



Species	
Bacteroides intestinalis	
Bacteroides thetaiotaomicron	
Atopostipes sp._MOT-200	
Haemophilus parainfluenzae	
Kosakonia sacchari	
Corynebacterium marinum	
Clostridium disporicum	
Microbacterium hydrocarbonoxydans	
Corynebacterium amycolatum	
Enterococcus avium	
Odoribacter laneus	
Collinsella aerofaciens	
Erysipelatoclostridium [Clostridium] saccharogumia	
Phocaeicola vulgatus	
Enterobacter cancerogenus	
Shigella sonnei	
Shigella flexneri	
Escherichia coli	
Ligilactobacillus murinus	
Blautia glucerasea	
Cutibacterium acnes	
Granulicatella adiacens	
Bacteroides xylanisolvens	
Bifidobacterium pseudolongum	
Rothia kristinae	
Longicatena caecimuris	
Lactobacillus intestinalis	
Shigella dysenteriae	
Rothia aerolata	
Eisenbergiella massiliensis	
Staphylococcus ureilyticus	
Actinidia eriantha	
Fusicatenibacter saccharivorans	
Parasutterella excrementihominis	
Ruthenibacterium lactatiformans	
Stenotrophomonas pavanii	
Streptococcus agalactiae	
Veillonella parvula	
Enterococcus faecalis	
Bilophila wadsworthia	
Streptococcus sp._HMT_064	
Streptococcus mutans	
Mammaliococcus sciuri	
Alistipes finegoldii	
Romboutsia ilealis	
Parabacteroides goldsteinii	
Staphylococcus haemolyticus	
Akkermansia muciniphila	
Adlercreutzia equolifaciens	
Monoglobus pectinilyticus	
Bacteroides uniformis	
Rothia nasimurium_nov_97.228%	
Streptococcus sp._MOT-045_nov_97.764%	
Anaerotignum aminivorans_nov_92.812%	
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%	
Butyricicoccus 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%	
Ruminococcus albus_nov_91.045%	
Acutalibacter muris_nov_95.661%	
Oscillospiraceae_[G-1] bacterium_MOT-147_nov_95.218%	
Erysipelatoclostridium [Clostridium] saccharogumia_nov_96.042%	
Anaerostipes hadrus_nov_92.754%	
Fusicatenibacter saccharivorans_nov_97.479%	
Faecalimonas umbilicata_nov_96.033%	
Oscillibacter valericigenes_nov_94.595%	
Coprococcus eutactus_nov_95.085%	
Lachnospiraceae_[G-4] bacterium_MOT-169_nov_97.065%	
Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.340%	
Mediterraneibacter [Ruminococcus] gnavus_nov_94.990%	
Caproiciproducens galactitolivorans_nov_91.083%	
Arthrospira platensis_nov_90.531%	
Lachnospira multipara_nov_84.298%	
Propioniferax innocua_nov_95.207%	
Blautia hydrogenotrophica_nov_93.582%	
Muribaculaceae_[G-2] bacterium_MOT-104_nov_86.228%	
Faecalicatena fissicatena_nov_96.042%	
Eubacterium coprostanoligenes_nov_91.845%	
Blautia schinkii_nov_94.340%	
Duncaniella freteri_nov_89.024%	
Lachnospiraceae_[G-6] bacterium_MOT-171_nov_96.226%	
Alistipes finegoldii_nov_94.835%	
Blautia producta_nov_96.017%	
Ruminococcus albus_nov_92.225%	
Chryseobacterium culicis_nov_97.280%	
Blautia producta_nov_95.178%	
Staphylococcus argenteus_aureus_roterodami	
Streptococcus parasanguinis_parasanguinis_clade_721	
Veillonella dispar_parvula	
Staphylococcus saprophyticus_xylosus	

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Samples