BACTERIAL VAGINOSIS AND HIGH-RISK HUMAN PAPILLOMAVIRUS COINFECTION AMONG AFRICAN AMERICAN WOMEN IN THE UNITED STATES

Purnima Madhivanan*, Makiella Coudray, Daniel Ruiz-Perez, Brett Colbert, Karl Krupp, Hansi Kumar, Giri Naranatham, Kalai Mathee, Florida International University, Epiology, Miami, USA; Florida International University, Bioinformatics Research Group, Miami, USA; Florida International University, Department of Biological Sciences, College of Arts and Sciences, Miami, USA; Florida International University, Department of Health Promotion and Disease Prevention, Robert Stempel College of Public Health, Miami, USA.

**Results**

Eighty reproductive age African American women were included in the analysis. The point prevalence of HRHPV was 48.1% (95% CI: 37–59%). The mean age of the participants was 21.4 years (SD: 2.11), 81.2% graduated high school. Prior antibiotic use was low (3.8%), and 75% were not treated for BV during their lifetime. Among those who had been treated previously for BV, most women were treated ≥ five times (60%). According to Nugent Scores, 70% had BV, 13.7% had intermediate and 16.3% had healthy vaginal flora. Among HRHPV positive women, 66.7% were infected with single HRHPV genotype, 33.3% with multiple HRHPV genotypes. Concurrent HRHPV and BV infection was found among 33.3% of the sample. However, there was no significant difference between the prevalence of HRHPV among women with and without BV.

**Conclusion**

Co-occurrence of HRHPV and BV among this group of young African American women was relatively high. Considering that these conditions are very common among women worldwide, further research in this field is imperative. More studies are needed to accurately evaluate temporal sequence of acquisition of both conditions in any attempt to establish a causal relationship.

**Disclosure**

No significant relationships.

CO-OCCURRENCE OF BACTERIAL VAGINOSIS AND TRICHOMONAS VAGINALIS AMONG YOUNG AFRICAN AMERICAN WOMEN

Purnima Madhivanan*, Makiella Coudray, Daniel Ruiz-Perez, Brett Colbert, Karl Krupp, Hansi Kumar, Giri Naranatham, Kalai Mathee, Florida International University, Epiology, Miami, USA; Florida International University, Bioinformatics Research Group, Miami, USA; Florida International University, Department of Biological Sciences, College of Arts and Sciences, Miami, USA; Florida International University, Department of Health Promotion and Disease Prevention, Robert Stempel College of Public Health, Miami, USA; Florida International University, Biomolecular Sciences Institute, Miami, USA.

**Background**

While the etiology of bacterial vaginosis (BV) is still not known, it is described as a polymicrobial condition that lacks lactic-acid producing Lactobacillus species with an overgrowth of anaerobic bacteria and elevated vaginal pH. This study aims to evaluate the relationship between BV assessed by Nugent scoring of vaginal smear and Trichomonas vaginalis infection among African American young women in the U.S.

**Methods**

 Stored vaginal swabs from a previously completed clinical trial were acquired for this study. The kinds of bacteria present in the samples were identified by classifying 16S rRNA gene sequences using high-throughput pyrosequencing. Vaginal smears were also categorized by the Nugent Gram stain score (0–3, normal; 4–6, intermediate state; 7–10, BV). TV genotyping was performed using quantitative polymerase chain reaction, performed using TaqMan probes in a customized plate (Thermo Fisher Scientific; Waltham, Massachusetts). Descriptive statistics were conducted to determine the odds of TV infection among women with BV.

**Results**

This study included 80 African American reproductive age women with a mean age of 21.4 years (SD: 2.11 years). Most (81.2%) women had graduated high school. 70% (95% CI: 37–59%) had BV, 13.7% had intermediate and 16.3% had healthy vaginal flora. TV was diagnosed among 11.1% (95% CI: 4–8%) of the women. Prior antibiotic use was low (3.8%), and 75% of women had not been treated for BV during their lifetime. Among those who were previously treated for BV, 60% were treated five or more times. Douching was reported by 49% of the sample. 35% of TV cases had concurrent BV, while 11.1% of TV cases also had intermediate vaginal flora. There were no associations with prior antibiotic use, hormonal contraception, douching or prior treatment.

**Conclusion**

Young African American women of reproductive age found to have abnormal vaginal flora should be screened for Trichomonas vaginalis infection.

**Disclosure**

No significant relationships.

OVERLAP BETWEEN AMSEL’S CRITERIA, NUGENT’S GRAM STAIN SCORE, AND VAGINAL MICROBIOTA COMMUNITY STATE TYPES

Antonio Salas, Khalil Ghanem, Rebecca Brotman, Jacques Ravel, Susan Tuddenham, Johns Hopkins University School of Medicine, Baltimore, USA; University of Maryland, Institute of Genome Sciences, Baltimore, USA.

**Background**

Bacterial vaginosis (BV) is characterized by low-Lactobacilli and increased anaerobes. BV can be determined by clinical assessment (Amsel’s criteria) or microscopy (Nugent scoring); molecular methods are also under study. We investigated concordance between Amsel-BV, Nugent-BV and low-Lactobacilli vaginal microbiota identified via 16S rRNA gene sequencing.

**Methods**

Vaginal swabs and clinical data were collected from women enrolled in a longitudinal study. Amsel’s criteria were determined clinically and Nugent scoring (Nugent-BV=7–10, intermediate=4–6) was determined by microscopy. Vaginal microbiota were characterized using 16S rRNA gene sequencing and categorized into 7 community state types (CSTs): 4 dominated by Lactobacillaceae (CST I, II, III and V), and 3 by Streptococcus spp. (CST VI), Bifidobacterium spp. (CST VII), or a variety of anaerobes (CST IV).

**Disclosure**

No significant relationships.