Cpn60 Species Characterization of the Vaginal Microbiome of Women with Recurrent Bacterial Vaginosis Vogue1B2.01.61-Vogue1B2.01.56-Voque1B2.01.38-Vogue1B2.01.26-Vogue1B2.01.64-Vogue1B2.01.62-Vogue1B2.01.10-Vogue1B2.01.52-Vogue1B2.01.35-Vogue1B2.01.21-Strongue 1B2.01.08-Vogue 1B2.01.58-Vogue 1B2.01.50-Vogue 1B2.01.29-○ Vogue1B2.01.28 Vogue1B2.01.23-Vogue1B2.01.12-Vogue1B2.01.07-Vogue1B2.01.06-Vogue1B2.01.01-Vogue1B2.01.19-Vogue1B2.01.63-Vogue1B2.01.37-Vogue1B2.01.15-Vogue1B2.01.09-25 50 75 100 **Species Proportion** Actinobacteria.sp.

Atopobium.vaginae

Clostridia.sp..BVAB2

Clostridium.genomosp..BVAB3

Escherichia.coli

Gardnerella.vaginalis.Group.A Lactobacillus.gasseri

Gardnerella.vaginalis.Group.B Lactobacillus.iners

Lactobacillus.iners

Lactobacillus.iners

Lactobacillus.iners

Cardnerella.vaginalis.Group.D 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