



- Faecalicatena fissicatena_nov_96.042%
- Oscillibacter valericiogenes_nov_94.595%
- Ruthenibacterium lactatiformans
- Oscillospiraceae_[G-1] bacterium_MOT-147_nov_95.218%
- Alistipes finegoldii_nov_94.835%
- Glucerberacter canis_nov_95.388%
- Erysipelatoclostridium [Clostridium] saccharogumia_nov_96.042%
- Agathobaculum desmolans_nov_86.345%
- Faecalimonas umbilicata_nov_96.033%
- Clostridium disporicum
- Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.340%
- Eubacterium xylanophilum_nov_91.313%
- Lachnospira multipara_nov_84.298%
- Frisingicoccus caecimuris_nov_96.451%
- Eubacterium coprostanoligenes_nov_91.845%
- Acutalibacter muris_nov_95.661%
- Anaerotignum aminivorans_nov_92.812%
- Coprococcus eutactus_nov_92.704%
- Blautia hydrogenotrophica_nov_93.582%
- Mediterraneibacter [Ruminococcus] gnavus_nov_94.990%
- Odoribacter laneus
- Bilophila wadsworthia
- Lachnospiraceae_[G-6] bacterium_MOT-171_nov_96.226%
- Coprococcus eutactus_nov_95.085%
- Bacteroides intestinalis
- Eubacterium coprostanoligenes_nov_91.379%
- Blautia schinkii_nov_94.340%
- Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.130%
- Oscillospiraceae_[G-6] bacterium_MOT-153_nov_97.831%
- Caproiciproducens galactitolivorans_nov_91.083%
- Turcibacter sanguinis_nov_96.137%
- Blautia faecicola_nov_93.111%
- Ruminococcus albus_nov_92.225%
- Bacteroides xylanisolvans
- Eisenbergiella massiliensis
- Bacteroides thetaiotaomicron
- Parabacteroides goldsteinii
- Duncaniella freteri_nov_89.024%
- Rothia kristinae
- Corynebacterium marinum
- Ruminococcus albus_nov_91.045%
- Blautia producta_nov_96.017%
- Butyricicoccus pullicaecorum_nov_87.216%
- Monoglobus pectinilyticus
- Blautia producta_nov_95.178%
- Anaerostipes hadrus_nov_92.754%
- Longibaculum muris_nov_92.754%
- Adlercreutzia equolifaciens
- Alistipes finegoldii
- Blautia glucerasea_nov_97.899%
- Propioniferax innocua_nov_95.207%
- Stenotrophomonas pavanii
- Romboutsia ilealis
- Parasutterella excrementihominis
- Erysipelatoclostridium [Clostridium] saccharogumia
- Shigella sonnei
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_86.228%
- Bacteroides uniformis
- Enterococcus avium
- Corynebacterium amycolatum
- Staphylococcus saprophyticus_xylosus
- Chryseobacterium culicis_nov_97.280%
- Microbacterium hydrocarbonoxydans
- Enterococcus faecalis
- Staphylococcus haemolyticus
- Kosakonia sacchari
- Phocaeicola vulgatus
- Enterobacter cancerogenus
- Cutibacterium acnes
- Shigella dysenteriae
- Atopostipes sp._MOT-200
- Granulicatella adiacens
- Shigella flexneri
- Arthrospira platensis_nov_90.531%
- Escherichia coli
- Actinidia eriantha
- Streptococcus agalactiae
- Akkermansia muciniphila
- Lachnospiraceae_[G-4] bacterium_MOT-169_nov_97.065%
- Fusicatenibacter saccharivorans
- Longicatena caecimuris
- Fusicatenibacter saccharivorans_nov_97.479%
- Streptococcus sp._HMT_064
- Veillonella parvula
- Collinsella aerofaciens
- Blautia luti_nov_96.008%
- Staphylococcus ureilyticus
- Staphylococcus argenteus_aureus_rotterdamii
- Mammalicoccus sciuri
- Rothia aerolata
- Blautia glucerasea
- Veillonella dispar_parvula
- Ligilactobacillus murinus
- Lactobacillus intestinalis
- Streptococcus parasanguinis_parasanguinis_clade_721
- Haemophilus parainfluenzae
- Bifidobacterium pseudolongum
- Streptococcus sp._MOT-045_nov_97.764%
- Rothia nasimurium_nov_97.228%
- Streptococcus mutans

Species

Samples