

repeats within the *arp* gene and restriction fragment length polymorphism analysis of *tpv* subfamily II genes) into sequence types.

**Methods** *T. pallidum* PCR positive ulcer samples from 219 patients diagnosed in Denmark between 2009 and 2013 were identified for the study. At present, 16 specimens have been analysed and included in the study. Molecular typing was performed by sequence analysis of a 400-base pair region of *tp0548* and will be supplemented with the CDC typing system.

Clinical data were obtained from the Danish National Syphilis Reporting System where patients diagnosed with early syphilis in Denmark are registered.

**Results** Sequence analysis of *tp0548* revealed three sequence types designated f, g, and c, among the 16 patients. Four patients had *tp0548* sequence type f. These patients were all men, and comprised both heterosexual men and MSM. One of the patients was originally from Iran. Eleven patients, including one female, had *tp0548* sequence type g. These patients all reported Denmark as country of origin and the majority were inhabitants of Copenhagen. One patient had *tp0548* sequence type c. This patient was a heterosexual man originally from Pakistan.

**Conclusion** These preliminary results show that a minimum of three *tp0548* sequence types are prevalent in Denmark. To further differentiate between strain types, the clinical samples will be characterised using the CDC typing system. This study only included patients with lesions which limit our results to patients with early syphilis.

#### P1.024 TRICHOMONAS VAGINALIS DETECTION AND CHARACTERIZATION FROM WOMEN ATTENDING AN ANTI-RETROVIRAL CLINIC IN PRETORIA, SOUTH AFRICA

doi:10.1136/sextrans-2013-051184.0245

<sup>1</sup>I Rukasha, <sup>1,2</sup>M M Ehlers, <sup>1,2</sup>M M Kock. <sup>1</sup>University of Pretoria, Pretoria, South Africa; <sup>2</sup>National Health Laboratory Service, Pretoria, South Africa

**Background** Trichomoniasis caused by *Trichomonas vaginalis* affects both men and women, although the clinical presentation differs. Infected women are more likely to have symptoms compared to infected men. Approximately 75% to 100% of men are asymptomatic in comparison to approximately 50% to 75% of women who are identified as asymptomatic. In South Africa, data on the prevalence and detection of *T. vaginalis* is well-documented; however, data on the molecular characterization of *T. vaginalis* is limited. The study aimed to detect and characterise *T. vaginalis* isolates from HIV positive women attending an anti-retroviral clinic.

**Methods** Self-collected vaginal swabs from 380 HIV positive women were included in the study. *Trichomonas vaginalis* was detected using wet mount microscopy, culture and a commercial PCR kit. The genetic relatedness of 92 culture positive *T. vaginalis* isolates was determined. Five primers (TV1, TV2, TV3, TV5 and TV6) were used for the RAPD assay. The PCR-IGS-RFLP products were digested with five enzymes, namely: *AluI*, *HinfI*, *RsaI*, *Sau3AI* and *Tsp509*.

**Results** A total of 8% (30/380) of specimens were positive for *T. vaginalis* using microscopy, 24% (92/380) of specimens were positive using culture and 31% (118/380) of the specimens were positive using the commercial PCR kit. RAPD assay analysis showed a high level of genetic diversity between the *T. vaginalis* isolates. The dendrograms obtained from the RAPD data grouped the 92 *T. vaginalis* isolates into between nine to 24 clusters with 70% similarity used to define clusters, while the PCR-IGS RFLP assay results for the isolates were genetically indistinguishable.

**Conclusion** The PCR assay was the most sensitive diagnostic tool for the detection of *T. vaginalis*. A high prevalence of *T. vaginalis*, consisting of different strain types, was detected in the HIV positive women included in the study.

#### P1.025 EFFECTS OF ABNORMAL VAGINAL FLORA AND MENOPAUSE ON CERVICOVAGINAL FLUID VISCOSITY

doi:10.1136/sextrans-2013-051184.0246

<sup>1</sup>C A Chappell, <sup>2</sup>L C Rohan, <sup>2</sup>L Wang, <sup>2</sup>L Meyn, <sup>2</sup>B J Moncla, <sup>1,2</sup>K Bunge, <sup>2</sup>S L Hillier. <sup>1</sup>University of Pittsburgh, Pittsburgh, PA, United States; <sup>2</sup>Magee-Womens Research Institute, Pittsburgh, PA, United States

**Background** Sexually transmitted pathogens, including HIV, are increased among women having abnormal vaginal flora. The mucus gel layer is a component of innate mucosal immunity, and the presence of mucus may contribute to the viscosity of genital tract fluid. Little is known about the impact of reproductive hormones, menopause and vaginal flora patterns on the viscosity of cervicovaginal fluid.

**Methods** Vaginal swabs and cervicovaginal lavage (CVL) were collected from 134 healthy asymptomatic post-menopausal women (n = 23), women in the follicular (n = 26) or proliferative (n = 19) phase, and women using levonorgestrel IUDs (n = 28), DMPA (n = 13) or combined oral contraceptives (n = 25). Vaginal smears were evaluated using the Nugent criteria. The viscosity (centipoise, cP) of each sample was measured in triplicate using a Cambridge MicroSample Viscometer. Student's t-test, analysis of variance, and linear regression were used to assess statistical significance.

**Results** The mean CVL viscosity among the 84 women having a Nugent score < 4 was 1.51(± 0.48) vs. 1.26(± 0.29) cP for the 50 women with abnormal vaginal flora (p = 0.001). There was no difference in CVL viscosity for women with intermediate vs. BV flora (p > 0.99). Similarly, there was no difference in viscosity among women at different phases of their menstrual cycle, nor among women using different hormonal contraceptive methods. However, the CVL of 23 postmenopausal women was less viscous compared to the 111 premenopausal women, 1.16(± 0.26) vs. 1.47(± 0.44) cP respectively (p = 0.001). In a linear regression model, abnormal flora (p = 0.01) and postmenopausal status (p = 0.005) were independently associated with decreased CVL viscosity.

**Conclusion** Abnormal flora and being post-menopausal are independently associated with decreased CVL viscosity. Even though phase of menstrual cycle and hormonal contraceptive use has been posited to have an impact on cervical mucus, these data suggest that these factors do not have a measurable impact of vaginal fluid viscosity.

#### P1.026 VALIDITY OF URINE-BASED DIAGNOSIS OF BACTERIAL VAGINOSIS

doi:10.1136/sextrans-2013-051184.0247

<sup>1</sup>H Verstraelen, <sup>2</sup>A Swidsinski. <sup>1</sup>Ghent University, Ghent, Belgium; <sup>2</sup>Laboratory for Molecular Genetics, Polymicrobial Infections and Bacterial Biofilms, Berlin, Germany

**Background** Current gold standard diagnosis of bacterial vaginosis (BV) relies on categorising Gram-stained vaginal smears through Nugent scoring. We have recently described an alternative method of diagnosis, based on visualisation through fluorescence-in-situ-hybridisation (FISH) of the BV biofilm on desquamated vaginal epithelial cells present in urine sediments.

**Methods** A vaginal swab and a first void urine specimen were obtained from 72 pregnant women attending the antenatal clinic. The vaginal swab was used to assess the vaginal microbiota status through Nugent scoring. The urine specimen served for FISH-based diagnosis of the Gardnerella dominated polymicrobial adherent biofilms attached to desquamated vaginal epithelial cells.

**Results** Among the 12 women with BV on urine assessment, 10 had BV according to Nugent's score and 2 had intermediate microbiota. Presence of Gardnerella in a planktonic mode of growth occurred with 8 women and all have Nugent scores ≤ 3. Among the 52 women in which no Gardnerella could be documented through FISH, 2 had a Nugent score of ≥ 7, 2 a Nugent score of 4 to 6, and the remainder a Nugent score of ≤ 3. Accordingly, when comparing