



- Lactobacillus rogosae
- Enterocloster multispecies_sppn13_2_nov_94.470%
- Hungatella xylanolytica
- Dysosmobacter welbionis_nov_93.953%
- Fusicatenibacter saccharivorans
- Roseburia inulinivorans
- Parabacteroides merdae
- Phascolarctobacterium faecium
- Bilophila wadsworthia
- Hungatella xylanolytica_nov_97.674%
- Sporobacter termitidis_nov_92.130%
- Alistipes onderdonkii
- Subdoligranulum variabile
- Ruminococcus albus_nov_90.187%
- Parasutterella excrementihominis
- Bacteroides fragilis
- Bacteroides faecichinchillae_faecis_thetaiotaomicron
- Bacteroides xylanisolvans
- Bacteroides koreensis_kribbi
- Ruminococcus champanellensis_nov_97.674%
- Faecalibacterium prausnitzii_nov_96.744%
- Bacteroides eggerthii
- Ruminiclostridium cellobioparum_nov_89.352%
- Butyrivibrio crossotus
- Alistipes putredinis
- Monoglobus pectiniilyticus_nov_90.698%
- Phocaeicola massiliensis
- Barnesiella intestinihominis
- Bacteroides cellulosilyticus_timonensis
- Bacteroides caccae
- Bacteroides stercoris
- Ruminiclostridium cellobioparum_nov_92.056%
- Oscillibacter valericigenes_nov_93.056%
- Blautia wexlerae
- Roseburia faecis
- Oscillospiraceae_[G-6] bacterium_MOT-153_nov_97.674%
- Prevotella stercorea
- Roseburia intestinalis
- Muribaculum intestinale_nov_85.116%
- Faecalibacterium prausnitzii_nov_97.209%
- Eubacteriales_[G-1] bacterium_MOT-144
- Massiliiprevotella massiliensis
- Prevotella copri_nov_97.674%
- Bacteroides uniformis
- Phocaeicola dorei
- Escherichia_Pseudoescherichia_Shigella boydii_coli_fergusonii_flexn
- Akkermansia muciniphila
- Faecalibacterium prausnitzii
- Phocaeicola vulgatus
- Prevotella copri

Samples