Cpn60 Species Characterization of the Vaginal Microbiome of Women with Recurrent Bacterial Vaginosis Voque1B2.01.61-Vogue1B2.01.10-Voque1B2.01.62-Vogue1B2.01.37-Vogue1B2.01.35-Vogue1B2.01.29-Vogue1B2.01.26-Vogue1B2.01.11-Vogue1B2.01.64-Vogue1B2.01.56-\$\text{Vogue1B2.01.52-}
\text{Vogue1B2.01.38-}
\text{Vogue1B2.01.19-}
\text{Vogue1B2.01.06-}
\text{Vogue1B2.01.50-} ○ Vogue1B2.01.23 Vogue1B2.01.21-Vogue1B2.01.09-Vogue1B2.01.08-Vogue1B2.01.07-Vogue1B2.01.01-Vogue1B2.01.63-Vogue1B2.01.58-Vogue1B2.01.28-Vogue1B2.01.15-Voque1B2.01.12-25 50 75 100 **Species Proportion** Actinobacteria.sp.

Atopobium.vaginae

Clostridia.sp..BVAB2

Clostridium.genomosp..BVAB3

Escherichia.coli

Gardnerella.vaginalis.Group.A

Lactobacillus.gasseri

Lactobacillus.iners

Lactobacillus.iners

Lactobacillus.iners

Lactobacillus.jensenii

Megasphaera.sp..genomosp..type.1 Other.Streptococcus Other.Bacteroidetes Other.Clostridium Prevotella.amnii Other.Firmicutes Prevotella.timonensis Other.Lactobacillus Streptococcus.devriesei Other.Prevotella Eukaryote Lactobacillus.crispatus Other.Bacteria Other.Proteobacteria