



Barnesiella intestinihominis	Species
Blautia faecis	
Roseburia faecis	
Phascolarctobacterium faecium	
Roseburia intestinalis	
Eubacteriales_[G-1] bacterium_MOT-144	
Bacteroides xylanisolvens	
Subdoligranulum variabile	
Parasutterella excrementihominis	
Bacteroides stercoris	
Bacteroides ovatus	
Fusicatenibacter saccharivorans	
Bacteroides fragilis	
Blautia obeum	
Hungatella xylanolytica	
Alistipes onderdonkii	
Blautia wexlerae	
Bacteroides caccae	
Akkermansia muciniphila	
Alistipes putredinis	
Ruminococcoides bili	
Bilophila wadsworthia	
Phocaeicola dorei	
Bacteroides uniformis	
Phocaeicola vulgatus	
Phocaeicola massiliensis	
Faecalibacterium prausnitzii	
Prevotella copri	
Parabacteroides merdae	
Solibaculum mannosilyticum_nov_93.925%	
Faecalibacterium prausnitzii_nov_97.209%	
Akkermansia muciniphila_nov_97.209%	
Muribaculum intestinale_nov_85.116%	
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%	
Hungatella xylanolytica_nov_97.674%	
Faecalibacterium prausnitzii_nov_96.744%	
Prevotella copri_nov_97.674%	
Dysosmobacter welbionis_nov_93.953%	
Ruminiclostridium cellobioparum_nov_89.352%	
Hungatella effluvii_nov_96.744%	
Bacteroides faecichinchillae_faecis_thetaiotaomicron	
Escherichia_Pseudoescherichia_Shigella boydii_coli_fergusonii_flexner	
Bacteroides koreensis_kribbi	
Bacteroides cellulosilyticus_timonensis	
Enterocloster multispecies_sppn13_2_nov_94.470%	

F3410872.S95
F3410872.S90
F3410872.S83
F3410872.S76
F3410872.S73
F3410872.S54
F3410872.S43
F3410872.S32
F3410872.S94
F3410872.S88
F3410872.S77
F3410872.S70
F3410872.S67
F3410872.S62
F3410872.S55
F3410872.S52
F3410872.S50
F3410872.S45
F3410872.S40
F3410872.S38
F3410872.S36
F3410872.S34
F3410872.S31

Samples