179 our meta-analyses.

180 We included several publications that were not part of the most recently published systematic 181 reviews on the topic: in males, we included an additional nine records of prevalence,^{8,24,29,31,36,40,42,45,51} three of incidence,^{8,45,48} and four of clearance^{8,32,45,48} that were not 182 included in Zhu's 2017 review and meta-analysis¹⁴. In females, we added three records^{28,36,45} that 183

184 were absent in Morris' 2019 review.¹¹ The addition of new records did not result in different 185 conclusions than those of previous reviews, but rather provided further and more detailed evidence 186 for the same interpretations, especially for the varying levels of protection MC confers at different 187 anatomical sites of the penis.

188 The biological mechanism by which MC is suggested to protect against HPV infections is still 189 unclear; the prevailing theories suggesting differences in keratinization and in the local immune 190 environment of the penis as plausible. It was originally thought that the glans of the circumcised penis is more keratinized than that of the uncircumcised penis⁵³ and less vulnerable to the 191 192 acquisition of sexually transmitted infections during sexual intercourse. However, anatomic and 193 histological studies have failed to find consistent results on the differences in keratinization 194 between the glans of circumcised and uncircumcised males.⁵⁴ MC has also been postulated to 195 change the local immune environment of the penis through changes in the microbiome and 196 immune cell density. Removal of the foreskin eliminates the anaerobic environment of the preputial cavity.⁵⁵ The Ugandan trial of MC found that circumcised males had a decreased total 197 198 bacterial load and reduced biodiversity in their microbiota,⁵⁶ whereas a 2017 study of 51 females 199 showed that those who were HPV-positive were more likely to have a diverse array of facultative and strict anaerobic bacteria in their vaginal microbiome.⁵⁷ MC may protect against HPV by 200 201 reducing the diversity of anaerobic bacteria in the penile microbiota. Finally, different anatomical sites of the penis have different distributions of immune cells.⁵⁸ The removal of the foreskin and 202 203 the immune cells within it may result in different cytokine environments and inflammatory 204 responses to pathogen entry, both of which are associated with the risk of HPV infections.^{54,59-61} 205 Our review had many strengths. We searched a diverse array of databases and validated our search 206 strategy with a librarian. We did not apply study design or language restrictions and we included

both males and females, multiple HPV infection-related outcomes, different HPV risk groupings,
and different anatomical sampling sites.

209 Our review also had several limitations. We included the term "circumcision" in our search 210 strategy and may not have captured records that measured MC and HPV infection without directly 211 assessing their association. We were unable to consider other factors that may play a role in MC's 212 association with HPV infection, such as method of MC, whether MC was performed before or 213 after sexual debut, and number of sexual partners, as these variables were not collected in the vast 214 majority of the included studies. Only three of the 25 unique study populations included in our 215 review came from RCTs, which limited our ability to assess causality. However, it is noteworthy that all RCTs assessing HPV infections in males^{8,9,30,48} found a protective effect of MC at the glans 216 217 for prevalence, incidence and clearance of all HPV types, including hrHPV, and all estimates but 218 one were statistically significant. 219 In conclusion, results from our systematic review and meta-analysis support that MC protects

against HPV infections in a diverse population of males, particularly at the glans, and that
protection may be passed on to female partners. MC may be a viable preventive strategy for HPV
infections, especially in regions with a high burden of HPV-associated cancers and where the HPV
vaccine is not commercially available.

Contributors: SBS and ELF conceptualized the project. SBS and CL conducted the literature search, screening, and data extraction. CL and SBS performed data analysis. SBS drafted the manuscript. CL, MZ, and ELF critically reviewed the manuscript. ELF and MZ provided supervision and guidance. All authors had access to all the data in the study and had final responsibility for the decision to submit for publication. Both SBS and CL directly accessed and verified the underlying data reported in the manuscript.

Declaration of interests: ELF and MZ hold a patent related to the discovery "DNA methylation markers for early detection of cervical cancer", registered at the Office of Innovation and Partnerships, McGill University, Montreal, Quebec, Canada (October 2018). ELF has served as consultant to Roche, Merck, and BD on HPV diagnostics and prevention. The other authors declare no competing interests.

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First author (year)	Country(ies), years conducted	World Bank economic classification	Study population	Number enrolled	Circumcision assessment	Sites sampled	HPV DNA genotyping method
Randomized	controlled trials						
Gray (2010)	Uganda, 2003– 2006	Low-income country	Males enrolled in the Rakai-1 trial	840	Randomized and verified by a clinician	Glans	MY09/11 PCR
Smith (2021)	Kenya, 2002– 2005	Middle-income country	Males enrolled in the Kisumu circumcision trial	2,193	Randomized and verified by a clinician	Inner foreskin, glans, outer foreskin, shaft	GP5+/6+ PCR
Tobian (2009)	Uganda, 2003– 2007	Low-income country	Males enrolled in the Rakai-1 and Rakai-2 trials	3,393	Randomized and verified by a clinician	Foreskin, glans	PGMY09/11 PCR
Tobian (2012)	Uganda, 2002– 2009	Low-income country	HIV-positive and negative males enrolled in the Rakai-1 and Rakai-2 trials	776ª	Randomized and verified by a clinician	Glans	PGMY09/11 PCR
Wawer (2011)	Uganda, 2003– 2007	Low-income country	Female partners of males enrolled in the Rakai-1 and Rakai-2 trials	1,245	Randomized and verified by a clinician	Vagina	MY09/11 PCR
Cohort studie	25						
Albero (2013) ^b	Brazil, Mexico, United States, 2005–2009	Predominantly high-income countries	Males from the general population, universities, and organized healthcare systems	3,969	Clinical exam	Foreskin, glans, shaft, scrotum	PGMY09/11 PCR

Table 1: Characteristics of included records according to study design

Albero (2014)	Brazil, Mexico, United States, 2005–2009	Predominantly high-income countries	Males from the general population, universities, and organized healthcare systems	4,003	Clinical exam	Foreskin, glans, shaft, scrotum	PGMY09/11 PCR
Hernandez (2008) ^b	United States, 2004–2006	High-income country	Male university students in Hawaii	379	Clinical exam	Foreskin, glans, shaft, scrotum	PGMY09/11 PCR
Hernandez (2010)	United States, 2004–2006	High-income country	Male university students in Hawaii	357	Clinical exam	Foreskin, glans, shaft, scrotum	PGMY09/11 PCR
Lajous (2005)	Mexico, 2002– 2005	Middle-income country	Healthy military males	1,030	Self-report	Urethra, glans, shaft, scrotum	MY09/11 PCR
		Males from the general population	285	Self-report	Foreskin, glans, shaft, scrotum	PGMY09/11 PCR	
Nielson (2009)	United States, 2002–2005	High-income country	Males from the general population	463	Self-report	Foreskin, urethra, glans, shaft, scrotum, perianal area, anus	PGMY09/11 PCR
Partridge (2007)	United States, 2003–2006	High-income country	Male university students in Washington	240	Clinical exam	Foreskin, urethra, glans, shaft, scrotum	PGMY09/11 PCR
Shapiro (2022)	Canada, 2005– 2011	High-income country	Female university students in Montreal and their male sexual partners	826	Clinical exam	Foreskin, glans, shaft	PGMY09/11 PCR

VanBuskirk (2011)	United States, 2003–2009	High-income country	Male university students in Washington	477	Clinical exam	Foreskin, glans, shaft, scrotum	MY09/11 PCR
Cross-section	nal studies						
Baldwin (2004)	United States, 2000–2001	High-income country	Males attending an STI clinic	393	Clinical exam	Glans	PGMY09/11 PCR
Bleeker (2005)	Netherlands, 1995–2002	High-income country	Males from a non-STI dermatology clinic and male partners of females with CIN	356	Clinical exam	Foreskin, glans	GP5+/6+ PCR
Castellsagué (2002)	Spain, Colombia, Brazil, Thailand, Philippines, 1985–1993	Predominantly middle-income countries	Male partners of case females with cervical cancer and healthy control females	1,913	Clinical exam	Urethra, glans	MY09/11 PCR
Contreras (2008)	Mexico, 2005– 2006	Middle-income country	Females with rheumatoid arthritis	61	Self-report	Cervix	CpI/CpIIG PCR
Da Rocha (2015)	Brazil, 2011– 2013	Middle-income country	Males from an STI clinic, a dermatology clinic, a university, and a factory	261	Self-report	Glans	MY09/11 PCR
Hebnes (2021)	Denmark, 2006–2007	High-income country	Military males	2,460	Clinical exam	Preputial cavity, glans, shaft, scrotum, perineum	SPF10 PCR
Mbulawa (2009)	South Africa, NR	Middle-income country	Sexually active Black heterosexual couples	254 ^a	Self-report	Foreskin, glans, shaft	PGMY09/11 PCR
Obiri- Yeboah (2017)	Ghana, NR	Middle-income country	Females attending an HIV or medical outpatient clinic	170ª	Partner report	Cervix	RT-PCR with type-specific primers

Ogilvie (2009)	Canada, NR	High-income country	Heterosexual males attending an STI clinic	262	Clinical exam	Foreskin, glans, shaft, scrotum	Amplicor® primer PCR
Olesen (2019)	Tanzania, 2009	Middle-income country	Males from urban and rural areas	1,902ª	Clinical exam	Foreskin, glans, shaft	PGMY09/11 PCR
Rocha (2012)	Brazil, NR	Middle-income country	Heterosexual couples in which the female HPV-related cervical lesions	43	Clinical exam	Foreskin, glans	GP5+/6+ PCR
Rombaldi (2006)	Brazil, 2003– 2004	Middle-income country	Male sexual partners of females with CIN	99	Self-report	Foreskin, urethra, glans, shaft	MY09/11 PCR
Roura (2012)	Spain, 2007– 2008	High-income country	Females attending cervical cancer screening	3,261	Partner report	Cervix	SPF10 PCR
Shin (2004)	South Korea, 2002	High-income country	Male university students	381	Self-report	Urethra, glans, shaft, scrotum	SPF10 PCR
Svare (2002)	Denmark, 1993	High-income country	Males attending an STI clinic	198	Self-report	Glans, shaft, scrotum, perianal area	GP5+/6+ PCR
Vaccarella (2006)	Mexico, 2003– 2004	Middle-income country	Males requesting a vasectomy	779	Clinical exam	Glans, shaft, scrotum	MY09/11 PCR
Vardas (2011)	18 countries in Africa, Asia- Pacific, Europe, Latin America, and North America, NR	Predominantly middle-income countries	Heterosexual males with 1–5 female lifetime sexual partners	3,463	Clinical exam	Penis (specific sites NR), scrotum, perianal area	RT-PCR with type-specific primers

Abbreviations: NR, not reported; STI: sexually transmitted infection

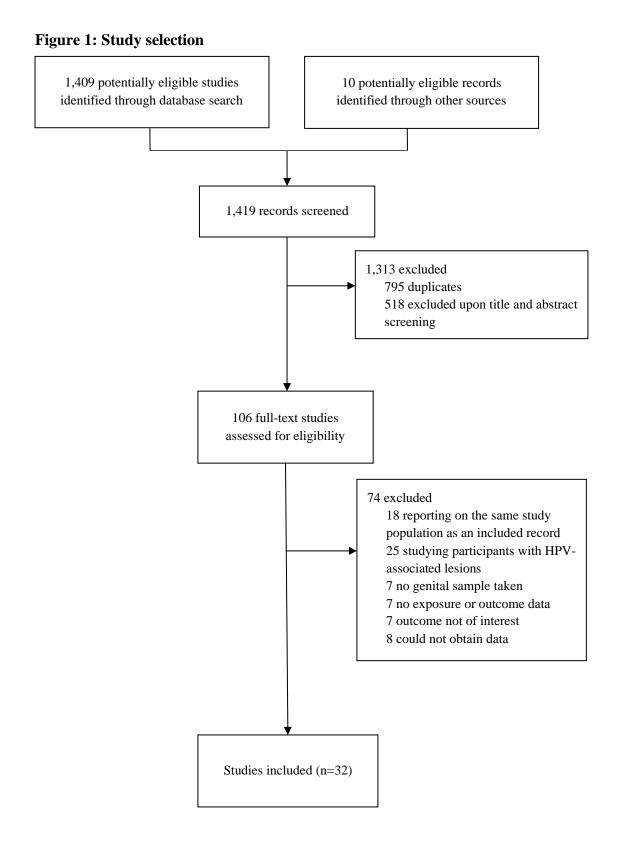
^a Only HIV-negative males were included

^b Cohort study analyzed cross-sectionally

	Number of	Univari	ate
Potential effect modifier	studies (%)	OR (95% CI)	p-value for modifier
Year			
2009 or earlier	15 (62.5)	1.00 (reference)	
2010 or later	9 (37.5)	1.28(0.88 - 1.86)	0.19
Site, n (%)			
Combined/shaft-only	15 (62.5)	1.00 (reference)	
Glans	9 (37.5)	0.68(0.48-0.98)	0.04
Economic development, n (%)			
High-income country	12 (50.0)	1.00 (reference)	
Low- or middle-income country	12 (50.0)	0.87 (0.61 - 1.24)	0.45
Control for confounding, n (%)			
Yes	17 (70.8)	1.00 (reference)	
No	7 (29.2)	1.54(0.89-2.67)	0.12

Table 2. Meta-regression of studies assessing the association between male circumcision and HPV prevalence in males

Abbreviations: CI, confidence interval; HPV, human papillomavirus; OR, odds ratio



Glans Cross-sectional Any HPV Glans OR 0.37 (0.16, 0.85) Castellsque 2002 Cross-sectional Any HPV Glans OR 0.44 (0.23, 0.83) Bidkin 2004 Cross-sectional Any HPV Glans OR 0.44 (0.23, 0.83) Bleeker 2005 Cross-sectional Any HPV Glans OR 0.51 (0.27, 0.97) Bleeker 2005 Cross-sectional Any HPV Glans OR 0.52 (0.04, 7.03) Da Rocha 2012 Cross-sectional Any HPV Glans OR 0.52 (0.04, 7.03) Bocha 2012 Cross-sectional Any HPV Glans OR 0.45 (0.34, 0.61) Test of θ = 0: z = -5.26, p = 0.00 Smith 2021 RCT Any HPV Glans RR 0.48 (0.41, 0.56) 2 Smith 2021 RCT Any HPV Glans RR 0.70 (0.53, 0.92) Heterogeneity: r ⁴ = 0.06, F = 82.20%, H ² = 5.62 D.57 (0.39, 0.82) D.57 (0.39, 0.82) D.57 (0.39, 0.82) Vardas 2011 Cross-sectional Any HPV Combined OR 0.70 (0.48, 0.99) D.70 (0.48, 0.99) D.70 (0.48, 0.99) D.70 (0.48, 0.99) D.70 (0.48, 0.99)	- · · ·						Effect esti		
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Baldwin 2004 Cross-sectional Any HPV Glans OR	Castellsague 2002	Cross-sectional	Any HPV	Glans	OR		0.37 (0.16,	0.85)	1139
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Heterogeneity: $\tau^2 = 0.00$, $H^2 = 0.00\%$, $H^2 = 1.00$ 0.45 (0.34, 0.61) Test of $\theta = 0$: $z = -5.26$, $p = 0.00$ 0.45 (0.34, 0.61) Smith 2021 RCT Any HPV Glans RR 0.48 (0.41, 0.56) 0.56) Tobian 2009 RCT Any HPV Glans RR 0.70 (0.53, 0.92) 0.57 (0.39, 0.82) Heterogeneity: $\tau^2 = 0.06$, $I^2 = 82.20\%$, $H^2 = 5.62$ 0.57 (0.39, 0.82) 0.57 (0.39, 0.82) Combined or shaft-only Vardas 2011 Cross-sectional Any HPV Combined OR 0.70 (0.49, 0.99) 0.70 (0.49, 0.99) Uscarella 2006 Cross-sectional Any HPV Combined OR 0.87 (0.52, 1.45) 0.48 (0.30, 0.77) Vaccarella 2006 Cross-sectional Any HPV Combined OR 0.81 (0.56, 1.17) 0.43 (0.40) Nielson 2009 Cross-sectional Any HPV Combined OR 0.83 (0.37, 1.07) 0.81 (0.65, 1.17) Shapiro 2022 Cohort Any HPV Combined OR 0.63 (0.43, 0.40) 0.53 (0.28, 1.00) Shapiro 2022 Crobort Any HPV Combined OR 0.61 (0.67, 1.42) 0.63 (0.37, 1.07) Mulawa 2009 Cross-sectional Any HPV Combined OR 0.54 (0.20, 1.42) 0.	Da Rocha 2015	Cross-sectional	Any HPV	Glans	OR —		0.52 (0.04,	7.03)	182
Test of $\theta = 0$: $z = -5.26$, $p = 0.00$ Smith 2021 RCT Any HPV Glans R 0.48 (0.41, 0.56) 2 0.70 (0.53, 0.92) Tobian 2009 RCT Any HPV Glans R 0.70 (0.53, 0.92) 0.57 (0.39, 0.82) Test of $\theta = 0$: $z = -2.99$, $p = 0.00$ Combined or shaft-only Vardas 2011 Cross-sectional Any HPV Combined OR 0.70 (0.69, 1.18) 0.70 (0.49, 0.99) 0.57 (0.39, 0.82) Vardas 2011 Cross-sectional Any HPV Combined OR 0.70 (0.49, 0.99) 0.70 (0.49, 0.99) 0.70 (0.49, 0.99) 0.70 (0.49, 0.99) 0.70 (0.49, 0.99) 0.70 (0.49, 0.99) 0.70 (0.49, 0.99) 0.70 (0.49, 0.99) 0.70 (0.49, 0.99) 0.70 (0.49, 0.99) 0.70 (0.49, 0.99) 0.70 (0.49, 0.99) 0.70 (0.49, 0.99) 0.70 (0.49, 0.99) 0.70 (0.49, 0.99) 0.70 (0.49, 0.99) 0.70 (0.49, 0.99) 0.70 (0.49, 0.99) 0.70 (0.49, 0.99) 0.70 (0.40, 0.81) 0.70 (0.49, 0.99) 0.70 (0.49, 0.99) 0.70 (0.49, 0.99) 0.70 (0.49, 0.99) 0.70 (0.40, 0.81) 0.70 (0.40, 0.81) 0.70 (0.40, 0.81) 0.70 (0.40, 0.81) 0.70 (0.40, 0.81) 0.70 (0.40, 0.81) 0.70 (0.40, 0.81) 0.70 (0.40, 0.81) 0.70 (0.40, 0.81) 0.70 (0.40, 0.81) 0.70 (0.40, 0.81) 0	Rocha 2012	Cross-sectional	Any HPV	Glans	OR –		— 1.32 (0.06,	28.87)	37
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Tobian 2009 RCT Any HPV Glans RR 0.70 (0.53, 0.92) Heterogeneity: $\tau^2 = 0.06$, $l^2 = 82.20\%$, $H^2 = 5.62$ 0.57 (0.39, 0.82) 0.57 (0.39, 0.82) Combined or shaft-only Vardas 2011 Cross-sectional Any HPV Combined OR 0.70 (0.69, 1.18) Hebnes 2021 Cross-sectional Any HPV Combined OR 0.70 (0.69, 1.18) Lajous 2005 Cohort Any HPV Combined OR 0.87 (0.52, 1.45) Vaccarella 2006 Cross-sectional Any HPV Combined OR 0.88 (0.30, 0.77) Vaccarella 2006 Cross-sectional Any HPV Combined OR 0.81 (0.56, 1.17) Shapiro 2022 Cohort Any HPV Combined OR 0.81 (0.56, 1.17) Shin 2004 Cross-sectional Any HPV Combined OR 0.81 (0.56, 1.17) Mbulawa 2009 Cross-sectional Any HPV Combined OR 0.63 (0.37, 1.07) Mbulawa 2009 Cross-sectional Any HPV Combined OR 0.54 (0.20, 1.42) Oglivie 2009 Cross-sectional Any HPV Combined OR 0.54 (0.20, 1.42) Oglivie 2009 Cross-sectional Any HPV Combined OR 0.54 (0.20, 0.63) Rombaldi	Test of $\theta = 0$: z = -5.26,	p = 0.00							
Heterogeneity: $r^2 = 0.06$, $l^2 = 82.20\%$, $H^2 = 5.62$ 0.57 (0.39, 0.82) Test of $\theta = 0$: $z = -2.99$, $p = 0.00$ Combined or shaft-only Vardas 2011 Cross-sectional Any HPV Combined OR 0.90 (0.69, 1.18) Hebnes 2021 Cross-sectional Any HPV Combined OR 0.70 (0.49, 0.99) Olesen 2019 Cross-sectional Any HPV Combined OR 0.87 (0.52, 1.45) Lajous 2005 Cohort Any HPV Combined OR 0.87 (0.52, 1.45) Vaccarella 2006 Cross-sectional Any HPV Combined OR 0.83 (0.28, 1.00) Shairo 2022 Cohort Any HPV Combined OR 0.81 (0.56, 1.17) Shairo 2022 Cohort Any HPV Combined OR 0.81 (0.56, 1.17) Shairo 2022 Cohort Any HPV Combined OR 0.81 (0.56, 1.17) Mbulawa 2009 Cross-sectional Any HPV Combined OR 0.84 (0.20, 1.42) Ogilvic 2009 Cross-sectional Any HPV Combined OR 0.54 (0.20, 1.42) Ogilvic 2009 Cross-sectional Any HPV Combined OR 0.54 (0.20, 1.42) Ogilvic 2009 Cross-sectional Any HPV Combined OR 0.50 (0.60, 0.63) Romb	Smith 2021	RCT	Any HPV	Glans	RR		0.48 (0.41,	0.56)	2193
Test of $\theta = 0: z = -2.99, p = 0.00$ Combined or shaft-only Vardas 2011 Cross-sectional Any HPV Combined OR 0.90 (0.69, 1.18) Hebnes 2021 Cross-sectional Any HPV Combined OR 0.70 (0.49, 0.99) Olesen 2019 Cross-sectional Any HPV Combined OR 0.87 (0.52, 1.45) Lajous 2005 Cohort Any HPV Combined OR 0.87 (0.52, 1.45) Vaccarella 2006 Cross-sectional Any HPV Combined OR 0.83 (0.30, 0.77) Vaccarella 2006 Cross-sectional Any HPV Combined OR 0.81 (0.56, 1.17) Shin 2009 Cross-sectional Any HPV Combined OR 0.81 (0.56, 1.17) Shin 2004 Cross-sectional Any HPV Combined OR 0.81 (0.56, 1.17) Shin 2004 Cross-sectional Any HPV Combined OR 0.81 (0.56, 1.17) Mbulawa 2009 Cross-sectional Any HPV Combined OR 0.84 (0.20, 1.42) Oglivic 2009 Cross-sectional Any HPV Combined OR 0.54 (0.20, 1.42) Oglivic 2009 Cross-sectional Any HPV Combined OR 0.50 (0.66, 0.63) Rombaldi 2006 Cross-sectional Any HPV Combined OR 0.20 (0.06, 0.63) <	Tobian 2009	RCT	Any HPV	Glans	RR	· · · · ·	0.70 (0.53,	0.92)	520
Combined or shaft-onlyVardas 2011Cross-sectional Any HPVCombined OR $0.90 (0.69, 1.18)$ Hebnes 2021Cross-sectional Any HPVCombined OR $0.70 (0.49, 0.99)$ Olesen 2019Cross-sectional hrHPVCombined OR $0.87 (0.52, 1.45)$ Lajous 2005CohortAny HPVCombined OR $0.87 (0.52, 1.45)$ Vaccarella 2006Cross-sectional Any HPVCombined OR $0.87 (0.52, 1.45)$ Nielson 2009Cross-sectional Any HPVCombined OR $0.83 (0.28, 1.00)$ Shapiro 2022CohortAny HPVCombined OR $0.81 (0.56, 1.17)$ Shin 2004Cross-sectional Any HPVCombined OR $0.81 (0.56, 1.17)$ Shin 2004Cross-sectional Any HPVCombined OR $0.81 (0.56, 1.17)$ Mbulawa 2009Cross-sectional Any HPVCombined OR $0.63 (0.37, 1.07)$ Mbulawa 2009Cross-sectional Any HPVCombined OR $0.54 (0.20, 1.42)$ Ogilvie 2009Cross-sectional Any HPVCombined OR $0.54 (0.20, 1.42)$ Svare 2002Cross-sectional Any HPVCombined OR $0.20 (0.06, 0.63)$ Rombaldi 2006Cross-sectional Any HPVCombined OR $0.20 (0.06, 0.63)$ Heterogeneity: $\tau^2 = 0.14$, I ² = 67.13%, H ² = 3.04 $0.96 (0.91, 1.01)$ Test of $\theta = 0: z = -2.98, p = 0.00$ Any HPVCombined PR $0.96 (0.92, 1.01)$ Albero 2013CohortAny HPVCombined PR $0.96 (0.92, 1.01)$ Smith 2021RCTAny HPVShaftRR $0.96 (0.92, 1.01)$	Heterogeneity: τ ² = 0.06	δ, I² = 82.20%, H² = 5.6	62			\blacklozenge	0.57 (0.39,	0.82)	
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Hernandez 2008CohortAny HPVShaftOR0.63 (0.37, 1.07)Mbulawa 2009Cross-sectionalAny HPVCombinedOR0.54 (0.20, 1.42)Ogilvie 2009Cross-sectionalAny HPVCombinedOR1.14 (0.67, 1.94)Svare 2002Cross-sectionalAny HPVCombinedOR0.20 (0.06, 0.63)Rombaldi 2006Cross-sectionalAny HPVCombinedOR2.07 (0.46, 9.35)Heterogeneity: $\tau^2 = 0.14$, $l^2 = 67.13\%$, $H^2 = 3.04$ \bullet \bullet \bullet 0.96 (0.91, 1.01)Test of $\theta = 0$: $z = -2.98$, $p = 0.00$ ChortAny HPVCombinedPR0.96 (0.91, 1.01)Albero 2013CohortAny HPVShaftRR1.05 (0.84, 1.31)2Smith 2021RCTAny HPVShaftRR0.96 (0.92, 1.01)	•					-		,	413
Mbulawa 2009 Cross-sectional Any HPV Combined OR 0.54 (0.20, 1.42) Ogilvie 2009 Cross-sectional Any HPV Combined OR 1.14 (0.67, 1.94) Svare 2002 Cross-sectional Any HPV Combined OR 0.20 (0.06, 0.63) Rombaldi 2006 Cross-sectional Any HPV Combined OR 0.20 (0.06, 0.63) Heterogeneity: $\tau^2 = 0.14$, $l^2 = 67.13\%$, $H^2 = 3.04$ 0.66 (0.50, 0.87) 0.66 (0.50, 0.87) Test of $\theta = 0$: $z = -2.98$, $p = 0.00$ Any HPV Combined PR 0.96 (0.91, 1.01) Albero 2013 Cohort Any HPV Shaft RR 1.05 (0.84, 1.31) 0.96 (0.92, 1.01) Heterogeneity: $\tau^2 = 0.00$, $l^2 = 0.00\%$, $H^2 = 1.00$ 0.96 (0.92, 1.01) 0.96 (0.92, 1.01) 0.96 (0.92, 1.01)							•	,	368
Ogilvie 2009 Cross-sectional Any HPV Combined OR 1.14 (0.67, 1.94) Svare 2002 Cross-sectional Any HPV Combined OR 0.20 (0.06, 0.63) Rombaldi 2006 Cross-sectional Any HPV Combined OR 2.07 (0.46, 9.35) Heterogeneity: $\tau^2 = 0.14$, $l^2 = 67.13\%$, $H^2 = 3.04$ 0.66 (0.50, 0.87) 0.66 (0.50, 0.87) Test of $\theta = 0$: $z = -2.98$, $p = 0.00$ Albero 2013 Cohort Any HPV Combined PR 0.96 (0.91, 1.01) Smith 2021 RCT Any HPV Shaft RR 1.05 (0.84, 1.31) 2.09 (0.92, 1.01)						-84	•		334
Svare 2002 Cross-sectional Any HPV Combined OR 0.20 (0.06, 0.63) Rombaldi 2006 Cross-sectional Any HPV Combined OR 0.20 (0.06, 0.63) Heterogeneity: $\tau^2 = 0.14$, $l^2 = 67.13\%$, $H^2 = 3.04$ 0.66 (0.50, 0.87) Test of $\theta = 0$: $z = -2.98$, $p = 0.00$ 0.00 (0.06, 0.63) Albero 2013 Cohort Any HPV Combined PR 0.96 (0.91, 1.01) Smith 2021 RCT Any HPV Shaft RR 1.05 (0.84, 1.31) Heterogeneity: $\tau^2 = 0.00$, $l^2 = 0.00\%$, $H^2 = 1.00$ 0.96 (0.92, 1.01) 0.96 (0.92, 1.01)								1.42)	298
Rombaldi 2006Cross-sectional Any HPVCombined OR2.07 (0.46, 9.35)Heterogeneity: $τ^2 = 0.14$, $l^2 = 67.13\%$, $H^2 = 3.04$ 0.66 (0.50, 0.87)Test of $θ = 0$: $z = -2.98$, $p = 0.00$ 0.06 (0.91, 1.01)Albero 2013CohortAny HPVSmith 2021RCTAny HPVHeterogeneity: $τ^2 = 0.00$, $l^2 = 0.00\%$, $H^2 = 1.00$ 0.96 (0.92, 1.01)	•	Cross-sectional	Any HPV			-		1.94)	262
Heterogeneity: $\tau^2 = 0.14$, $l^2 = 67.13\%$, $H^2 = 3.04$ Test of $\theta = 0$: $z = -2.98$, $p = 0.00$ Albero 2013 Cohort Any HPV Combined PR 0.96 (0.91, 1.01) Combined PR 0.96 (0.91, 1.01) Combined PR 1.05 (0.84, 1.31) Combined PR 1.05 (0.84, 1.31) Combined PR 1.05 (0.92, 1.01)							-	0.63)	198
Test of $\theta = 0: z = -2.98, p = 0.00$ Albero 2013 Cohort Any HPV Combined PR 0.96 (0.91, 1.01) 1.01) Smith 2021 RCT Any HPV Shaft RR 1.05 (0.84, 1.31) 1.01) Heterogeneity: $\tau^2 = 0.00, l^2 = 0.00\%, H^2 = 1.00$ 0.96 (0.92, 1.01)			•	Combined	OR		•	,	99
Albero 2013 Cohort Any HPV Combined PR 0.96 (0.91, 1.01) 1.01 (0.96) Smith 2021 RCT Any HPV Shaft RR 1.05 (0.84, 1.31) 1.03 (0.96, 0.92, 1.01) Heterogeneity: $\tau^2 = 0.00$, $I^2 = 0.00\%$, $H^2 = 1.00$ 0.96 (0.92, 1.01) 0.96 (0.92, 1.01)	Heterogeneity: $\tau^2 = 0.14$	I, I² = 67.13%, H² = 3.0)4			•	0.66 (0.50,	0.87)	
Smith 2021RCTAny HPVShaftRR $1.05(0.84, 1.31)$ $2.000(0.92, 1.01)$ Heterogeneity: $\tau^2 = 0.00$, $I^2 = 0.00\%$, $H^2 = 1.00$ $0.96(0.92, 1.01)$	Test of $\theta = 0$: $z = -2.98$,	p = 0.00							
Heterogeneity: $\tau^2 = 0.00$, $I^2 = 0.00\%$, $H^2 = 1.00$ 0.96 (0.92, 1.01)	Albero 2013	Cohort	Any HPV	Combined	PR		0.96 (0.91,	1.01)	3969
	Smith 2021	RCT	Any HPV	Shaft	RR	<u>.</u>	1.05 (0.84,	1.31)	2193
Test of $\theta = 0$: $z = -1.40$, $p = 0.16$	Heterogeneity: $\tau^2 = 0.00$), l² = 0.00%, H² = 1.00)				0.96 (0.92,	1.01)	
	Test of $\theta = 0$: z = -1.40,	p = 0.16							
0.05 0.50 1.50 5.00						0.50.1.50.5.00			

Figure 2: Studies of male circumcision and HPV prevalence in males by sampling site

Abbreviations: CI, confidence interval; HPV, human papillomavirus; hrHPV, high-risk HPV; OR, odds ratio; PR, prevalence ratio; RCT, randomized controlled trial; RR, risk ratio

Design	0					Effect estimate	
	Uncogenic	ity Site N	leasur	е		(95% CI)	n
RCT	hrHPV	Glans	IRR			0.70 (0.55, 0.89)) 776
RCT	hrHPV	Glans	IRR			0.67 (0.50, 0.90)) 448
H ² = 1.00				•		0.69 (0.57, 0.83))
= 0.00							
RCT	Any HPV	Glans	HR	-8-		0.51 (0.43, 0.61)) 1196
					_		
	-			-	-		
	-				-		
				_			
	-			8			
	Any HPV	Combined	HR				
						1.04 (0.94, 1.16))
: 0.39							
Cohort	Any HPV	Combined	IRR			0.77 (0.37, 1.60)) 55
Cohort	Any HPV	Combined	OR			1.12 (0.45, 2.79)) 210
				0.50	1.50	5.00	
	RCT , H ² = 1.00 = 0.00 RCT Cohort RCT Cohort Cohort , H ² = 1.00 = 0.39 Cohort	RCThrHPV $H^2 = 1.00$ $H^2 = 1.00$ RCTAny HPVRCTAny HPVCohortAny HPVCohorthrHPVCohortAny HPVCohortAny HPVCohortAny HPV $H^2 = 1.00$ H^2 CohortAny HPV	RCThrHPVGlans $H^2 = 1.00$ RCTAny HPVGlansRCTAny HPVGlansCohortAny HPVCombinedRCTAny HPVShaftCohortAny HPVCombinedCohortAny HPVCombinedCohortAny HPVCombinedCohortAny HPVCombined $Any HPV$ CombinedCombined	RCThrHPVGlansIRR $, H^2 = 1.00$ IRRIRRRCTAny HPVGlansIRRRCTAny HPVGlansIRRCohortAny HPVShaftIRRCohorthrHPVShaftIRRCohortAny HPVCombinedIRRCohortAny HPVCombinedIRRCohortAny HPVCombinedIRRCohortAny HPVCombinedIRRCohortAny HPVCombinedIRR $, H^2 = 1.00$ IRRIRR	RCT hrHPV Glans IRR , H² = 1.00 = 0.00 RCT Any HPV Glans HR RCT Any HPV Glans HR Cohort Any HPV Shaft HR RCT Any HPV Shaft HR Cohort Any HPV Combined HR ., H² = 1.00	RCT hrHPV Glans IRR H ² = 1.00 RCT Any HPV Glans HR RCT Any HPV Glans HR Cohort Any HPV Shaft HR Cohort hrHPV Combined HR Cohort Any HPV Combined IRR	RCT hrHPV Glans IRR 0.67 (0.50, 0.90) , H ² = 1.00 0.69 (0.57, 0.83) = 0.00 RCT Any HPV Glans HR RCT Any HPV Glans HR 0.51 (0.43, 0.61) Cohort Any HPV Combined HR 1.08 (0.91, 1.28) RCT Any HPV Shaft HR 1.01 (0.87, 1.17) Cohort Any HPV Combined HR 1.10 (0.83, 1.46) Cohort Any HPV Combined HR 1.10 (0.83, 1.46) Cohort Any HPV Combined HR 1.10 (0.60, 2.01) , H ² = 1.00 .039 .077 (0.37, 1.60) 1.04 (0.94, 1.16) Cohort Any HPV Combined IRR

Figure 3: Studies of male circumcision and HPV incidence in males by sampling sit	e

Abbreviations: CI, confidence interval; HPV, human papillomavirus; HR, hazard ratio; hrHPV, high-risk HPV; IRR, incidence rate ratio; RCT, randomized controlled trial;

Study	Design	Oncogenic	ity Site N	leasure		Effect estimate (95% CI)	n
Glans							
Smith 2021	RCT	Any HPV	Glans	HR		1.90 (1.49, 2.42)	2032
Hernandez 2010	Cohort	Any HPV	Glans	HR		1.69 (1.02, 2.79)	357
Heterogeneity: $\tau^2 = 0.00$, I ² = 0.00%,	H ² = 1.00			•	1.86 (1.49, 2.31)	
Test of θ = 0: z = 5.56, p	= 0.00						
Tobian 2012	RCT	hrHPV	Glans	RR	-	1.48 (1.26, 1.74)	776
Gray 2010	RCT	hrHPV	Glans	CRR	-	1.39 (1.17, 1.65)	448
Combined or shaft-only							
Albero 2014		-	Combined		I _	0.95 (0.88, 1.02)	
Smith 2021	RCT	Any HPV		HR		2.19 (1.34, 3.58)	624
Hernandez 2010		Any HPV		HR —		0.94 (0.63, 1.41)	357
Lu 2009		•	Combined	HR		3.10 (1.19, 8.10)	285
Heterogeneity: $\tau^2 = 0.24$		$H^2 = 7.63$				1.41 (0.82, 2.42)	
Test of θ = 0: z = 1.25, p	= 0.21						
Shapiro 2022	Cohort	Any HPV	Combined			0.81 (0.52, 1.25)	131
				0.50	1.50 5.	00	

	· · · · · ·		e in males by sampling	· · · • 4 · ·
HIGHTE 4. MURIES OF	r male circiimcición	and HPV clearanc	e in males hy sampling	T CITA
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Abbreviations: CI, confidence interval; CRR, clearance rate ratio; HPV, human papillomavirus; HR, hazard ratio; hrHPV, high-risk HPV; RCT, randomized controlled trial; RR, risk ratio

Study	Design O	ncogenicity	Measure		Effect esti (95% 0		n
Prevalence, any HPV							
Roura 2012	Cross-sectional	Any HPV	OR	-	0.80 (0.59,	1.08)	2735
Shapiro 2022	Cohort	Any HPV	OR	- +	1.05 (0.75,	1.46)	413
Mbulawa 2009	Cross-sectional	Any HPV	OR		0.42 (0.14,	1.23)	202
Contreras 2008	Cross-sectional	Any HPV	OR		9.00 (1.23,	65.93)	61
Heterogeneity: $\tau^2 = 0.0$	2, I ² = 23.90%, H ² =	1.31		•	0.91 (0.68,	1.21)	
Test of $\theta = 0$: $z = -0.67$, p = 0.50						
Wawer 2011	RCT	Any HPV	RR		0.81 (0.72,	0.92)	1032
Prevalence, hrHPV							
Roura 2012	Cross-sectional	hrHPV	OR	-8-	0.70 (0.49,	0.99)	2735
Shapiro 2022	Cohort	hrHPV	OR	-	1.06 (0.74,		413
Heterogeneity: $\tau^2 = 0.0$				▲	0.86 (0.57,		
Test of $\theta = 0$: $z = -0.73$,	,	
Wawer 2011	RCT	hrHPV	RR	-	0.72 (0.60,	0.86)	1032
Obiri-Yeboah 2017	Cross-sectional	hrHPV	RR	-8-	0.50 (0.29,	0.87)	170
Heterogeneity: τ ² = 0.0	2, l² = 34.99%, H² =	1.54			0.66 (0.49,	0.89)	
Test of $\theta = 0$: z = -2.68	, p = 0.01			•			
Prevalence, IrHPV Roura 2012	Cross-sectional		OR	_	0.70 (0.40	0.99)	0705
		IrHPV	OR	1	0.70 (0.49, 1.07 (0.72,		413
Shapiro 2022 Heterogeneity: τ ² = 0.0	Cohort		UR	X	0.86 (0.56,	,	413
Test of $\theta = 0$: $z = -0.74$					0.000 (0.000)	,	
Wawer 2011	RCT	IrHPV	RR	-	0.77 (0.66,	0.90)	1032
Incidence, any HPV							
Shapiro 2022	Cohort	Any HPV	IBB		0.59 (0.16,	2,19)	61
		,			,	,	
Incidence, hrHPV							
Wawer 2011	RCT	hrHPV	IRR		0.77 (0.63,	0.94)	1051
Shapiro 2022	Cohort	hrHPV	IRR		0.80 (0.12,	5.36)	32
Heterogeneity: $\tau^2 = 0.0$ Test of $\theta = 0$: $z = -2.64$.00		•	0.77 (0.63,	0.93)	
Incidence, IrHPV							
Wawer 2011	RCT	IrHPV	IRR		0.83 (0.69,	1.00)	1051
Shapiro 2022	Cohort	IrHPV	IRR		0.23 (0.03,	1.65)	30
Heterogeneity: τ ² = 0.3	1, I ² = 37.97%, H ² =	1.61		•	0.65 (0.24,	1.75)	
Test of $\theta = 0$: z = -0.86	, p = 0.39						
Clearance, hrHPV				L			
Wawer 2011	RCT	hrHPV	RR		1.12 (1.02,	1.22)	1051
				0.501.50 5.00			

Figure 5: Studies of male circumcision and various HPV outcomes in females

Abbreviations: CI, confidence interval; HPV, human papillomavirus; hrHPV, high-risk HPV; IRR, incidence rate ratio; lrHPV, low-risk HPV; OR, odds ratio; RCT, randomized controlled trial; RR, risk ratio