



- Acinetobacter septicus
- Phocaeicola vulgatus
- Lachnospiraceae_[G-3] bacterium_MOT-168
- Bacteroides caecimuris
- Staphylococcus saprophyticus
- Turicimonas muris
- Alistipes sp._MOT-127
- Mammaliicoccus sciuri
- Adlercreutzia equolifaciens
- Bacteroides uniformis
- Bifidobacterium pseudolongum
- Faecalibaculum rodentium
- Ligilactobacillus murinus
- Parabacteroides goldsteinii
- Lactobacillus johnsonii
- Erysipelatoclostridium [Clostridium] cocleatum
- Dubosiella newyorkensis
- Clostridium disporicum
- Serratia marcescens
- Enterococcus faecalis
- Mammaliicoccus lentus
- Lachnospiraceae_[G-3] bacterium_MOT-168_nov_95.000%
- Alistipes putredinis_nov_92.213%
- Alistipes shahii_nov_93.827%
- Faecalicoccus acidiformans_nov_89.600%
- Parafannyhessea umbonata_nov_92.161%
- Parvibacter caecicola_nov_97.609%
- Lachnospiraceae_[G-3] bacterium_MOT-168_nov_93.737%
- Adlercreutzia caecimuris_nov_93.290%
- Faecalicatena fissicatena_nov_93.528%
- Alistipes putredinis_nov_92.813%
- Parasutterella excrementihominis_nov_94.578%
- Lawsonia intracellularis_nov_89.336%
- Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.351%
- Alistipes senegalensis_nov_96.920%
- Ihubacter massiliensis_nov_94.572%
- Lachnospiraceae_[G-2] bacterium_HMT_096_nov_91.632%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_85.887%
- Turicibacter sanguinis_nov_95.923%
- Duncaniella freteri_nov_87.375%
- Faecalicatena fissicatena_nov_95.407%
- Glucerbacter canis_nov_93.501%
- Lachnospiraceae_[G-7] bacterium_MOT-172_nov_91.097%
- Anaeromassilibacillus senegalensis_nov_92.489%
- Oscillospiraceae_[G-3] bacterium_MOT-150_nov_93.125%
- Staphylococcus saprophyticus_xylosus
- Serratia marcescens_nematodiphila
- Olsenella multispecies_sppn1_2_nov_91.966%
- Faecalicatena multispecies_sppn2_2_nov_92.067%
- Faecalicatena multispecies_sppn3_2_nov_93.737%

Species

- F29493.S08
- F29493.S07
- F29493.S06
- F29493.S05
- F29493.S12
- F29493.S11
- F29493.S10
- F29493.S09

Samples