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

Purnima Madhivanan*, Makella Coudray, Daniel Ruiz-Perez, Brett Colbert, Karl Krupp, Hansi Kumar, Giri Narasimhan, Florida International University, Epidemiology, 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: Bacterial vaginosis (BV) increases the risk of many sexually transmitted infections. The co-occurrence of persistent BV and high-risk HPV (HRHPV) increases the risk of developing cervical cancer. This study aims to investigate the co-occurrence of HRHPV and BV among young women in the US.

Methods: Stored vaginal swabs were acquired from a previously completed clinical trial. The kinds of bacteria present in the samples were identified by classifying 16S rRNA gene sequences in each sample using high-throughput pyrosequencing. HPV genotyping was performed using quantitative polymerase chain reaction using TaqMan probes in a customized plate (Thermo Fisher Scientific; Waltham, Massachusetts). BV was classified using Nugent Scores of Gram Stain.

Results: Eighty reproductive age African American (AA) women were included in the analysis. The point prevalence of HRHPV was 48.1% (95% CI: 37.5–59%). The mean age of the participated women 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 5–6 times (60%). According to Nugent Scores, 70% had BV; 13.7% had intermediate flora and 16.3% were healthy. 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.

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

Antonio Saleas, Khail 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: 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 Gram stain 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: 37.5–59%) 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. 55% of TV cases had concurrent BV, while 11.1% of TV cases also had intermediate vaginal flora. There were no association 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.
Results 110 samples, ~10 from each CST, were selected. 5.5% (n=6) had AmS-BV, 32.7% (n=36) Nugent-BV, and 36.4% (n=40) had low-Lactobacillus CSTs (IV, VI, VII); 8.2% had symptoms. Among AmS-BV samples, 83.3% had Nugent-BV, 16.7% had intermediate Nugent score and all were CST IV. 86.1% of women with Nugent-BV and 85% of women with low-Lactobacillus CSTs did not have AmS-BV. 22.2% of those with Nugent-BV did not have low-Lactobacillus CSTs; of these 50% were CST III (L. iners-dominated). 46.7% of CST-III had a vaginal pH ≥ 4.5, and 13.3% had a Nugent BV score. 22 samples had a vaginal pH=4.5 and a normal Nugent score: 45.5% were in CST III, 9.1% in low-Lactobacillus CSTs. 30.6% women with Nugent-BV had a vaginal pH < 4.5; of these 45.5% were in CST IV.

Conclusion Nugent score and low-Lactobacillus CST were concordant. L. iners-dominated CSTs often had normal Nugent scores and high pH. Among mostly asymptomatic women, a large proportion with low-Lactobacillus CSTs did not have AmS-BV. Future studies assessing long term clinical outcomes will be needed to determine whether molecular methods provide added actionable or prognostic information.

Disclosure No significant relationships.