

## Pathogenic lower genital tract organisms in HIV-infected and uninfected women, and their association with postpartum infectious morbidity

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**Objectives.** To determine the prevalence of vaginal pathogens during pregnancy and their impact on postpartum infectious morbidity among antiretroviral-naïve HIV-infected, and HIV-uninfected, women.

**Methods.** Vaginal swabs were obtained during early labour by speculum examination prior to digital vaginal examination, and sent for microscopy and culture. Women were assessed for infectious complications within 24 - 72 hours of delivery, and up to 2 weeks postpartum.

**Results.** Laboratory results were available for 801 women who delivered vaginally (418 HIV infected and 383 uninfected). The baseline characteristics of the two groups were comparable, and the median CD4 count for HIV-infected women ( $N=391$ ) was 416/ $\mu$ l. Fifty-five per cent (54.8%) of women had positive cultures

(439/801), more among those who were HIV infected than uninfected (60% v. 49.1%,  $p=0.002$ ). Women with positive cultures had slightly higher rates of infectious morbidity than those without (20.5% v. 15.2%,  $p=0.052$ ). *Trichomonas vaginalis* and group B streptococci were significantly associated with sepsis ( $p=0.023$  and  $<0.001$ , respectively), whereas the presence of *Candida* species seemed to be protective (relative risk 0.69,  $p=0.014$ ).

**Conclusion.** The study shows that a high proportion of pregnant women have pathogenic organisms in the lower genital tract that are associated with development of postpartum infectious morbidity.

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A balanced colonisation by endogenous vaginal microflora of the lower female genital tract consists of predominantly *Lactobacillus* species,<sup>1</sup> co-existing with vaginal anaerobes and aerobes such as *Streptococcus agalactiae* (e.g. group B streptococci)<sup>2</sup> and the

*Bacteroides* family.<sup>3</sup> Disturbing this microbial milieu results in overgrowth of bacterial vaginosis-related organisms which are associated with a 3 - 6 times greater risk of postpartum endometritis.<sup>4,5</sup> However, the role of endogenous lower genital tract pathogens, especially bacterial vaginosis organisms, and their impact on postpartum infectious morbidity, is controversial.<sup>6,7</sup> Other vaginal organisms associated with postpartum infectious morbidity include *Ureaplasma urealyticum*, group B streptococcus (GBS) and other sexually transmitted infections (STIs) such as *Trichomonas vaginalis*, *Neisseria gonorrhoeae* and *Chlamydia trachomatis*. Case detection and treatment of the latter STIs is often technically difficult and expensive. Furthermore, the prevalence of genital tract infections is probably higher among HIV-positive women because of their associated immunosuppression. Studies have shown perturbations in the vaginal flora of HIV-infected women, with approximately 20% of HIV-infected women having *Lactobacillus*-dominant vaginal flora.<sup>8,9</sup> Cu-Uvin *et al.* found no difference in the prevalence rates of

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lower genital tract infections among HIV-infected women compared with HIV-uninfected women.<sup>10</sup> These conflicting findings led us to determine the prevalence of lower genital tract infections during pregnancy and their impact on postpartum infectious morbidity among HIV-infected and uninfected women.

## Methods

### Study population

Our study was nested within a prospective trial of 1 404 women (716 HIV positive compared with 688 HIV negative) over a 2-year period.<sup>11,12</sup> Women of 18 years and older, and who were at 36 weeks' gestation and greater, in whom vaginal delivery was anticipated, were enrolled after obtaining consent. In addition to standard antenatal tests, pre- and post-test HIV counselling and HIV rapid testing were offered to all the women. CD4 lymphocyte counts were performed in the HIV-infected women. Vaginal swab specimens were obtained during early labour by speculum examination prior to digital vaginal examination. Since the parent study involved randomising HIV-infected women to receive either a *stat* dose of cefoxitin or placebo, the specimens were taken before the administration of the study drug. The vaginal swab specimens were rolled onto a glass slide for the determination of a bacterial vaginosis (BV) score using Nugent's criteria,<sup>13</sup> and then transported in Ames medium (for detection of Gram-negative and positive organisms) and Diamond's medium (for detection of trichomoniasis). In the laboratory, the swabs were also inoculated on cefoxitin CLED plates, bacitracin choc plate and laked blood cefoxitin, for the detection of any significant isolates, and an AMS plate was inoculated to detect the presence of any antimicrobial substances. Significant isolates were identified, and susceptibilities were determined. Screening tests for syphilis (rapid plasma reagin) were also performed, and presence or absence of vulvo-vaginal warts was noted during speculum examination. Swab specimens for *N. gonorrhoeae* and *Chlamydia* species were not taken.

Demographic details and obstetric information were obtained, including: the number of vaginal examinations before and after the rupture of membranes, duration of ruptured membranes, and mode of delivery. All deliveries were conducted by professional midwives or doctors. Women were assessed for infectious complications within 24 - 72 hours of delivery, at 1 week (range 6 - 10 days), 2 weeks (range 11 - 17 days) and 6 weeks (40 - 49 days) postpartum. Infection occurring

at any time point up to 2 weeks was recorded as a positive result. Only participants who had all microbiological laboratory results available were analysed.

### Statistical analysis

Data were captured on Epi-Info 6 and imported for analyses into SPSS version 15.0 (SPSS Inc., Chicago, Ill.). A *p*-value <0.05 was considered statistically significant. Pearson's chi-square analysis was used to compare categorical variables between groups. Quantitative variables were compared between treatment groups using independent *t*-tests or non-parametric Mann-Whitney tests as appropriate. Generalised linear models estimated crude and adjusted relative risks. The dependent variable was modelled as binary with a log link to estimate relative risks. For multivariable analysis, the independent variables were chosen as those which were statistically significant at the 90% level of confidence on crude analysis. They were entered as main effects. Relative risks (RR) and 95% confidence intervals (CI) were reported. Since HIV status and mode of delivery were possible confounders, these were entered into the model, and adjusted relative risks were calculated.

## Results

Laboratory results were available for 801 (418 HIV infected and 383 uninfected) of the 885 who had vaginal deliveries (including episiotomy and assisted deliveries). Baseline characteristics of the two groups were comparable, with no statistical differences in mean age, parity and body weight. The median CD4 count for HIV-infected women (*N*=391) was 416/μl (range 24 - 1 393), with 49/391 (12.5%) having CD counts <200/μl.

Overall, 439/801 (54.8%) had positive cultures, which was significantly more among HIV-infected women (251/418 (60%)) than HIV uninfected (188/383 (49.1%), *p*=0.002). *T. vaginalis*, BV and *Candida* species were significantly more prevalent among HIV-infected than uninfected women, as was vulvo-vaginal warts (Table I). Of these organisms, only *T. vaginalis* was significantly associated with presence of vaginal discharge on enquiry (*p*<0.001) (the *p*-value was 0.391 for BV and 0.080 for *Candida* species). Other organisms that were cultured included streptococci (groups B and F), as well as Gram-negative bacilli, none of which was significantly different among the HIV-infected and uninfected groups. Syphilis serology

Table I. Vaginal pathogens according to HIV status

	HIV status				RR (95% CI)	<i>p</i> -value
	Positive		Negative			
	Count	%	Count	%		
Positive cultures	251	60.0	188	49.1	1.22 (1.08 - 1.39)	0.002
Bacterial vaginosis	161	38.5	109	28.5	1.35 (1.11 - 1.65)	0.003
<i>T. vaginalis</i>	56	13.4	30	7.8	1.29 (1.08 - 1.53)	0.011
Group B streptococci	10	2.4	10	2.6	0.92 (0.39 - 2.18)	0.843
<i>Candida</i>	164	39.2	115	30	1.31 (1.08 - 1.59)	0.006
Group F streptococci	6	1.4	4	1.0	1.37 (0.39 - 4.83)	0.619
Gram-negative bacilli	104	24.9	91	23.8	1.05 (0.82 - 1.34)	0.712
Syphilis serology	26	6.5	14	3.7	1.74 (0.92 - 3.28)	0.082
Vulvo-vaginal warts	14	3.3	4	1.0	3.21 (1.06 - 9.66)	0.028

RR = relative risk for presence of pathogens in HIV-positive compared with HIV-negative women.

reactivity was also slightly more among HIV-infected women than among those who were uninfected ( $p=0.082$ ).

**Association between vaginal organisms and development of postpartum infectious complications**

The overall puerperal sepsis rate in this cohort was 18.1% (145/801) over the 2 weeks of follow-up. Women with positive cultures had higher rates of infectious morbidity than those without (90/439 (20.5%) v. 55/362 (15.2%),  $p=0.052$ ).

HIV status was not associated with development of sepsis ( $p=0.391$ ) in crude or multivariable analysis and was therefore omitted from the multivariable model. Mode of delivery, *Candida*, *T. vaginalis*, Gram-negative bacilli and group B streptococci were added to the multivariable model as independent variables. After adjustment for confounding effects, assisted vaginal delivery or episiotomy posed a significantly higher risk for sepsis than non-assisted normal vaginal delivery (NVD) ( $p=0.001$ ). The presence of *Candida* species seemed to be protective (RR 0.66, 95% CI 0.47 - 0.92) (Table II), while *T. vaginalis*, Gram-negative bacilli and group B streptococci were all significant risk factors for sepsis.

**Discussion**

In this cohort, more than 50% of women had culture-positive pathogenic organisms detected during early labour, which represents

a very high rate of microbial pathogens in the lower genital tract of pregnant women who were asymptomatic, with the exception of a possible vaginal discharge. However, similar rates have been reported.<sup>14,15</sup> In a similar population, one or more STIs was found to be present in 16.5% of asymptomatic antenatal patients.<sup>15</sup> Though not statistically significant, postpartum infectious morbidity was higher in women with positive cultures ( $p=0.052$ ). Furthermore, in our study, positive cultures were significantly higher among HIV-infected (60%) than HIV-uninfected women (49.1%,  $p=0.002$ ). The difference according to HIV status was expected and is in keeping with other reports of increased rates of lower genital tract pathogens among HIV-infected women.<sup>16</sup>

A higher incidence of infectious morbidity among HIV-infected women was expected in view of their higher rate of positive cultures. The lack of influence of HIV status on the development of postpartum infectious morbidity in our study has also been noted in other reports;<sup>11,12</sup> this is probably because most HIV-infected women in our study were immunocompetent, with mean CD4 counts >350.

The most common organisms detected in our study were associated with BV, with rates in keeping with other sub-Saharan Africa rates of up to 50%.<sup>8,17,18</sup> Although organisms associated with BV were the most prevalent, there was surprisingly no association between presence of BV and postpartum infectious morbidity. The vast literature showing an association of BV with pregnancy complications has mainly focused on preterm labour and delivery,

**Table II. Association between development of infectious morbidity and presence of organisms**

Parameter	Infectious morbidity: N (%)	p-value	Crude RR (95% CI)	Adjusted RR* (95% CI)	p-value
<b>HIV</b>					
Positive	71 (17.0)	0.391	1.14 (0.85 - 1.53)		
Negative	74 (19.3)				
<b>Mode of delivery</b>					
Assisted/episiotomy	88 (23.3)	<0.001	1.74 (1.28 - 2.35)	1.71 (1.26 - 2.31)	0.001
Normal delivery	57 (13.4)				
<b>Syphilis</b>					
Positive	6 (15)	0.568	0.81 (0.38 - 1.71)		
Negative	137 (18.6)				
<b>Vaginal warts</b>					
Positive	4 (22.2)	0.550	1.23 (0.51 - 2.97)		
Negative	141 (18)				
<b>Candida</b>					
Positive	39 (14)	0.027	0.69 (0.49 - 0.96)	0.66 (0.47 - 0.92)	0.014
Negative	106 (20.3)				
<b>Bacterial vaginosis</b>					
Positive	47 (17.4)	0.716	0.94 (0.69 - 1.29)		
Negative	98 (18.5)				
<b>Group F strep.</b>					
Positive	3 (30)	0.399	1.67 (0.64 - 4.36)		
Negative	142 (18)				
<b>T. vaginalis</b>					
Positive	26 (30.2)	0.002	1.82 (1.27 - 2.61)	1.49 (1.06 - 2.11)	0.023
Negative	119 (16.6)				
<b>Gram-negative bacilli</b>					
Positive	44 (22.6)	0.063	1.35 (0.99 - 1.86)	1.48 (1.09 - 2.01)	0.013
Negative	101 (16.7)				
<b>Group B strep.</b>					
Positive	11 (55)	<0.001	3.21 (2.10 - 4.90)	2.62 (1.76 - 3.91)	<0.001
Negative	134 (17.2)				

\*Adjusted for mode of delivery, *Candida*, *T. vaginalis*, Gram-negative bacilli, and group B Streptococci.

and subsequently postpartum endometritis.<sup>19</sup> We postulate that the postpartum endometritis in those studies is a continuation of the preceding inflammatory process that led to miscarriage or preterm labour, and asymptomatic women who proceed to term are less likely to experience BV-associated complications in the postpartum period. This observation could explain the lack of association between the presence of BV and postpartum infectious morbidity in our cohort of otherwise asymptomatic women, recruited at  $\geq 36$  weeks' gestation.

We showed a significant association of postpartum infectious morbidity and presence of *T. vaginalis*. This organism colonises the lower genital tract, and is therefore not expected to be associated with upper genital disease (above the level of the internal cervical os). *T. vaginalis* has been documented in the upper genitalia and is associated with pelvic inflammatory disease in those who are HIV infected,<sup>20</sup> but not among those without HIV infection. Although small numbers were *T. vaginalis* positive, our study also found a significant association between *T. vaginalis* and postpartum infections in the whole cohort ( $p=0.010$ ), and particularly among HIV-infected women (relative risk (RR) 3.36,  $p=0.009$ ) rather than HIV-uninfected women (RR 1.29,  $p=0.599$ ). This association between HIV and *T. vaginalis*-associated morbidity may suggest an altered host-microbe relationship, resulting in an apparent increased risk of upper genital tract infection.<sup>20</sup> Since the parent study involved randomising HIV-infected women to a single dose of 2 grams of cefoxitin and placebo, cefoxitin administration was found to significantly reduce the development of infectious morbidity in women who had *T. vaginalis* (RR 0.18, 95% confidence interval 0.03 - 1.22,  $p=0.024$ ). Cefoxitin has little effect on anaerobes such as *T. vaginalis*, and it therefore raises the possibility that its effect is more on the 'secondary invaders' that cause infection in mucosa that is already compromised by the presence of *T. vaginalis*. This relationship is well known in the development of pelvic inflammatory disease in non-pregnant women.<sup>21</sup>

As expected, the presence of GBS was strongly associated with development of postpartum infections. However, the prevalence rate of GBS in our study was much lower than the 15% prevalence rate reported previously from the same institution.<sup>22</sup> This may be due to swabs only being taken from the cervico-vaginal area in this study, whereas Naidoo *et al.* noted that a combination of vaginal and introital swabs increased the yield for GBS by 25%.<sup>22</sup>

*Candida* species were present in at least 30% of our cohort; however, most women were asymptomatic. A surprising finding in our study was that the presence of *Candida* organisms protected against the development of postpartum infectious morbidity, which persisted even after controlling for other variables. Further studies are needed to confirm this.

In conclusion, HIV-positive women did not have an increased risk of postpartum infectious morbidity despite higher rates of

pathogens in the lower genital tract, a finding which may be due to the reasonably intact immune system of the study cohort. However, the presence of *T. vaginalis*, Gram-negative bacilli and group B streptococci was associated with a significantly increased risk of postpartum morbidity, irrespective of HIV status.

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