Introduction In 2015, the estimated HIV prevalence in Brazil was 0.4%. This figure has been stable in recent years, but it can mask disparities among regions. In this study we present the spatial distribution of the variation in HIV incidence rates (IR) in Brazilian municipalities from 2009–14 and assess the existence of spatial clustering of increase or decrease in these IR.

Methods We used the AIDS reporting system (Sinan) and programmatic data on Viral Load (VL) exams and ARVs. The diagnosis date was used as the earliest among 1st detectable VL, 1st ARV dispensation or diagnosis date in Sinan. Annual IR were generated by municipality. To smooth the IR, we used 3 year averages and applied the local empirical Bayesian method. To assess IR time trends, we calculated the percent IR variations in the period. For spatial statistical analysis, a simple adjacency matrix was generated, and Global and Local Moran’s I autocorrelation tests were applied.

Results The Global Moran’s I for the IR variation was 0.42 (p<0.001) which points to spatial clustering. We generated 2 maps, one for the percent IR variation and another to represent the statistically significant high-high and low-low clusters. In the 1st, we observed that most municipalities in the North (N) and Center-West (CW) present increases in the period; in the Southeast (SE), the state of Sao Paulo (SP) reveals the most relevant decreases in the country; the Northeast (NE), South (S) and some areas of the SE show mixed patterns. The 2nd map makes regional disparities even clearer. There are big clusters of increasing IR in most states of the N, and smaller ones in areas of the CW and NE. Several clusters of declining IR are seen in SP, Minas Gerais (SE), Santa Catarina, Rio Grande do Sul (S), and areas of the NE.

Conclusion Spatial dependency in HIV IR variations in Brazil was evidenced. The methods used in this study have proved useful in monitoring spatiotemporal trends, pointing out important regional differences. Similar analysis can be performed at state and city levels, contributing to improved diagnoses of local epidemics.


Maria Vono Tancredo, CSB Domingues, A Arya, MA Silva, MC Gianna. Centro de Referência e Treinamento DST/AIDS-SP, São Paulo – SP, Brazil

Introduction This study analysed the HIV/AIDS epidemic trends in the state of Sao Paulo, in adults, during the period of 2000 to 2015.

Methods Study conducted by polynomial regression model, with AIDS and HIV positive cases notification comparing by age group and exposure categories. The dependent variable were the annual number of cases, in each of the studied categories, and the independent variable was the time, (calendar years), concerning the study period. The goodness of fit via r² and p<0.05 were used to determine which models and data were most appropriate.

Results It was analysed 142,015 AIDS cases and 70,761 cases of HIV. The AIDS cases trends declined in the entire period, with speed=243 cases/year (p=0.001), but the HIV+ cases trends increased with speed=325 cases/year (p=0.003). From 2000 to 2007 AIDS decreased 30 cases/year (p=0.024) in the group 20–29 years and the 30–39 years group falled 26 cases/year (p=0.003). From 2007 to 2012 there was a growth of MSM with speed=116 cases/year (p=0.039). Among the HIV infection cases in the period 2000 to 2013, the trend was increasing among MSM with speed=337 cases/year (p=0.001) and the group 40–49 presented the highest velocity=8 cases/year (p=0.006). Between 2007 and 2015 the trend was increasing among MSM with speed=337 cases/year (p=0.001) and the...
main age group was 20–24 with speed=106 cases/year, (p=0.043).

Conclusion There was a decreasing trend of AIDS cases and growth in HIV+, with expressive speed among young MSM. This analysis points to the need for discussions of unprotected sexual practices, combined prevention and other actions aimed to controlling the epidemic in greater vulnerability groups.

P3.152 PREVALENCE OF BACTERIAL VAGINOSIS INFECTION AND ASSOCIATED FACTORS IN WOMEN WHO HAVE SEX WITH WOMEN

MTC Duarte, MAO Ignacio, J Andrade, APF Freitas, MG Silva. São Paulo State University, Brazil

Introduction The present research aims to fill a gap in the national and international literature concerning prevalences and factors related to Bacterial vaginosis (BV) in Women who Have Sex with Women (WSW).

Methods It is a cross sectional, analytical and non-ranomized study with 128 WSW from Botucatu - SP and surrounding regions who answered the call from social media, mass communication means, health services and friends or acquaintances from January to November,2015. Data was obtained by the researchers involved in the main study, through interviews ang gynaecological exam. The diagnostic of BV was abtainned through gram staining. Associations were estimated by multiple regression.

Results The prevalence of BV was 41,1%0 and factors associated were vaginal douching [OR=3.29 (IC:95%: 1.26–8.59) p=0014] and sex toys use[OR=2.34 (IC:95%: 1.00–5.50); p=0049].

Conclusion Considered as whole, these data lead to conclusion that the individuals of this study presented high vulnerability to STI/AIDS, as shown by the high prevalence of BV. This study clearly shows the need for a specific health assistance to these women, promoting prevention and education in a holisitic approach.

Spatial and temporal associations between congenital syphilis cases and epidemiological characteristics of infectious syphilis in England


Introduction There has been a rapid rise in infectious syphilis (IS) diagnoses in England since 2011 but congenital syphilis (CS) is rare. In 2016, 3 CS cases were diagnosed in geographically dispersed areas of England. Unusually, their mothers had tested negative at first trimester antenatal screen (screen-negative), indicating syphilis acquisition during pregnancy. Simulation modelling using historical CS cases indicated an event probability of 3%. We investigated the spatial and temporal relationship between screen-negative CS cases and IS epidemiology in the affected areas.

Methods Data from 01/2011-06/2016 were obtained from GUMCADv2, the national electronic surveillance system. England was divided into 3 syphilis epidemiological areas (SEAs): wider incident areas (WIAs; the 3 affected and immediate surrounding counties); endemic areas (with established epidemics in men who have sex with men-MSM) and non-incident non-endemic areas (NINEAs). Time-series analysis (TSA) was used to estimate IS outliers by gender, sexual orientation and SEA. Associations between IS characteristics and SEA (WIAs vs. NINEAs) were assessed using Pearson’s chi-square and Kruskal-Wallis tests. Mothers of CS cases were excluded from analyses.

Results In 2011–2016, IS rates/100,000 in WIAs rose in heterosexual women (1.3–3.0) and MSM (8.9–13.9) but fell in heterosexual men (3.7–3.0). In NINEAs, rates rose in heterosexual women (1.6–1.9), MSM (5.0–10.8) and heterosexual men (2.7–3.2). On TSA, IS cases significantly exceeded expected bounds in 2016 in heterosexual women in WIAs; no exceedance was seen in NINEAs. In 2016, heterosexual women with IS were more likely to be UK-born in WIAs than in NINEAs (78% vs. 39%; p<0.001). A greater proportion of MSM were bisexual in WIAs than in NINEAs (11% vs. 8%; p<0.001).

Conclusion Increased syphilis transmission in some sexual networks of MSM and a higher proportion of bisexual men in WIAs may have created more opportunities for IS acquisition in women. Efforts to raise awareness of the potential risk of acquiring syphilis during pregnancy are needed.


1Artin, 2Sawatzky, 3Lefebvre, 4Allen, 5Ndiau, 6Hoang, 7Horsman, 8Mulvey.
1Public Health Agency of Canada, Winnipeg, Canada; 2Laboratoire De Santé Publique Du Québec, Ste-Anne-De-Bellevue, Canada; 3Public Health Ontario, Toronto, Canada; 4Alberta Health Services, Edmonton, Canada; 5British Columbia Centres For Disease Control, Vancouver, Canada; 6Saskatchewan Disease Control Laboratory, Regina, Canada

Introduction Neisseria gonorrhoeae have acquired resistance to many antimicrobials including third generation cephalosporins and azithromycin, which are the current co-therapy recommended by the Canadian STI guidelines for gonorrhoea treatment. Minimum inhibitory concentrations (MIC) to azithromycin and molecular sequence types were determined for N. gonorrhoeae circulating in Canada.

Methods Between 2014 and 2015, N. gonorrhoeae strains isolated by Canadian provincial public health laboratories were submitted to the National Microbiology Laboratory (NML) (n=4,720) for azithromycin MIC determination (resistance MIC ≥2.0 mg/L) by agar dilution. N. gonorrhoeae multi-antigen sequence typing (NG-MAST) was used for molecular typing.

Results Azithromycin resistance was identified in 3.3% (127/3,809) and 4.7% (198/4,190) of N. gonorrhoeae in 2014 and 2015, respectively, a significant increase since 2013 (p<0.001). MICs ranged from 2 to 16 mg/L. The most common sequence types identified in 2014 were ST10451 (n=40), ST10567 (n=38) and ST11765 (n=10). ST11765 is closely related to ST10451, differing by 1 bp in the por allele. In 2015, the prevalent sequence types were ST12302 (n=110), ST10451 (n=34) and ST9047 (n=23). ST10451, identified in Quebec, Ontario and Alberta, was newly identified in 2014 and also identified in 2015. ST10451 is related to ST1407 (differing by 1 bp in the por allele) which is an internationally-