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## ***Mycoplasma genitalium* infection among HIV-infected pregnant African women and implications for mother-to-child transmission of HIV**

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### **Abstract**

**Objective:** Many sexually transmitted infections (STIs) increase risk of mother-to-child transmission (MTCT) of HIV, but the effect of *Mycoplasma genitalium* (MG) is not known. We hypothesized that MG infection would be common among HIV-infected pregnant women and could be associated with *in utero* and intrapartum MTCT.

**Design:** Observational case-cohort study

**Methods:** This study used specimens from a Kenyan perinatal MTCT cohort (1999-2005) involving HIV-infected women and their infants, who received short-course zidovudine for prevention of MTCT. Vaginal swabs collected at 32 weeks gestation were tested for MG using a transcription-mediated amplification assay. Infant perinatal HIV infection was determined at birth and 4 weeks of age by DNA PCR. Using a case-cohort design, a random sample was generated with 3:1 control: case ratio; prevalence and correlates of MG were assessed with Chi-squared and t-tests; predictors of infant outcomes were analyzed using logistic regression.

**Results:** Among 220 HIV-infected pregnant women evaluated, 47 women (21.4%) had MG. Antenatal MG infection was associated with higher HIV RNA in plasma (5.0 vs. 4.6 log<sub>10</sub>

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**Competing Interests:** Hologic (Gen-Probe) donated materials for *M. genitalium* laboratory testing, but was not involved in any other aspects of the research. P.A.T. has received remuneration for contract work, consulting, and speakers fees from Hologic and SpeedX. No other co-authors report any competing interests to disclose.

copies/ml in MG-positive vs. MG-negative women,  $p=0.02$ ) at 32 weeks. Women with MG were less likely to report prior STIs and genital ulcers (both  $p=0.05$ ). There was no association found between exposure to MG and perinatal MTCT (OR=0.72, 95% CI 0.35, 1.51,  $p=0.39$ ).

**Conclusions:** Vaginal MG infection was frequently detected among Kenyan HIV-infected pregnant women and was associated with higher plasma HIV levels, but was not associated with perinatal transmission of HIV.

### Keywords

*Mycoplasma genitalium*; HIV/AIDS; mother-to-child transmission of HIV; perinatal HIV

### Introduction:

Many sexually transmitted infections (STIs) increase risk of mother-to-child transmission (MTCT) of HIV. Despite rapid scale-up of interventions to eliminate MTCT, approximately 180,000 children acquired HIV in 2017.<sup>[1]</sup> STIs may facilitate perinatal transmission of HIV by increasing genital inflammation and HIV viral shedding; syphilis, gonorrhea (GC), and herpes simplex virus 2 (HSV-2) have all been implicated in MTCT of HIV.<sup>[2-5]</sup> *Mycoplasma genitalium* (MG), a pathogen of the genital tract of both men and women, is an emerging STI.<sup>[6]</sup> To our knowledge, the potential influence of MG in MTCT of HIV has not been investigated. MG infection is associated with increased vaginal cytokine levels and inflammation,<sup>[7]</sup> and women with a high burden of MG have been observed to have increased cervical HIV shedding,<sup>[8]</sup> providing a potential mechanism by which MG infection could lead to increased *in utero* and/or intrapartum MTCT of HIV.

MG has a wide range of presentations, from asymptomatic to severe, and if untreated, can persist for years. MG has been detected in 5%-20% of sexually active African adults<sup>[9]</sup>, and is more common among adults with HIV infection.<sup>[10]</sup> Further, MG is linked to clinical illness including acute and chronic non-gonococcal urethritis in men,<sup>[11]</sup> and urethritis,<sup>[12]</sup> cervicitis,<sup>[13]</sup> tubal infertility,<sup>[14, 15]</sup> endometritis,<sup>[16]</sup> pelvic inflammatory disease<sup>[17]</sup> and salpingitis<sup>[15, 18]</sup> in women. Longitudinal data link MG to female acquisition of HIV.<sup>[19, 20]</sup> Despite the spectrum of disease caused by MG, and, in select settings, a higher prevalence of MG than GC or chlamydia (CT),<sup>[9, 21, 22]</sup> MG is not routinely screened for in Africa, nor is it effectively treated by World Health Organization (WHO) syndromic management guidelines for STIs.<sup>[23]</sup> Antenatal screening for MG is not currently recommended, and there are no FDA approved commercial tests or point-of-care tests available.

Further evidence linking MG to MTCT could provide additional impetus for screening and treatment recommendations, as well as stimulating development of novel and inexpensive diagnostics and more effective treatments. We examined a historical perinatal HIV cohort of Kenyan women to determine the prevalence and correlates of MG infection, and to determine whether antenatal MG infection is associated with increased risk of MTCT.

**Methods:****Population:**

This study utilized specimens collected from a cohort of HIV-infected pregnant women enrolled and followed in Nairobi, Kenya from 1999-2003 to determine immunologic correlates of MTCT. [24, 25] Pregnant women were eligible for enrollment if HIV-seropositive, less than 28 weeks gestation, and willing to participate in longitudinal follow-up for 1-2 years.

**Ethics:**

All women gave written informed consent in English or Kiswahili, and the study was approved by the University of Washington Institutional Review Board and the University of Nairobi/Kenyatta National Hospital Ethical Review Committee.

**Clinical procedures:**

Enrolled women had a pelvic exam at 32 weeks gestation; vaginal Dacron swab specimens were collected and preserved in 1 mL of freezing medium (70% RPMI 1640 with L-glutamine, 25 mM HEPES, 20% fetal bovine serum, 10% DMSO) and cryopreserved at  $-80^{\circ}\text{C}$ . Women were provided with antenatal short-course zidovudine (ZDV) for prevention of MTCT, the standard of care at the time. Women were asked to deliver in a hospital facility, where infant blood was collected for HIV testing at birth; infants were seen and followed again at 4 weeks of age for HIV testing.

**Laboratory methods:**

One hundred microliters of the thawed cryopreserved vaginal swab specimens were subjected to testing for MG using the Aptima research-only transcription-mediated amplification (TMA) assay using primers targeting MG rRNA according to the manufacturer's directions (Hologic).<sup>[26]</sup> Since rRNA is present in multiple copies per cell, this potentially increases the sensitivity of MG detection over standard PCR techniques. Signals correspond to the hybridized target and express as relative light units (RLU), with RLU  $\geq 50,000$  considered positive, RLU  $< 5000$  considered negative, and the remainder intermediate. Intermediate samples were repeated twice, if repeats remained intermediate, the specimen was reported as "indeterminate."

CD4 counts were measured in Nairobi; in Seattle, HIV RNA levels were measured on serum and genital tract specimens using methods validated for HIV subtypes prevalent in Kenya. <sup>[27]</sup> Infant HIV status was determined by HIV DNA testing, confirmed by HIV RNA testing, using a prototype HIV-1 viral load assay (Gen-Probe/Hologic).<sup>[28]</sup> STI testing included: HSV-2 IgG antibody testing by Focus ELISA, confirmed by Western blot; syphilis testing by rapid plasma reagin (Becton Dickinson), confirmed by *Treponema pallidum* hemagglutination assay (Randox); *Trichomonas vaginalis* from vaginal swabs by in-pouch culture testing using the APTIMA platform (Gen-Probe/Hologic); CT and NG testing of cervical swabs with the PCR Amplicor CT/NG test (Roche); and bacterial vaginosis (BV) by Gram stain using the scoring method described by Nugent and Hillier<sup>[29]</sup>.

## Study Design:

The study used a case-cohort design, which creates a synthetic case-control study by randomly choosing specimens from a larger cohort, enabling cost-effective use of expensive laboratory assays.<sup>[30, 31]</sup> The primary analysis defined cases as mothers whose infants acquired HIV (with HIV DNA and RNA detected) within the first 4 weeks of life. The exposure of interest was maternal MG detected at 32 weeks gestation. Mother/infant pairs were excluded *a priori* if the gestation included twins, no infant HIV test results available, elective Caesarean section was performed, or vaginal swabs were unavailable. A random number generator was used to select eligible women from the parent cohort for MG testing. The size of the random sample was determined by multiplying the number of cases of MTCT in the cohort by 3, and adding an additional 10% in case of failed assays.<sup>[32]</sup> MG testing was also conducted on specimens from all women who transmitted HIV to infants perinatally if they were not already included in the random sample.

## Statistical Methods:

In the randomly-selected cohort, prevalence and sociodemographic and immunological correlates of MG infection were analyzed among women using univariate tests including Chi-squared or Fisher's exact test for categorical variables or Student t-tests for continuous variables, and correlates were confirmed with a logistic regression model. All other MTCT cases were then added to the random cohort and logistic regression was used to model the association between MG (exposure) and infant HIV (outcome). A multivariate logistic regression model was constructed, adjusting for known cofactors of maternal HIV transmission (maternal age, plasma HIV RNA level, CD4 count, trichomoniasis, BV, syphilis, CT) and any other correlates of MG infection identified in our random cohort. Each cofactor was added stepwise to the model and included in the final model if significantly associated ( $p < 0.1$ ).

## Sample size and power calculations:

The number of cases of infant HIV acquisition was fixed. Assuming 20% prevalence of MG infection among women, this study had 80% power to detect at least a 2.5-fold increased odds of MTCT among mothers with exposure to MG (2-sided  $\alpha = 0.05$ ).

## Results:

### Case-cohort selection

From the original cohort of 510 women, we identified 408 mother-infant pairs eligible for the primary analysis, which included 67 early infant HIV infections. We then randomly selected 222 samples from the parent cohort. This random sample included 33 women who transmitted HIV to their infants. The remaining 34 cases of MTCT were added to the random cohort for MG testing. In total, 256 swabs were tested for MG, with 254 results.

## Demographic and health data of women and infants in the cohort

MG results were obtained from 220 of the 222 randomly-selected cohort (two specimens failed the assay); baseline characteristics are detailed in Table 1. Prevalence of MG infection was high; 47 (21.4%) women had MG detected at 32 weeks gestation. The remainder tested negative; there were no intermediate results. Women had a median age of 25 years and a median of 8 years of education. The median CD4 count of women was 465 cells/mm<sup>3</sup> (interquartile range (IQR)=310-643 cells/mm<sup>3</sup>) and mean plasma HIV RNA level was 4.68 log<sub>10</sub> copies/ml (standard deviation (SD) = 0.89). Prior STIs were reported by 28 women (12.7%) and prior genital ulcer disease (GUD) was reported by 13 women (6%). STI testing in pregnancy revealed 7 women (3.2%) had syphilis; HSV-2 results were not available for all women, but of 137 women tested, 121 were seropositive (88%). Trichomoniasis was diagnosed in 41 women (18.6%), CT in 10 women (4.6%), GC in 2 women (<1%), and 80 women (38%) had BV on Gram stain. Among 220 randomly-selected mothers, there were 33 total perinatal infant HIV infections (15%), 16 (7.3%) detected at birth, reflecting *in utero* transmission, and 17 infants (7.7%) with no HIV detection at birth and HIV detection at 4 weeks of life, indicating intrapartum transmission.

## Correlates of maternal MG infection

Correlates of maternal MG infection are summarized in Table 2. Women with MG had similar demographic characteristics (age, education, parity, sex partners) as women without MG infection, but were less likely to report a history of STIs, including GUD ( $p=0.05$  for both). Mean plasma HIV RNA levels at 32 weeks gestation were significantly higher among MG-infected mothers, (5.0 log<sub>10</sub> copies/ml for mothers with MG v. 4.6 log<sub>10</sub> copies/ml for mothers without MG,  $p=0.02$ ). Mothers with MG were more likely to be diagnosed with trichomoniasis while pregnant (28% vs. 16%,  $p=0.07$ ). Multivariate modeling did not result in different associations.

## Case-cohort analysis of MG and MTCT:

The case-cohort analysis included the random sample and all cases of infant HIV ( $n=254$  mother-baby pairs). No association between maternal MG infection and infant HIV was seen in an unadjusted model (odds ratio (OR)=0.72 (95% confidence interval (CI) 0.36, 1.51,  $p=0.39$ ) (Table 3). Similarly, in an analysis adjusting for plasma HIV RNA level and BV status, known predictors of infant HIV acquisition, MG was not associated with increased odds of infant HIV acquisition (adjusted OR 0.55 (95% CI 0.25, 1.22,  $p=0.14$ ) and the model showed that maternal plasma HIV RNA level and maternal BV infection remained dominant associations with infant HIV acquisition. We also constructed unadjusted and adjusted models examining maternal MG and different timing of infant HIV acquisition, and did not observe any different associations of MG and *in utero* or intrapartum transmission. Additional multivariate models were run adjusting for significant correlates of MG infection identified in Table 2, including history of STI, history of genital ulcer disease, trichomoniasis in pregnancy, and marital status. Adjusting for these variables did not affect the odds ratios, and so these variables were not included in the final adjusted models.

## Discussion

We detected MG in more than 20% of pregnant women with HIV infection in this Kenyan cohort. MG was more commonly detected than GC, CT, and trichomoniasis; only HSV-2 and BV were more prevalent in this population. The prevalence of MG was higher than most previously reported cohorts, with the exception of cohorts of African women at high risk of HIV,<sup>[9]</sup> and represents an underappreciated burden of this infection in pregnant African women with HIV. Interestingly, MG was detected more often among women who did not self-report a history of bacterial STIs or prior treatment of STIs. This is in contrast to HSV-2 and other STIs which we typically detect more often in women reporting some STI history. Women with MG were also more likely to be diagnosed with trichomoniasis while pregnant (borderline significant  $p=0.07$ ). Other than trichomoniasis, women with MG had similar proportions of all other STIs detected in pregnancy compared to women without MG, which does not align with our expectation that behaviors that predisposed to MG acquisition would result in positive associations between MG and other STIs.

We did not observe an association between MG infection and early infant HIV infection, including both *in utero* or intrapartum HIV transmission. Although the study had sufficient power to detect a fairly large association ( $OR > 2.4$ ) between MG infection and MTCT, it lacked power to detect a smaller association. The direction of the association identified, however, with  $OR < 1$ , indicates that it is unlikely that MG increases MTCT. Our failure to find a strong relationship between MG infection and MTCT is, however, reassuring, as numerous barriers exist to antenatal screening and treatment of MG. There are currently no readily available, inexpensive screening methods to detect MG in antenatal or pre-pregnancy settings.<sup>[33]</sup> We also lack cheap and effective antibiotic therapy to treat MG, especially as macrolide antibiotics are generally avoided during pregnancy.<sup>[34]</sup> Further, treatment of MG is not always successful, antibiotic resistance to both azithromycin and fluoroquinolones is becoming more prevalent,<sup>[35]</sup> and relapse rates are common.<sup>[36]</sup> It is not known whether concurrent treatment of sexual partners is needed to eradicate MG, although best practices for STIs would presume that partner treatment would be recommended.

MG is linked to acquisition of HIV in women in sub-Saharan Africa<sup>[19, 20]</sup>, but it is not known whether women with MG transmit HIV more easily to male sexual partners. Our finding of high prevalence of MG in this cohort, and higher HIV plasma levels in women with MG infection, may be important for targeting efforts to prevent heterosexual HIV transmission.<sup>[37, 38]</sup> MG infection may continue to have public health significance in the fight against HIV. If, as we have shown here, women with MG have higher plasma HIV RNA levels, then it is biologically plausible that MG-infected women may be more likely to transmit HIV to partners, which would be another indication for improved diagnostic and treatment options for persons with MG. We do not have an explanation for our observation of increased HIV RNA in plasma but not cervicovaginal secretions. We expected the converse given that MG is a genital infection. However, our findings are from a small sample of women and should be evaluated in larger studies.

Our study has some limitations. The vaginal swab specimens used in this study were collected 15 years prior to MG testing, possibly leading to TMA target degradation over

time and underestimation of MG in our population. However, such degradation should not change the direction of an association. We used accepted definitions of timing of infant HIV transmission used in prior studies, however, some infants classified as intrapartum transmission may have acquired HIV from early breastfeeding in the first days of life, which would be unrelated to MG exposure. This is a known challenge in MTCT research and is not unique to this cohort. Cohort studies can suffer from longitudinal loss to follow-up, but we observed very low attrition in the first weeks of the cohort (2% by one month of life). MG status was ascertained at 28 weeks gestation, and some women may have acquired MG, or cleared MG, prior to delivery.<sup>[39]</sup> We included only HIV-seropositive women, which limits the generalizability of the findings.

Another limitation is that we lacked statistical power to detect low odds of increased MG exposure among mothers transmitting HIV perinatally. The cohort was studied when short-course ZDV monotherapy in late pregnancy was the standard of care and therefore had rates of HIV transmission to infants that are much higher than current perinatal trends; within this cohort, we have previously demonstrated statistically significant associations with MTCT and both HSV-2 and BV<sup>[5, 40]</sup>. We therefore present evidence that a relationship between MG and MTCT is weaker than the relationship seen with these known cofactors, and the relationship seen was in the direction of no risk (OR <1). While our lack of power is a limitation of this data, it is unlikely that prospective studies done in the current era of low MTCT rates could uncover a more precise or smaller association without requiring a prohibitively large cohort.

In conclusion, we detected a high prevalence of MG in pregnant HIV-infected Kenyan women and did not detect an association between MG and MTCT. A recent technical consultation convened to study MG outlined research gaps in our understanding of the public health implications of MG infection in pregnancy. Considering the high prevalence of MG in our population of pregnant women, we concur with this expert panel that additional research to determine the long term effects of maternal MG and the best treatment options in pregnancy should be a research priority.<sup>[41]</sup>

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Study design: ACR, KY, BAR, GJS

Parent cohort data collection: CF, RB, DMN, BAR, GJS

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**Table 1:**

Characteristics of a random sample of pregnant, HIV-seropositive women in Nairobi, Kenya, 1999-2005.

| Characteristic (n=220)   | N   | Median (IQR) or N (%) |
|--|-----|-----------------------|
| <b>Baseline (32 weeks gestation)</b>   |     |                       |
| Age (years)  | 220 | 24.5 (22-27)          |
| Education (years)  | 218 | 8 (7-12)              |
| Single   | 220 | 12 (5.5%)             |
| Employed   | 220 | 160 (72.7%)           |
| Age at first intercourse (years)   | 220 | 17 (16-19)            |
| Number of lifetime sex partners  | 220 | 3 (2-4)               |
| Parity   | 217 | 1 (1-2)               |
| History of tuberculosis  | 220 | 16 (17.3%)            |
| History of syphilis  | 220 | 20 (9.1%)             |
| History of STIs  | 220 | 28 (12.7%)            |
| History of GUD   | 220 | 13 (5.9)              |
| HSV-2 positive   | 137 | 121 (88.3)            |
| CD4 count (cells/mm <sup>3</sup> )   | 216 | 465 (310-643)         |
| HIV RNA (log <sub>10</sub> copies/ml)  | 210 | 4.68 (0.89)*          |
| <b>Pregnancy</b>   |     |                       |
| Diagnosed with tuberculosis  | 219 | 2 (0.9%)              |
| Diagnosed with syphilis  | 217 | 7 (3.2%)              |
| Diagnosed with genital ulcer disease   | 219 | 9 (4.1%)              |
| Diagnosed with bacterial vaginosis   | 209 | 80 (38.3%)            |
| Diagnosed with trichomoniasis  | 220 | 41 (18.6%)            |
| Diagnosed with gonorrhoea  | 219 | 2 (0.9%)              |
| Diagnosed with <i>Chlamydia</i>  | 219 | 10 (4.6%)             |
| Delivery vaginal HIV RNA (log <sub>10</sub> copies/ml)                       | 150 | 2.11 (1.70 – 3.13)    |
| Delivery plasma HIV RNA (log <sub>10</sub> copies/ml)                        | 169 | 4.15 (3.52-4.74)      |
| <i>M. genitalium</i> detected at antenatal visit                             | 220 | 47 (21.4%)            |
| <b>Infant characteristics</b>  |     |                       |
| All perinatal HIV: HIV detected at or before 4 weeks                         | 220 | 33 (15%)              |
| <i>In utero</i> transmission: HIV detected at delivery                       | 220 | 16 (7.3%)             |
| Intrapartum transmission: HIV not detected at birth, but detected at 4 weeks | 220 | 17 (7.7%)             |

IQR = inter-quartile range; STI = sexually transmitted infections; GUD = genital ulcer disease; HSV-2 = herpes simplex virus 2.

**Table 2:**

Correlates of *Mycoplasma genitalium* infection from random sample (n=220) of HIV-seropositive pregnant women.

| Characteristic                                      | N   | Women without <i>M. genitalium</i> infection (N=173) Mean (SD) or N (%) | Women with <i>M. genitalium</i> infection (N=47) Mean (SD) or N (%) | p-value* |
|---|-----|---|---|----------|
| <b>Baseline (32 weeks gestation)</b>                |     |   |   |          |
| Age (years)   | 220 | 25 (4.4)  | 25 (3.7)  | 0.70     |
| Education (years)                                   | 218 | 8.8 (2.7)   | 8.9 (2.9)   | 0.86     |
| Single  | 220 | 7 (4.1%)  | 5 (10.6%)   | 0.08     |
| Employed  | 220 | 127 (73.4%)   | 33 (70.2%)  | 0.66     |
| Age at first intercourse (years)                    | 220 | 17.4 (2.4)  | 17.3 (2.5)  | 0.81     |
| Lifetime sexual partners                            | 220 | 3 (1.6)   | 3 (1.5)   | 0.86     |
| Parity  | 217 | 1.5 (1.3)   | 1.6 (1.3)   | 0.69     |
| History of tuberculosis                             | 220 | 15 (8.7%)   | 1 (2.1%)  | 0.13     |
| History of syphilis                                 | 220 | 18 (10.4%)  | 2 (4.3%)  | 0.19     |
| History of any sexually transmitted infection (STI) | 220 | 26 (15%)  | 2 (4.3%)  | 0.05     |
| History of genital ulcer disease                    | 220 | 13 (7.5%)   | 0   | 0.05     |
| HSV-2 positive                                      | 137 | 91 (87.5%)  | 30 (91%)  | 0.60     |
| CD4 count (cells/mm <sup>3</sup> )                  | 216 | 479 (236)   | 522 (258)   | 0.29     |
| Plasma HIV RNA (log <sub>10</sub> copies/ml)        | 210 | 4.6 (0.92)  | 5.0 (0.73)  | 0.02     |
| <b>Pregnancy</b>                                    |     |   |   |          |
| Diagnosed with tuberculosis                         | 219 | 2 (1.2%)  | 0   | 0.46     |
| Diagnosed with syphilis                             | 217 | 6 (3.5%)  | 1 (2.1%)  | 0.63     |
| Diagnosed with genital ulcers                       | 219 | 7 (4.1%)  | 2 (4.3%)  | 0.96     |
| Diagnosed with BV                                   | 209 | 61 (37%)  | 19 (43.2%)  | 0.45     |
| Diagnosed with trichomoniasis                       | 220 | 28 (16.2%)  | 13 (27.7%)  | 0.07     |
| Diagnosed with gonorrhoea                           | 219 | 2 (1.2%)  | 0   | 0.46     |
| Diagnosed with chlamydia                            | 219 | 8 (4.7%)  | 2 (4.3%)  | 0.91     |
| <b>Delivery</b>                                     |     |   |   |          |
| Plasma HIV RNA (log <sub>10</sub> copies/ml)        | 169 | 4.0 (1.0)   | 4.3 (0.78)  | 0.16     |
| Vaginal HIV RNA (log <sub>10</sub> copies/ml)       | 150 | 2.4 (1.1)   | 2.6 (1)   | 0.45     |

Chi squared test or Fisher's exact test were used to compare categorical variables; t-tests were used to compare continuous variables. HIV RNA testing of plasma (n=169) and vaginal swabs (n=150) was not available at delivery for all mothers; similarly HSV-2 testing was not available for all mothers (n=137).

**Table 3:**

*Mycoplasma genitalium* and odds of perinatal, *in utero*, and intrapartum mother to child transmission of HIV

| Type of transmission             | Characteristic                              | Unadjusted OR (95%CI) | Adjusted OR (95%CI) |
|----------------------------------|---|-----------------------|---------------------|
| All perinatal HIV transmission   | Maternal MG infection                       | 0.72 (0.36 – 1.51)    | 0.55 (0.25 – 1.22)  |
|                                  | Plasma HIV RNA, log <sub>10</sub> copies/ml | 2.27 (1.47 – 3.50)    | 2.57 (1.62 – 4.06)  |
|                                  | Maternal BV                                 | 2.14 (1.20 – 3.81)    | 2.33 (1.26 – 4.34)  |
|                                  | Maternal CD4, cells/mm <sup>3</sup>         | 0.99 (0.99 – 1.00)    | -                   |
| <i>In utero</i> HIV transmission | Maternal MG infection                       | 0.63 (0.20 – 1.92)    | 0.49 (0.15 – 1.57)  |
|                                  | Plasma HIV RNA, log <sub>10</sub> copies/ml | 1.91 (1.07 – 3.42)    | 2.07 (1.13 – 3.80)  |
|                                  | Maternal BV                                 | 3.05 (1.33 – 6.98)    | 3.24 (1.38 – 7.58)  |
|                                  | Maternal CD4, cells/mm <sup>3</sup>         | 1.00 (0.99 – 1.00)    | -                   |
| Intrapartum HIV transmission     | Maternal MG infection                       | 0.80 (0.33 – 1.96)    | 0.63 (0.26 – 1.60)  |
|                                  | Plasma HIV RNA, log <sub>10</sub> copies/ml | 2.21 (1.31 – 3.72)    | 2.29 (1.35 – 3.86)  |
|                                  | Maternal BV                                 | 1.58 (0.78 – 3.21)    | 1.68 (0.80 – 3.56)  |
|                                  | Maternal CD4, cells/mm <sup>3</sup>         | 0.99 (0.99 – 1.00)    | -                   |

MG = *Mycoplasma genitalium* infection; BV = bacterial vaginosis, defined as Nugent score  $\geq 7$ ; OR = odds ratio; Perinatal HIV transmission is defined as any infant with HIV detected by 4 weeks of life; *In utero* HIV transmission is defined as HIV detected at birth in infants; intrapartum HIV transmission is defined as HIV not detected at birth, and detected at 4 weeks postpartum.

HIV plasma RNA level in log<sub>10</sub> copies/ml was measured at 28 weeks gestation.

Infants in whom *in utero* HIV transmission was detected at birth were excluded from the intrapartum HIV analysis.

Adjusted models include the following numbers of infants: Perinatal adjusted model, N=232, *in utero* adjusted model, N=232, intrapartum adjusted model, N=201.