These results show extreme vulnerability of HIV-positive women, which has increased the common obstetric risks of gestational process, the immune compromised. Policies directed to the health needs of HIV-positive women become crucial to prevent maternal-to-child transmission of HIV and other STIs.

Disclosure of interest statement The authors report no real or perceived vested interests that relate to this article that could be construed as a conflict of interest.

P16.09

COINFECTION WITH GONORRHOEA, SYPHILIS OR BOTH DOES NOT APPEAR TO AFFECT HIV TRANSMISSION TO THE SEXUAL CONTACTS OF HIV+ PATIENTS WITH UNDETECTABLE VIRAL LOADS

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Background Infection with gonorrhoea (GC) or syphilis is postulated to increase the transmission of HIV 2–5 times; however, studies were done before highly effective HIV therapy was available. In Philadelphia, partner services (PS) is performed, regardless of viral load (VL), for HIV+ patients who are newly diagnosed, STI coinfected, or who are reported as contacts to a patient with new STI or HIV. We hypothesised that STI coinfection would not affect HIV transmission among partners of patients with undetectable VL receiving PS.

Methods HIV+ Philadelphia residents receiving PS from January 2012–December 2014 with a VL within +/- 6 months of PS interview date, with or without STI within +/- 90 days of PS, were included. Partners not already known to be HIV+ who tested for HIV as part of PS were categorised into contacts of either 1) HIV+ patient, undetectable VL (<50 c/ml) or 2) HIV+ patient, detectable/unknown VL.

Results PS encounters were initiated 2,463 times; 80.9% of encounters resulted in interview and 2,106 partners were elicited. Of the 1,211 locatable partners not known to be HIV+, 668 (55.1%) were tested after PS. New HIV was diagnosed more often among patients of partners with detectable/unknown VL (57/435, 13.1%) versus those with undetectable VL (17/233, 7.3%). When patient VL was undetectable, there was little difference in HIV diagnoses among partners of patients with no STI (8/89, 9.0%), syphilis (7/90, 7.7%), GC (2/49, 4.1%), or syphilis/GC (0/5).

Conclusion Patients with undetectable VL who were coinfected with GC, syphilis or both did not transmit HIV to their named sexual contacts at a higher rate than those with HIV alone. Molecular sequencing data can add insight into actual transmission between partners. HIV/STI PS programs could consider deprioritizing the provision of PS to patients with undetectable VL regardless of STI coinfection.

Disclosure of interest None of the authors have any conflicts of interest to disclose.

P16.10

WORSE EPIDEMIC OF EARLY HIV INFECTION AMONG MEN WHO HAVE SEX WITH MEN IN CHINA: IMPLICATION FOR REAL TIME ACTION

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Background Recent upsurge of new HIV infections among men who have sex with men (MSM) is a major concern in China. Paucity of national-level information regarding the burden and predictors of this progressive epidemic of new infections called for a multi-centric, timely and comprehensive investigation.

Methods Mixed methodology was used to recruit MSM (MSM) from seven cities in China between 2012 and 2013. Early and established HIV infections were estimated by Western Blot and BED HIV-1 capture enzyme immunoassay. Syphilis and herpes simplex virus-2 (HSV-2) were also tested.

Results A total of 4496 eligible MSM were recruited. The majority was aged ≤35 years (77.5%), migrants (60.3%), never married (69.8%), and played receptive role in anal sex (70.5%). The HIV prevalence was 9.9%, and 41.9% were recently infected, with HIV incidence of 8.9/100 Person-Years (95% CI: 7.6–10.2). The prevalence of history HSV-2 and syphilis were 12.5% and 8.5%, respectively. Early HIV infection was associated with having multiple male partners (aOR = 1.4, 95% CI 1.1–1.9), recreational drug use (aOR = 2.2, 95% CI 1.6–3.0)), anal bleeding (aOR = 2.1, 95% CI 1.4–3.0), circumcision experience (aOR = 2.0, 95% CI 1.3–3.1), syphilis infection (aOR = 2.8, 95% CI 1.9–4.3) and history HSV-2 infection (aOR = 2.3, 95% CI 1.5–3.3).

Conclusion High rate of early HIV infection is potentially resulting in progressive deterioration of the overall HIV epidemic among MSM in China. Targeted interventions to address high-risk MSM including those having multiple partners, history of recreational drug use and syphilis or HSV-2 infection seemed to be the need of the hour.

P16.11

ESTIMATING THE DISTRIBUTION OF NEW HIV INFECTIONS BY KEY DETERMINANTS IN GENERALISED EPIDEMICS OF SUB-SAHARAN AFRICA USING A VALIDATED MATHEMATICAL MODEL

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Background Recent evidence suggests that sub-Saharan Africa (SSA) is facing a worsening HIV epidemic. There is currently a lack of high-quality data to estimate the geographic and population distribution of HIV incidence in SSA and how this is changing over time. There is also a lack of up-to-date methods to estimate the distribution of new HIV infections among key population subgroups and how this varies by geographic area, gender, and sexual orientation.

Methods We report a new mathematical model to estimate the distribution of new HIV infections by key determinants in SSA for the years 2002–2017. We used a validated mathematical model previously developed to estimate the number of new HIV infections by sex, age, and country for SSA. The model incorporates key determinants of HIV transmission, such as population demographics, sexual behaviour, and HIV incidence and prevalence.

Results The model was able to replicate well the distribution of new HIV infections by sex, age, and country for SSA for the years 2002–2017. The model also showed that new HIV infections are occurring at a much faster rate in SSA compared to previous estimates.

Conclusion The model can be used to estimate the distribution of new HIV infections by key determinants in SSA and to track changes over time. This can help inform targeted interventions to address the HIV epidemic in SSA.
Background Estimating the distribution of new HIV infections according to identifiable characteristics is a priority for programmatic planning in HIV prevention. We propose a mathematical modelling approach that uses robust data sources to estimate the distribution of new infections acquired in the generalised epidemics of sub-Saharan Africa and validate it against cohort data.

Methods We developed a predictive model that represents the population according to factors powerfully associated with risk: gender, marital status, geographic location, key risk behaviours (sex-work, injecting drug-use, male-to-male sex), sero-discordancy within couples, circumcision and ART status. Incidence inference methods are applied to estimate the short-term distribution of new infections by group. The model is applied within a Bayesian framework whereby regional demographic and epidemiological prior information is updated, where possible, with local data. We validated and trained the model against cohort data from Manicaland (Zimbabwe), Kisesa (Tanzania) and Rakai (Uganda). Building on the results from the acquisition model we infer likely sources of transmission. The model was applied to six countries in the region to investigate potential differences in incidence patterns.

Results Without training using the site-specific data, the model was able to predict the pattern of new infections with reasonable accuracy: 95% credible intervals were substantially overlapping and the rank ordering of groups with new infections was consistent. With training using group-specific data on new infections, the accuracy of predictions for subsequent rounds of data improved further and credible intervals narrowed. When applied to the six countries in the region the model showed variation in the distribution of infections between and within countries consistent with the data on prevalence.

Conclusions It is possible to accurately predict, the distribution of new HIV infections acquired using data routinely available in many countries in the Sub-Saharan African region. This validated tool can complement additional analyses on resource allocation and data collection priorities.

Declaration of conflicts of interest All authors declare having no conflicts of interest.

P16.12 OPTIMISATION HIV INVESTMENT IN SWAZILAND: MODELLING HIGH-LEVEL INTERVENTIONS

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Introduction Swaziland has the highest global HIV prevalence with an estimated 27% of people aged 15–49 living with HIV. To address this issue, the Government aimed to optimise HIV investment in the country. The present study was carried out to examine the risk factors of HIV infection among women who lived in an urban area in Swaziland.

Methods The Tanzania HIV/AIDS and Malaria Indicator Survey (2011–12) data has been used. The sample size for urban and rural women who were tested for HIV and ever had sex was 2227 and 6210 respectively. Bivariate and multivariate analyses like logistic regression analyses were used.

Results The present study found that rural women were significantly less likely to be HIV-infected compared to urban women (OR = 0.612, p < 0.00). About 10% urban women were HIV-infected whereas 5.8% women in rural areas were HIV positive. Those women who had more sex partners were significantly more likely to be HIV-infected compared to women who had one sex partner (OR = 4.49, p < 0.00).

Conclusion The present study results suggested that less-educated women, women belong to poor or poorer quintile, women spent night outside, had more sex partners were significantly more likely to have HIV infection among urban women compared to rural women. There is an urgent need for a short and effective program to control the HIV epidemic in urban areas of Tanzania especially for less-educated urban women.

Disclosure of interest statement No conflicts of interest.

P16.13 EPIDEMIOLOGY OF HIV PREVALENCE AMONG URBAN WOMEN IN TANZANIA

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Introduction Tanzania is the country that was the hardest hit by the HIV epidemic in Sub-Saharan Africa. The present study was carried out to examine the risk factors of HIV infection among women who lived in an urban area in Tanzania.

Methods The Tanzania HIV/AIDS and Malaria Indicator Survey (2011–12) data has been used. The sample size for urban and rural women who were tested for HIV and ever had sex was 2227 and 6210 respectively. Bivariate and multivariate analyses like logistic regression analyses were used.

Results The present study found that rural women were significantly less likely to be HIV-infected compared to urban women (OR = 0.612, p < 0.00). About 10% urban women were HIV-infected whereas 5.8% women in rural areas were HIV positive. Those women who had more sex partners were significantly more likely to be HIV-infected compared to women who had one sex partner (OR = 4.49, p < 0.00).

Conclusion The present study results suggested that less-educated women, women belong to poor or poorer quintile, women spent night outside, had more sex partners were significantly more likely to have HIV infection among urban women compared to rural women. There is an urgent need for a short and effective program to control the HIV epidemic in urban areas of Tanzania especially for less-educated urban women.

Disclosure of interest Nothing to Disclose.

P16.14 GEOGRAPHY OF AIDS IN THE STATE OF CEARÃ, BRAZIL

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Introduction The geographical distribution of AIDS subsidises the understanding of the relationship of the disease with socioeconomic and cultural characteristics, enabling local planning of...