Genus Composition of Bacterial Communities Across Different Sample Genus 1.00 -UCG-005 Faecalibacterium Subdoligranulum Facklamia Escherichia-Shigella Streptococcus Erysipelatoclostridium Staphylococcus 0.75 **-**Enterococcus Sellimonas Ruminococcus Eisenbergiella Genus Relative Abundance Rudaeicoccus DTU089 Dellaglioa Romboutsia Pygmaiobacter Corynebacterium 0.50 -Parasporobacterium Colidextribacter Murimonas Christensenellaceae R-7 group Millisia CHKCI002 Ligilactobacillus Butyricicoccus Lactobacillus Brachybacterium 0.25 -Lachnospiraceae UCG-008 Blautia Lachnospiraceae NC2004 group Asaccharospora Lachnoclostridium Alistipes Incertae Sedis Aerococcus 0.00 -HT002 [Ruminococcus] torques group ME16S12 ME16S3 ME16S1 ME16S10 ME16S2 ME16S11