



- Phocaeicola sartorii
- Clostridioides difficile
- Adlercreutzia equolifaciens
- Akkermansia muciniphila
- Bifidobacterium pseudolongum
- Ileibacterium valens
- Muribacter sp._MOT-143
- Ligilactobacillus murinus
- Erysipelotrichaceae_[G-1] bacterium_MOT-189
- Parabacteroides distasonis
- Staphylococcus ureilyticus
- Bacteroides acidifaciens
- Enterococcus faecalis
- Lachnospiraceae_[G-3] bacterium_MOT-168
- 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%
- Lachnospiraceae_[G-2] bacterium_HMT_096_nov_91.632%
- Duncaniella freteri_nov_86.263%
- Alistipes putredinis_nov_93.224%
- Glucerbacter canis_nov_93.305%
- Turicibacter sanguinis_nov_95.923%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.755%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_85.887%
- Lachnospiraceae_[G-3] bacterium_MOT-168_nov_94.792%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.200%
- Parafannyhessea umbonata_nov_92.161%
- Lachnospiraceae_[G-3] bacterium_MOT-168_nov_96.050%
- Duncaniella freteri_nov_88.531%
- Oscillospiraceae_[G-6] bacterium_MOT-153_nov_86.475%
- Alistipes putredinis_nov_92.418%
- Duncaniella freteri_nov_89.775%
- Eubacterium ventriosum_nov_93.320%
- Duncaniella freteri_nov_86.061%
- Alistipes senegalensis_nov_93.443%
- Anaeroplasma abactoclasticum_nov_86.538%
- Staphylococcus saprophyticus_xylosus
- Olsenella multispecies_sppn1_2_nov_91.966%

Species

F33955.S24
F33955.S23
F33955.S22
F33955.S21
F33955.S28
F33955.S27
F33955.S26
F33955.S25

Samples