



Acinetobacter septicus
Phocaeicola vulgatus
Lachnospiraceae_[G-3] bacterium_MOT-168
Turicimonas muris
Alistipes sp._MOT-127
Bacteroides acidifaciens
Adlercreutzia equolifaciens
Bacteroides uniformis
Bifidobacterium pseudolongum
Faecalibaculum rodentium
Ligilactobacillus murinus
Lactobacillus intestinalis
Parabacteroides goldsteinii
Lactobacillus johnsonii
Erysipelatoclostridium [Clostridium] cocleatum
Dubosiella newyorkensis
Clostridium disporicum
Staphylococcus ureilyticus
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%
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%
Peptostreptococcaceae_[G-5] bacterium_HMT_493_nov_90.000%
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%

F29493.S04

F29493.S03

F29493.S02

F29493.S01

F29493.S12

F29493.S11

F29493.S10

F29493.S09

Samples

Species