



- Phocaeicola sartorii
- Clostridioides difficile
- Porphyromonas sp._MOT-131
- Adlercreutzia equolifaciens
- Akkermansia muciniphila
- Bifidobacterium pseudolongum
- Ileibacterium valens
- Muribacter sp._MOT-143
- Carnobacteriaceae_[G-1] bacterium_MOT-197
- Ligilactobacillus murinus
- Erysipelotrichaceae_[G-1] bacterium_MOT-189
- Parabacteroides distasonis
- Staphylococcus ureilyticus
- Bacteroides acidifaciens
- Bacteroides uniformis
- Bacteroides caccae
- Enterococcus faecalis
- Lachnospiraceae_[G-3] bacterium_MOT-168
- Weeksellaceae_[G_1] bacterium_MOT-126
- Cutibacterium acnes
- Dubosiella newyorkensis
- Streptococcus acidominimus
- Rodentibacter pneumotropicus
- Eubacteriales_[G-1] bacterium_MOT-159
- Corynebacterium mastitidis
- Parabacteroides goldsteinii
- Streptococcus danieliae
- Streptococcus thoraltensis
- Mammalicoccus lentus
- Lactobacillus johnsonii
- Erysipelatoclostridium [Clostridium] cocleatum
- Faecalibaculum rodentium
- Lachnospiraceae_[G-14] bacterium_MOT-185_nov_96.781%
- Lachnospiraceae_[G-14] bacterium_MOT-185_nov_95.717%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_89.431%
- Anaerostipes caccae_nov_96.328%
- Lachnospiraceae_[G-2] bacterium_HMT_096_nov_91.632%
- Duncaniella freteri_nov_86.263%
- Alistipes putredinis_nov_93.224%
- Glucerbacter canis_nov_93.305%
- Turcibacter sanguinis_nov_95.923%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.755%
- Bacteroides uniformis_nov_95.893%
- Lachnospiraceae_[G-3] bacterium_MOT-168_nov_94.792%
- Parafannyhessea umbonata_nov_92.161%
- Duncaniella freteri_nov_88.531%
- Duncaniella freteri_nov_89.775%
- Alistipes senegalensis_nov_93.443%
- Staphylococcus saprophyticus_xylosus
- Olsenella multispecies_sppn1_2_nov_91.966%

Species

F33955.S28

F33955.S27

F33955.S26

F33955.S25

F33955.S32

F33955.S31

F33955.S30

F33955.S29

Samples