Strong association between higher-risk sex and HIV prevalence at the regional level: an ecological study of 27 sub-Saharan African countries [version 1; peer review: 2 approved]

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Abstract

Background: It is unclear why HIV prevalence varies by nearly two orders of magnitude between regions within countries in sub-Saharan Africa. In this ecological study, we assess if HIV prevalence by region is associated with any of four markers of higher risk sexual behavior: lifetime number of partners, multiple partners in past year, higher risk sex (defined as sex with non-cohabiting, non-marital partners) and age at debut.

Methods: We performed Pearson’s correlation between the 4 behavioral risk factors and HIV prevalence by region in 47 nationally representative surveys from 27 sub-Saharan African countries, separately by gender. In addition, principal components analysis was used to reduce the eight risk factors (four for each gender) to two principal components (PCs). Mixed effects linear regression was used to assess the relationship between the resulting two PCs and HIV prevalence after controlling for the prevalence of male circumcision.

Results: HIV prevalence varied by a median 3.7 fold (IQR 2.9-7.9) between regions within countries. HIV prevalence was strongly associated with higher risk sex and, to a lesser extent, the other risk factors evaluated. Both PCs were strongly associated with HIV prevalence when assessed via linear regression.

Conclusions: Differences in sexual behavior may underpin the large differences in HIV-prevalence between subpopulation within sub-Saharan African countries.

Keywords
HIV prevalence, ecological, sexual network, high risk sex, sub-Saharan Africa

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Approval Status ✓ ✓

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Abbreviations

DHS - Demographic Health Survey
PC - Principal components
PCA - Principal components analysis
STI - Sexually transmitted infection

Background

Whereas individual-level parameters may influence which individuals in a given population acquire infection, it is population-level parameters that affect the prevalence of infection.

Aral et al. (Aral et al., 2007)

Aral and others have argued persuasively that population-level parameters such as the structure of sexual networks determine the prevalence of sexually transmitted infections (STIs) (Aral et al., 2007; Morris et al., 2010; Morris et al., 2009). If this is true, then it follows that ecological studies are necessary to assess which markers of network structure are associated with higher HIV prevalence (Kenyon & Colebunders, 2012; Morris et al., 2010).

Whilst a few ecological studies have found a difference in sexual behavior could explain differences in HIV prevalence (De Walque, 2006; Kenyon et al., 2013; Kenyon & Colebunders, 2012; Morris et al., 2010; Morris et al., 2009), many have not (Auvert et al., 2001; Cleland & Ferry, 1995; Drain et al., 2004; Sawers & Stillwaggon, 2010; Wellings et al., 2006), and the meted cross-country comparisons; these are suboptimal because national populations may be constituted by several relatively separate sexual networks (Kenyon & Colebunders, 2013; Laumann & Youm, 1999). It would therefore be more appropriate to conduct these ecological assessments at the level of coherent sexual networks rather than national populations (Kenyon et al., 2015b). If there is both a high degree of sexual partner homophily by ethnic group, and large differences in HIV prevalence between ethnic groups within countries, then it would be appropriate to use ecological analyses to assess if differences in sexual network structure could explain the variations in HIV prevalence (Kenyon et al., 2013). This type of study from the USA, UK, Kenya, Uganda and South Africa has revealed positive ecological associations between various markers of sexual behavior (number of partners in past year and lifetime, partner concurrency and age of sexual debut), by ethnic group and HIV prevalence (Kenyon et al., 2014a; Kenyon et al., 2013; Kenyon et al., 2014b; Kenyon et al., 2014c; Kenyon et al., 2016b; Morris et al., 2009).

Results from our recent ecological analysis of factors associated with HIV prevalence in Ethiopia based on data from Demographic Health Surveys (DHS) implied that it may be more appropriate to evaluate differences by region than by ethnic group (Kenyon et al., 2015). For instance, 80 different ethnic groups have been grouped into 9 ethnically-based regions in Ethiopia’s DHS surveys (Central Statistical Agency [Ethiopian] & ICF International, 2012). Surveys from other sub-Saharan African countries also combine similar ethnic groups into regions and, as in Ethiopia, are designed to provide representative samples from each of these regions. We established that there was a high degree of sexual partner homophily by ethnic group region in Ethiopia (Kenyon et al., 2015b). There was also a high (seven-fold) difference in HIV prevalence between regions and this correlated closely at an ecological level with a number of behavioral variables such as lifetime number of partners, reporting sex with a non-marital, non-cohabiting partner, and age at first sex (Kenyon et al., 2015b). In this paper, we extend these earlier analyses to assess systematically if differences in HIV prevalence by region within countries in sub-Saharan Africa are associated with markers of sexual risk behavior.

Methods

Data source

Nationally representative HIV-serolinked DHS’s are available for 29 sub-Saharan African countries from MeasureDHS (http://www.measuredhs.com). We limited our analyses to the 47 surveys from 27 countries where the HIV prevalence varied between ro, 1/1 from Swaziland and 2/3 from Zimbabwe (2005 and 2010). The selected variables were downloaded in regional format for all HIV-serolinked surveys from included countries via the STATcompiler function of MeasureDHS.

Variables used in the analyses (number of surveys with missing data)

Behavioral risk factors:

Higher risk sex (0/47): Percentage of women/men who have had sex with a non-marital, non-cohabiting partner in the last 12 months of all women/men reporting sexual activity in the last 12 months

Multiple partners in past-year (0/47): Percentage of women/men aged 15–49 years who have had sexual intercourse with more than one partner in the last 12 months

Lifetime partners (11/47): Mean number of lifetime sexual partners among women/men who ever had sexual intercourse.

Women/men’s debut (5/47 for women and 30/47 for men): Median age at first sexual intercourse in years among women/men aged 20–49 years

Control variables:

Men circumcised (14/47): Percentage of men who report being circumcised.

Outcome variable:

HIV prevalence (0/47): Percentage of 15–49 year old men and women who tested positive for HIV.

These variables were selected for analysis based on associations with HIV from previous studies and coverage in a sufficient number of surveys (Kenyon et al., 2014a; Kenyon et al., 2013; Kenyon et al., 2015b; Mishra et al., 2009; Morris et al., 2009).

Statistical analysis

Our analysis used Pearson’s correlation to assess the association between lifetime partners, multiple partners in the past year, high-risk sex, sexual debut and HIV prevalence (15–49 years, men and women combined). The analyses were conducted by region within each survey and done separately for women...
and men. A Pearson’s $r \geq 0.3$ or $\leq -0.3$ was considered positive or negative, respectively. For each risk factor, we also compared the fold-difference in the prevalence of the risk factor between the regions with the highest and lowest HIV prevalence in each survey.

**Multivariable analysis.** Pearson’s correlation was used to assess the associations between the four behavioral risk factors separately between women and men. Principal components analysis (PCA) was used to reduce seven of the eight risk factors to two principal components. Sexual debut of men was not included in this process as data was missing for 30/47 surveys.

Mixed effects linear regression with a random intercept for individual surveys was then used to assess the relationship between HIV prevalence and these two principal components, controlling for prevalence of circumcision. The following sensitivity analyses were conducted. 1. Analyses stratified by gender. 2. Using principal components calculated excluding the lifetime partners of women and men variables (as these were missing in 11 surveys).

The study was conceived as being exploratory rather than hypothesis testing. Because of this and the fact that the sample sizes in each survey were relatively small, we did not use Bonferroni corrections (Armstrong, 2014; Perneger, 1998; Rothman, 1990). A P-value below 0.05 was regarded as statistically significant. The analyses were conducted using STATA 13.0 (College Station, TX). Each DHS received ethical committee clearance for data analyses such as the one performed here. Consequently, no specific ethics committee approval was necessary for this study.

**Results**

A total of 47 HIV-serolinked surveys from 27 countries were included. This sample included countries with a wide range of national HIV prevalences in 15–49 year olds, ranging from 0.4% (95% confidence interval [CI] 0.2-0.5%) in Niger 2012 to 13.8 (95% CI, 12.9-14.8%) in Zimbabwe 2015. The HIV prevalence differed a median 3.8-fold (interquartile range [IQR] 2.9-8.5) between regions within surveys (Table 1).

**Higher-risk sex**

The prevalence of higher-risk sex between regions within surveys varied widely in women (median 4.8-fold difference [IQR 2.3-8.1]) and men (median 2.5-fold difference [IQR 1.8-5.5]).

Women: There was a positive association between higher risk sex and HIV in 39/47 surveys, of which 26 were statistically significant. There were no surveys where the association was negative (Table 1; Figure 1). Within surveys, the region with the highest HIV prevalence had a median 2.5 (IQR 1.4 to 6.5) times increased prevalence of higher-risk sex for women than the lowest HIV prevalence population.

Men: The association was positive in 38/47 surveys, 22 of which were significant, and negative in none of the surveys. In each survey, the region with the highest HIV prevalence had a median 1.9 (IQR 1.2 to 2.9) times higher prevalence of higher risk sex for men than the lowest HIV prevalence population.

**Number of lifetime sex partners**

The reported number of lifetime partners between regions in each survey varied by a median 1.8-fold (IQR 1.5-2.4) for women and 2.5-fold (2.0-3.3) for men.

**Women:** There was a positive association between number of lifetime partners and HIV prevalence in 18/36 surveys, of which 9 were statistically significant (Table 1, Figure 2). Of the three surveys for which the association was negative, this was statistically significant for one (Namibia, 2013). Within surveys, the region with the highest HIV prevalence had a median 1.3 (IQR 1.0 to 1.5) times higher prevalence of lifetime partners for women than the lowest HIV prevalence population.

**Men:** The association was positive in 22/36 surveys (significant in 13 surveys). In the three surveys where the association was negative, this was only significant in the case of Namibia 2013. Within each survey, the region with the highest HIV prevalence had a median 1.5 (IQR 1.1 to 2.1) times higher prevalence of lifetime partners for men than the lowest HIV prevalence population.

**Multiple partners in past-year**

The percent with multiple partners in the past year varied between regions by a median 6.5-fold (IQR 3.5-13.7) for women and 2.8-fold (IQR 2.0-4.8) for men.

**Women:** There was evidence of a positive association in 30/47 surveys but this association was only significant in five surveys (Table 1, Figure 3). The association was negative in three surveys, one of which was significant (Namibia, 2013). In each survey, the region with the highest HIV prevalence had a median 2.1 (IQR 1.4 to 4.1) times higher prevalence of multiple partners for women than the lowest HIV prevalence population.

**Men:** The association was positive in 21/47 surveys, of which four were statistically significant and negative in 10 surveys, of which one was significant – Chad 2014. Within surveys, the region with the highest HIV prevalence had a median 1.3 (IQR 1.0 to 2.1) times higher prevalence of multiple partner for men than the lowest HIV prevalence population.

**Men’s and women’s debut**

**Women:** There was evidence of a positive association between women’s age at sexual debut and HIV prevalence in 22/42 surveys, but this association was only significant in three surveys (Chad, 2014; Guinea, 2012; Niger, 2012; Table 1, Figure 4). The association was negative in six surveys, of which this was significant in two (Kenya, 2008; Senegal, 2010).

**Men:** The association was positive in 1/17 surveys, which was statistically significant (Gabon, 2012) and negative in three surveys, none of which was significant.

**Associations between behavioral risk factors**

Within both women and men, the three behavioral risk factors (lifetime partners, past year partners and high-risk sex) were
Table 1. Correlations between HIV prevalence and risk factors in 47 sub-Saharan African Demographic Health Surveys.

<table>
<thead>
<tr>
<th>Country</th>
<th>Year</th>
<th>N regions</th>
<th>HIV ratio*</th>
<th>Women</th>
<th>Men</th>
</tr>
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<td></td>
<td></td>
<td></td>
<td></td>
<td>Sexual Debut</td>
<td>Higher risk Sex</td>
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<td>42</td>
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<td>0.56</td>
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</tr>
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<td>-0.19</td>
<td>-0.09</td>
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positively associated with one another (Table 2). Within men (but not women) these three risk factors were negatively associated with age at first sex.

**Multivariable analyses**

*Principal components:* Two principal components were able to explain 80% of the variation in the seven variables. The component loadings of the principal components are depicted in Figure 5. Principal component one (PC1) represented a summation of the variables high-risk sex (women and men), lifetime partners (women and men) and multiple partners (women and men), minus the variable debut age women (Figure 5).

In the unadjusted model, PC1 and PC2 were associated with HIV prevalence (coeff. 0.51, 95% CI 0.31-0.71, and coeff. 0.72, 95% CI 0.38-1.05, both P<0.005, respectively; Table 3). Adjusting for the prevalence of circumcision made little difference to the strength of the association (Table 3).

Sensitivity analyses repeating the analyses stratified by gender, using the principal components that were calculated excluding the lifetime partners (women and men) variables in the analyses made little difference to the results (Table 3).

**Discussion**

We found wide variations in the prevalence of HIV and the four behavioral risk factors evaluated. All the risk factors, excluding women’s men’s debut, were positively associated with one another and with HIV prevalence. Combining the four risk factors via principal components analysis generated two components which were strongly associated with HIV prevalence.

There are a number of interpretations of and limitations to our results. Demographic and Health Surveys are not designed to accurately assess sensitive information such as sexual behavior (Beaclair et al., 2013; Morris et al., 2014). As such, they have been shown to underestimate the prevalence of behaviors that may be particularly sensitive to a respondent bias, such as number of partners (Glick & Sahn, 2007; Kenyon et al., 2013; Morris et al., 2014). There is, however, no evidence that we could find that this bias would operate differentially by ethnic or regional group (Johnson et al., 2009; Kenyon et al., 2014a). This bias should thus result in underestimated the prevalence of reported risk behavior, but this effect should not differ by region. We did not control for a broad range of variables that may confound our results. Age of respondents, which is known to affect HIV prevalence and sexual behaviors such as lifetime partner number, was not controlled for. However, the DHS sampling strategy typically produces populations that do not differ by age between regions (Central Statistical Agency [Ethiopia] & ICF International, 2012; De Walque, 2006; Kenyon et al., 2013; Kenyon et al., 2015). We also decided not to control for condom usage or upstream determinants such as socioeconomic status. In keeping with a number of other African studies (Kenyon et al., 2014a; Kenyon et al., 2013; Kenyon et al., 2015b), we found that condom usage tended to be higher in areas more affected by HIV (data not shown). As a result, we did not control for condom usage.

The relationship between socioeconomic status and HIV is complex, with most evidence pointing to a positive association between wealth and HIV in sub-Saharan Africa (Hajizadeh et al., 2014; Mishra et al., 2009; Susser, 1994). We found the same to be true at regional level (data not shown). Because our conceptual framework (Figure 6) conceived the upstream determinants as operating via sexual behavior, we considered it would be inappropriate to control for these in our model. The surveys were done at different stages in the HIV epidemics of

<table>
<thead>
<tr>
<th>Country</th>
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<th>Higher risk Sex</th>
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<th>Multiple partners past year</th>
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<td>Zambia</td>
<td>2013</td>
<td>10</td>
<td>2.8</td>
<td>0.38</td>
<td>0.50</td>
<td>0.38</td>
<td>0.68</td>
<td>0.12</td>
<td>0.64</td>
<td>0.19</td>
<td>0.06</td>
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<tr>
<td>Zimbabwe</td>
<td>2015</td>
<td>10</td>
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<td>-0.23</td>
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<td>0.80</td>
<td>0.67</td>
<td>0.69</td>
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<tr>
<td>Median (IQR)</td>
<td>10</td>
<td>2</td>
<td>3.8</td>
<td>(0.38-8.5)</td>
<td>0.61</td>
<td>(0.38-0.79)</td>
<td>0.31</td>
<td>(0.07-0.62)</td>
<td>-0.08</td>
<td>(-0.26-0.12)</td>
<td>0.63</td>
<td>(0.35-0.80)</td>
</tr>
</tbody>
</table>

*HIV ratio: HIV prevalence in highest HIV prevalence region/ HIV prevalence in lowest HIV prevalence region in each survey.

P<0.05 represented by bold font.
Figure 1. Scatterplots of HIV prevalence (15–49 year olds) versus high-risk sex behavior stratified by men/women and high/low HIV prevalence in 47 sub-Saharan African Demographic Health Surveys.
Figure 2. Scatterplots of HIV prevalence (15–49 year olds) versus lifetime number of sex partners stratified by men/women and high/low HIV prevalence in 47 sub Saharan African Demographic Health Surveys.
Figure 3. Scatterplots of HIV prevalence (15–49 year olds) versus multiple partners past year stratified by men/women and high/low HIV prevalence in 47 sub Saharan African Demographic Health Surveys.
Figure 4. Scatterplots of HIV prevalence (15–49 year olds) versus median age of sexual debut stratified by men/women and high/low HIV prevalence in 47 sub-Saharan African Demographic Health Surveys.
<table>
<thead>
<tr>
<th></th>
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<td>Burkina Faso</td>
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<td>11 (1)</td>
<td>12 (1)</td>
<td>8 (1)</td>
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<td>8 (1)</td>
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<td>7 (4)</td>
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<td>7 (4)</td>
<td>8 (4)</td>
<td>7 (4)</td>
</tr>
<tr>
<td>Cote d'Ivoire</td>
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<td>12 (5)</td>
<td>10 (4)</td>
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<td>12 (5)</td>
<td>13 (5)</td>
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<td>6 (4)</td>
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<td>12 (4)</td>
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<td>9 (4)</td>
<td>9 (4)</td>
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<tr>
<td>Nigeria</td>
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<td>9 (4)</td>
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<td>9 (4)</td>
</tr>
<tr>
<td>Senegal</td>
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<td>9 (4)</td>
<td>9 (4)</td>
<td>9 (4)</td>
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<td>South Africa</td>
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<td>12 (4)</td>
<td>12 (4)</td>
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<td>12 (4)</td>
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<tr>
<td>Tanzania</td>
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<td>9 (4)</td>
<td>9 (4)</td>
<td>9 (4)</td>
<td>9 (4)</td>
<td>9 (4)</td>
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<td>9 (4)</td>
<td>9 (4)</td>
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</tbody>
</table>

Table 2. Correlations between behavioral risk factors in 47 sub-Saharan African Demographic Health Surveys.
### Correlations with HIV Prevalence

<table>
<thead>
<tr>
<th>Country</th>
<th>Year</th>
<th>HIV Prevalence</th>
<th>Debut vs Lifetime</th>
<th>Debut vs Multiple</th>
<th>Circumcision vs Multiple</th>
<th>Circumcision vs Past partners</th>
</tr>
</thead>
<tbody>
<tr>
<td>Mali</td>
<td>2006</td>
<td>10</td>
<td>0.33</td>
<td>0.73</td>
<td>0.47</td>
<td>0.26 (-0.31)</td>
</tr>
<tr>
<td></td>
<td>2009</td>
<td>11</td>
<td>0.53</td>
<td>0.22</td>
<td>0.60</td>
<td>0.50 (-0.36)</td>
</tr>
<tr>
<td>Namibia</td>
<td>2011</td>
<td>13</td>
<td>0.32</td>
<td>0.13</td>
<td>0.59</td>
<td>0.66 (0.35)</td>
</tr>
<tr>
<td>Niger</td>
<td>2012</td>
<td>8</td>
<td>0.57</td>
<td>0.42</td>
<td>0.36</td>
<td>0.21 (-0.17)</td>
</tr>
<tr>
<td>Rwanda</td>
<td>2014</td>
<td>10</td>
<td>0.35</td>
<td>0.05</td>
<td>0.95</td>
<td>0.89 (-0.47)</td>
</tr>
<tr>
<td>Senegal</td>
<td>2010</td>
<td>11</td>
<td>0.77</td>
<td>0.70</td>
<td>0.80</td>
<td>0.71 (-0.51)</td>
</tr>
<tr>
<td>Tanzania</td>
<td>2008</td>
<td>31</td>
<td>0.83</td>
<td>0.03</td>
<td>0.80</td>
<td>0.87 (-0.60)</td>
</tr>
<tr>
<td>Uganda</td>
<td>2014</td>
<td>9</td>
<td>0.85</td>
<td>0.43</td>
<td>0.90</td>
<td>0.94 (-0.18)</td>
</tr>
<tr>
<td>Zimbabwe</td>
<td>2001</td>
<td>9</td>
<td>0.87</td>
<td>0.63</td>
<td>0.68</td>
<td>0.83 (-0.39)</td>
</tr>
<tr>
<td>Zambia</td>
<td>2013</td>
<td>10</td>
<td>0.28</td>
<td>0.08</td>
<td>0.60</td>
<td>0.38 (-0.05)</td>
</tr>
<tr>
<td>Zimbabwe</td>
<td>2015</td>
<td>10</td>
<td>0.32</td>
<td>0.13</td>
<td>0.59</td>
<td>0.66 (0.35)</td>
</tr>
</tbody>
</table>

### Median (IQR)

<table>
<thead>
<tr>
<th>Country</th>
<th>Median (IQR)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Mali</td>
<td>3.8 (2.9-4.7)</td>
</tr>
<tr>
<td>Namibia</td>
<td>0.50 (0.40-0.60)</td>
</tr>
<tr>
<td>Niger</td>
<td>0.79 (0.60-0.80)</td>
</tr>
<tr>
<td>Rwanda</td>
<td>0.74 (0.60-0.80)</td>
</tr>
<tr>
<td>Senegal</td>
<td>0.21 (-0.05-0.40)</td>
</tr>
<tr>
<td>Tanzania</td>
<td>0.79 (0.60-0.80)</td>
</tr>
<tr>
<td>Uganda</td>
<td>0.74 (0.60-0.80)</td>
</tr>
<tr>
<td>Zimbabwe</td>
<td>0.79 (0.60-0.80)</td>
</tr>
<tr>
<td>Zambia</td>
<td>0.79 (0.60-0.80)</td>
</tr>
</tbody>
</table>

F1000Research 2018, 7:1879 Last updated: 06 JUN 2022
Figure 5. Loadings of the 7 behavioral risk factors in principal components 1 and 2.

Table 3. Mixed effects multiple regression with HIV-prevalence by region as outcome variable [Coefficient (95% CI)].

<table>
<thead>
<tr>
<th>PC</th>
<th>Including lifetime sex partners</th>
<th>Excluding lifetime sex partners</th>
</tr>
</thead>
<tbody>
<tr>
<td>PC1</td>
<td>0.51 (0.31-0.71)**</td>
<td>0.76 (.49-1.03)**</td>
</tr>
<tr>
<td>PC2</td>
<td>0.72 (0.38-1.05)**</td>
<td>0.58 (0.14-1.0)**</td>
</tr>
<tr>
<td>Circumcision</td>
<td>-</td>
<td>-0.09 (-.11-.07)**</td>
</tr>
</tbody>
</table>

*P<0.05, **P<0.005. PC, principal component.

the various countries. A wide range of studies have found that populations in sub-Saharan Africa have responded to the HIV epidemic by reducing a range of risk behaviors including casual partners and partner numbers (Glick & Sahn, 2007; Hajizadeh et al., 2014; Halperin et al., 2011; Kirby, 2008). By preferentially affecting persons with higher numbers of partners, AIDS mortality may also reduce the average number of partners reported in high prevalence settings (Chesson et al., 2003; Kenyon et al., 2016a). Not controlling for age of the epidemic, behavior change and AIDS mortalities, effect on partner number should, however, serve to dilute any relationship between sexual risk behaviors and HIV prevalence. We did not assess heterogeneity in HIV prevalence and high-risk behaviors within regions or the extent to which sexual networks were coterminous with regions within countries. Once again unconsidered heterogeneity would be expected to reduce the strength of the association between HIV and risk-behavior.

Because HIV prevalence is a product of a number of risk factors operating over decades, one would not expect to consistently find a linear relationship between HIV prevalence and any particular risk factor by region. For this reason, in addition to the Pearson’s correlations, we compared the prevalence of each of the risk factors between the lowest and highest HIV prevalence region in each survey. These provided results commensurate with those provided by linear regression. If non-participation in the surveys or the HIV-testing component were not random this could result in non-response biases. However, response rates in
Figure 6. Conceptual framework for understanding how risk factors could come together to produce sexual networks that are higher risk for HIV transmission in certain populations than others, using the example of Ethiopian regions. (A) Men in Gambela (HIV prevalence 7%) report a higher number of lifetime sex partners (both median and percent with more than 6 partners – vertical red line) than those in the Somali region (HIV prevalence 0.9%) (Reproduced from [14] with permission). (B) This in conjunction with other determinants of enhanced network connectivity will result in a larger reachable path for HIV in Gambela than Somali. The reachable path is represented by the edges between nodes in the sexual networks. (C) The prevalence of other risk factors such as condom usage and circumcision influence the probability of HIV transmission per sexual contact (probability of transmission is depicted as proportional to edge width between nodes). (D) The composite of A, B and C determine the sexual network transmission index, which combines the reachable path determined by A and B with the probability of transmission (C) and results in a higher HIV prevalence (red nodes) in Gambela than Somali (E). Within both Gambela and Somali there is a stepwise increase in HIV prevalence with increasing number of partners, but given the higher sexual network transmission index in Gambela, the probability of HIV is higher for someone with a low number of partners in Gambela than someone with a high number of partners in Somali. (F) represents the HIV prevalence by number of lifetime sex partners for men 15–49 years from all DHS countries with available data. Graphic reproduced from [20] with permission. Somali most closely approximates India and Gambela, Tanzania.
most of the surveys were high. This study involves the ecological analysis of cross sectional data. As such we can only describe associations found and not attribute causation.

Convincing evidence of a strong link between rate of partner change and risk of STIs including HIV has been provided by empirical individual level studies, modelling studies, and the theoretical importance of rate of partner change in the formula for the basic reproductive number (Auvert et al., 2001; Chen et al., 2007; Mishra et al., 2009; Xu et al., 2006). Women reporting sex with a non-marital, non-cohabiting person have also been shown to be more likely to be HIV positive in all of 18 DHS’s where this was assessed (Mishra et al., 2009). The same was true for men in only 1/18 surveys (Vietnam) (Mishra et al., 2009). These findings at an individual level reduce the chance that the association we found between these same factors and HIV at an ecological level is due to an ecological inference fallacy. Furthermore, studies from nationally representative samples in several sub-Saharan African countries have found that region and/or ethnic group remain strongly associated with HIV after controlling for a range of standard individual level risk factors (adjusted odds ratios of up to 14.3 (95% CI 6.1- 33.4) (Barankanira et al., 2016; Fraser-Hutt et al., 2011; Johnson & Way, 2006; Johnson & Budlender, 2002; Mermin et al., 2008; Oluoch et al., 2011). This suggests that there is a risk factor at the level of region and/or ethnic group that was not adequately controlled for in these analyses. This risk factor could be a wide range of factors, such as HSV-2 prevalence (Weiss et al., 2001), composition of the vaginal microbiome (Buve et al., 2014), host genetic factors (Ramsuran et al., 2011) or differential network connectivity (Halpern et al., 2011; Kenyon et al., 2013; Kirby, 2008). Our ecological study is unable to assess which of these factors is responsible for the variations in HIV-prevalence by region. However, our results are compatible with the network-connectivity theory (Kenyon et al., 2017). In brief, this theory posits that populations vary in how connected their sexual networks are (determined by factors such as rate of partner change, concurrency). This, combined with the prevalence of other risk factors that affect the probability of transmission per contact (such as condom use, circumcision and STI prevalence), determines the prevalence of HIV (Kenyon et al., 2017) (see Figure 6 for a more detailed explanation). Why was higher risk sex the behavioral risk factor most strongly associated with HIV-prevalence? Having sex with a non-marital, non-cohabiting person may be an independent risk factor but it may also be associated with another risk behavior that was either unmeasured or inaccurately measured and this other risk behavior is a driver of HIV transmission. Given that the DHS methodology is poor at ascertaining socially sensitive information such as respondent concurrency, questions regarding less sensitive information (such as if a partner was non-marital, non-cohabiting) may be more accurately ascertained.

Conclusion

The available evidence, including the results from this study, suggests that a variety of combinations of behavioural and other risk factors result in high HIV prevalence (Buve, 2006; Kenyon et al., 2014a; Kenyon et al., 2013; Kenyon et al., 2015b; Laumann & Youm, 1999). A striking finding of this study was the strong positive associations between lifetime partners, multiple partners and higher risk sex and the negative association with debut in men. These associations suggest that these risk factors may be underpinned by a common factor. Evidence of varying strengths has been advanced for a wide range of upstream determinants including demographic, socioeconomic and norm-related factors (Abramsky et al., 2014; Barnighausen et al., 2007; Bowleg et al., 2011; Carter et al., 2007; Chen et al., 2007; Goebach et al., 2002; Kenyon et al., 2014a; Kenyon et al., 2015a; Leclerc-Madlala, 2009; Mulawa et al., 2016; Richardson et al., 2014; Yamanis et al., 2016). More research is required to better delineate the relationship between these upstream factors, sexual-behaviors and the resulting sexual-networks and HIV prevalence. Future behavioral and HIV surveys would be strengthened by using audio-computer-assisted self-interview technology to collect sexual behavioral information (Le & Vu, 2012). In countries with large variations in HIV–prevalence thought could be given to using the sexual behaviour of the lower HIV-prevalence communities as positive examples for what could be achieved with behavior change in high HIV-prevalence communities (The Positive Deviance Approach in Female Genital Mutilation Eradication, 1999).

Data availability

The datasets analyzed during the current study are available in the MEASURE DHS repository, (http://www.measuredhs.com). Access to the dataset requires registration, and is granted to those that wish to use the data for legitimate research purposes. A guide for how to apply for dataset access is available at: https://dhsprogram.com/data/Access-Instructions.cfm. The exact datasets analyzed are detailed in Table 1.

Grant information

The author(s) declared that no grants were involved in funding this study.

Acknowledgements

We would like to thank MEASURE DHS for access to the DHS data of the 47 surveys included.

References


Auvert B, Ballard R, Campbell C, et al.: HIV infection among youth in a South

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![Version 1]

Reviewer Report 06 February 2019

https://doi.org/10.5256/f1000research.18706.r43673

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Brian G. Williams
South African Centre for Epidemiological Modelling and Analysis (SACEMA), Stellenbosch University, Stellenbosch, South Africa

The authors carry out a detailed analysis of the association between risky sex and HIV prevalence at the regional level. They show that the prevalence of HIV is associated with risky sex. While this result is not unexpected it is important to quantify the relationship and to examine how the prevalence of HIV and risky sex vary among regions within countries.

However, in my view a critical factor in determining the timing and peak prevalence of HIV epidemics depends critically on how people move. It is trivial to observe that sexually transmitted infections depend on direct, intimate contact between two individuals. If people don’t move then, by definition, the epidemic cannot spread.

DHS studies do not, of course, capture information on migration, and in particular oscillating migration, as is common in South Africa and its neighbouring countries and territories, and a direct consequence of the needs of the mining industry for a cheap supply of very large numbers of unskilled men rooted in the system of Apartheid. I would recommend reading this paper in parallel with the paper by John Hargrove "Migration, mines and mores: the HIV epidemic in southern Africa" published in the South African Journal of Science.

References

Is the work clearly and accurately presented and does it cite the current literature?  
Yes

Is the study design appropriate and is the work technically sound?  
Yes
Are sufficient details of methods and analysis provided to allow replication by others?
Yes

If applicable, is the statistical analysis and its interpretation appropriate?
Yes

Are all the source data underlying the results available to ensure full reproducibility?
Yes

Are the conclusions drawn adequately supported by the results?
Yes

Competing Interests: No competing interests were disclosed.

Reviewer Expertise: Mathematics, Statistics, Epidemiology, HIV, TB

I confirm that I have read this submission and believe that I have an appropriate level of expertise to confirm that it is of an acceptable scientific standard.

Reviewer Report 09 January 2019
https://doi.org/10.5256/f1000research.18706.r42625

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Nico Nagelkerke
Department of Medical Microbiology, University of Manitoba, Winnipeg, Canada

Kenyon and his coauthors explore whether self reported sexual “risk” taking on an aggregate level is associated with HIV prevalence. They present cogent arguments, supported by an extensive theoretical literature on sexual networks and propagation of STIs along such networks, why this is a valuable approach. I agree, but before reading their paper would have been pessimistic about the outcome because of the well-known misreporting of sexual behavior. That they nonetheless find some moderate to strong associations between (aggregated) self-reported risk behavior and HIV prevalence is not just a (methodological) justification of their approach but also (finally perhaps) a rationale for asking such behavior questions (in DHS and other surveys).

The paper is well written with appropriate figures and tables. All potential weaknesses were - as far as I could tell - recognized, acknowledged, and addressed by the authors. As regards to the statistical methods: although alternative approaches were possible, their approach was not flawed, and the choice of methods seemed appropriate.

Is the work clearly and accurately presented and does it cite the current literature?
Yes

Page 19 of 20
Is the study design appropriate and is the work technically sound?  
Yes

Are sufficient details of methods and analysis provided to allow replication by others?  
Yes

If applicable, is the statistical analysis and its interpretation appropriate?  
Yes

Are all the source data underlying the results available to ensure full reproducibility?  
Yes

Are the conclusions drawn adequately supported by the results?  
Yes

Competing Interests: No competing interests were disclosed.

Reviewer Expertise: Biostatistics, Epidemiology and Infectious disease modeling

I confirm that I have read this submission and believe that I have an appropriate level of expertise to confirm that it is of an acceptable scientific standard.

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