



Bacteroides intestinalis	
Bacteroides thetaiotaomicron	
Haemophilus parainfluenzae	
Kosakonia sacchari	
Clostridium disporicum	
Corynebacterium amycolatum	
Enterococcus avium	
Odoribacter laneus	
Collinsella aerofaciens	
Erysipelatoclostridium [Clostridium] saccharogumia	
Phocaeicola vulgaris	
Enterobacter cancerogenus	
Shigella sonnei	
Escherichia coli	
Ligilactobacillus murinus	
Blautia glucerasea	
Granulicatella adiacens	
Bacteroides xylanisolvans	
Bifidobacterium pseudolongum	
Longicatena caecimuris	
Lactobacillus intestinalis	
Shigella dysenteriae	
Eisenbergiella massiliensis	
Triticum aestivum	
Staphylococcus ureilyticus	
Fusicatenibacter saccharivorans	
Parasutterella excrementihominis	
Streptococcus agalactiae	
Veillonella parvula	
Enterococcus faecalis	
Streptococcus sp._HMT_064	
Streptococcus mutans	
Mammaliococcus sciuri	
Alistipes finegoldii	
Romboutsia ilealis	
Parabacteroides goldsteinii	
Staphylococcus haemolyticus	
Akkermansia muciniphila	
Adlercreutzia equolifaciens	
Monoglobus pectinilyticus	
Bacteroides uniformis	
Lachnospiraceae_[G-3] bacterium_MOT-168_nov_94.154%	
Niveispirillum fermenti_nov_85.242%	
Rothia nasimurium_nov_97.228%	
Streptococcus sp._MOT-045_nov_97.764%	
Anaerotruncum aminivorans_nov_92.812%	
Kiloniella majae_nov_85.144%	
Blautia luti_nov_96.008%	
Frisingicoccus caecimuris_nov_96.451%	
Blautia glucerasea_nov_97.899%	
Eubacterium xylanophilum_nov_91.313%	
Turicibacter sanguinis_nov_96.137%	
Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.130%	
Eubacterium coprostanoligenes_nov_91.379%	
Coprococcus eutactus_nov_92.704%	
Agathobaculum desmolans_nov_86.345%	
Butyricoccus pullicaecorum_nov_87.216%	
Oscillospiraceae_[G-6] bacterium_MOT-153_nov_97.831%	
Longibaculum muris_nov_92.754%	
Glucerbacter canis_nov_95.388%	
Blautia faecicola_nov_93.111%	
Oscillospiraceae_[G-4] bacterium_MOT-151_nov_90.496%	
Lachnoclostridium [Clostridium] herbivorans_nov_92.723%	
Lachnospiraceae_[G-3] bacterium_MOT-168_nov_93.750%	
Ruminococcus albus_nov_91.045%	
Lawsonibacter asaccharolyticus_nov_90.909%	
Acutalibacter muris_nov_95.661%	
Barnesiella viscericola_nov_90.447%	
Fusicatenibacter saccharivorans_nov_89.474%	
Oscillospiraceae_[G-1] bacterium_MOT-147_nov_95.218%	
Lachnospiraceae_[G-9] bacterium_MOT-174_nov_96.855%	
Erysipelatoclostridium [Clostridium] saccharogumia_nov_96.042%	
Anaerostipes hadrus_nov_92.754%	
Fusicatenibacter saccharivorans_nov_97.479%	
Faecalimonas umbilicata_nov_96.033%	
Lachnospiraceae_[G-11] bacterium_MOT-178_nov_94.929%	
Oscillospiraceae_[G-4] bacterium_MOT-151_nov_92.946%	
Anaerotruncum rubiinfantis_nov_88.820%	
Oscillospiraceae_[G-3] bacterium_MOT-150_nov_91.097%	
Zhongshania marina_nov_79.765%	
Pseudoflavonifractor phocaeensis_nov_94.191%	
Pseudoflavonifractor phocaeensis_nov_92.340%	
Lachnospiraceae_[G-3] bacterium_MOT-168_nov_94.154%	
Oscillospiraceae_[G-3] bacterium_MOT-150_nov_90.702%	
Coprococcus eutactus_nov_95.085%	
Lachnospiraceae_[G-4] bacterium_MOT-169_nov_97.065%	
Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.340%	
Muribaculaceae_[G-2] bacterium_MOT-104_nov_86.228%	
Blautia schinkii_nov_94.340%	
Duncaniella freteri_nov_89.024%	
Alistipes finegoldii_nov_94.835%	
Blautia producta_nov_96.017%	
Ruminococcus albus_nov_92.225%	
Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.473%	
Blautia producta_nov_95.178%	
Spongibacter marinus_nov_80.142%	
Staphylococcus argenteus_aureus_roterodami	
Streptococcus parasanguinis_parasanguinis_clade_721	
Veillonella dispar_parvula	
Kordiimonas multispecies_sppn3_2_nov_79.121%	

Species

Samples