



Species	
Barnesiella intestinihominis	
Butyrivibrio crossotus	
Blautia faecis	
Roseburia faecis	
Phascolarctobacterium faecium	
Eubacteriales_[G-1] bacterium_MOT-144	
Subdoligranulum variabile	
Parasutterella excrementihominis	
Roseburia inulinivorans	
Lactobacillus rogosae	
Bacteroides stercoris	
Bacteroides ovatus	
Fusicatenibacter saccharivorans	
Bacteroides fragilis	
Blautia luti	
Alistipes onderdonkii	
Blautia wexlerae	
Bacteroides caccae	
Akkermansia muciniphila	
Alistipes putredinis	
Alistipes shahii	
Ruminococcoides bili	
Massiliiprevotella massiliensis	
Phocaeicola dorei	
Bacteroides uniformis	
Phocaeicola vulgatus	
Prevotella stercorea	
Faecalibacterium prausnitzii	
Prevotella copri	
Bacteroides eggerthii	
Parabacteroides merdae	
Ruminococcus albus_nov_90.187%	
Faecalibacterium prausnitzii_nov_97.209%	
Barnesiella intestinihominis_nov_97.674%	
Aestuariuspira insulae_nov_85.000%	
Oscillibacter valericigenes_nov_93.056%	
Sporobacter termitidis_nov_92.130%	
Monoglobus pectinilyticus_nov_90.698%	
Ruminococcus champanellensis_nov_97.674%	
Ruminiclostridium cellobioparum_nov_92.056%	
Oscillospiraceae_[G-6] bacterium_MOT-153_nov_97.674%	
Faecalibacterium prausnitzii_nov_96.744%	
Dysosmobacter welbionis_nov_93.953%	
Ruminiclostridium cellobioparum_nov_89.352%	
Bacteroides faecichinchillae_faecis_thetaiotaomicron	
Escherichia_Pseudoescherichia_Shigella boydii_coli_fergusonii_flexnerii	
Buttiauxella_Enterobacter_Erwinia_Klebsiella_Kluysv ...(10 species)	
Bacteroides koreensis_kribbi	
Bacteroides cellulosilyticus_timonensis	
Enterocloster multispecies_sppn13_2_nov_94.470%	

F3410872.S92
F3410872.S79
F3410872.S66
F3410872.S61
F3410872.S66
F3410872.S44
F3410872.S24
F3410872.S22
F3410872.S05
F3410872.S98
F3410872.S93
F3410872.S82
F3410872.S74
F3410872.S69
F3410872.S59
F3410872.S30
F3410872.S27
F3410872.S25
F3410872.S20
F3410872.S18
F3410872.S16
F3410872.S14
F3410872.S12
F3410872.S10
F3410872.S07
F3410872.S04
F3410872.S01

Samples