



- Anaerostipes hadrus
- Anaerobutyricum hallii
- Enterocloster multispecies_sppn13_2_nov_94.470%
- Dysosmobacter welbionis_nov_93.953%
- Bilophila wadsworthia
- Hungatella effluvii_nov_96.744%
- Blautia faecis
- Subdoligranulum variabile
- Fusicatenibacter saccharivorans
- Lactobacillus rogosae
- Bacteroides caccae
- Eubacterium ruminantium_nov_96.279%
- Sporobacter termitidis_nov_92.130%
- Parabacteroides merdae
- Bacteroides koreensis_kribbi
- Roseburia intestinalis
- Clostridium oceanicum_nov_91.705%
- Bacteroides stercoris
- Phascolarctobacterium faecium
- Ruminiclostridium cellobioparum_nov_92.056%
- Ruminococcoides bili
- Faecalibacterium prausnitzii_nov_96.744%
- Oscillibacter valericigenes_nov_93.056%
- Bacteroides cellulosilyticus_timonensis
- Faecalibacterium prausnitzii_nov_97.209%
- Parasutterella excrementihominis
- Barnesiella intestinihominis
- Alistipes putredinis
- Monoglobus pectinilyticus_nov_90.698%
- Aestuariaispira insulae_nov_85.000%
- Bacteroides fragilis
- Bacteroides eggerthii
- Blautia wexlerae
- Bacteroides ovatus
- Buttiauxella_Enterobacter_Erwinia_Klebsiella_Kluyv ...(10 species)
- Oscillospiraceae_[G-6] bacterium_MOT-153_nov_97.674%
- Phocaecicola plebeius
- Alistipes onderdonkii
- Eubacteriales_[G-1] bacterium_MOT-144
- Roseburia faecis
- Ruminococcus champanellensis_nov_97.674%
- Prevotella copri
- Bacteroides faecichinchillae_faecis_thetaiotaomicron
- Escherichia_Pseudoescherichia_Shigella boydii_coli_fergusonii_flexn
- Phocaecicola dorei
- Akkermansia muciniphila
- Bacteroides uniformis
- Faecalibacterium prausnitzii
- Phocaecicola vulgatus
- Akkermansia muciniphila_nov_97.209%

Samples