



- Porphyromonas sp._MOT-131
- Pediococcus pentosaceus
- Streptococcus acidominimus
- Streptococcus danieliae
- Eubacteriales_[G-4] bacterium_MOT-164
- Enterococcus faecalis
- Corynebacterium stationis
- Mammaliococcus lentus
- Bifidobacterium pseudolongum
- Mollicutes_[G-1] bacterium_MOT-186
- Lactococcus lactis
- Lactobacillus johnsonii
- Streptococcus thoraltensis
- Stenotrophomonas [Pseudomonas] hibiscicola
- Staphylococcus ureilyticus
- Weissella cibaria
- Staphylococcus saprophyticus
- Anaerococcus sp._HMT_290
- Triticum aestivum
- Muribacter sp._MOT-143
- Rodentibacter pneumotropicus
- Streptococcus thermophilus
- Moraxella osloensis
- Enterococcus gallinarum
- Limosilactobacillus reuteri
- Staphylococcus equorum
- Massilia aurea
- Psychrobacter alimentarius
- Dubosiella newyorkensis
- Atopostipes sp._MOT-201
- Ligilactobacillus animalis
- Lachnospiraceae_[G-14] bacterium_MOT-185
- Ligilactobacillus murinus
- Eubacteriales_[G-2] bacterium_MOT-162
- Cutibacterium granulosum
- Corynebacterium mastitidis
- Cutibacterium acnes
- Clostridium disporicum
- Stenotrophomonas maltophilia
- Streptomyces aculeolatus
- Lachnospiraceae_[G-11] bacterium_MOT-177
- Pseudomonas helleri
- Lactcaseibacillus rhamnosus
- Jeotgalicoccus halotolerans
- Levilactobacillus brevis
- Gemella sp._MOT-033
- Corynebacterium ammoniagenes
- Secundilactobacillus paracollinoides
- Leptothrix sp._HMT_025
- Weissella paramesenteroides
- Delftia acidovorans
- Carnobacteriaceae_[G-1] bacterium_MOT-197
- Rothia nasimurium
- Pelomonas saccharophila
- Ralstonia sp._HMT_406
- Actinidia eriantha
- Akkermansia muciniphila
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_88.105%
- Duncaniella freteri_nov_93.293%
- Turicibacter sanguinis_nov_95.923%
- Enterococcus faecalis_nov_95.825%
- Fusobacterium varium_nov_96.696%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.867%
- Oribacterium parvum_nov_89.770%
- Duncaniella freteri_nov_89.775%
- Alistipes senegalensis_nov_93.443%
- Erysipelotrichaceae_[G-1] bacterium_MOT-189_nov_88.798%
- Peptococcus sp._HMT_168_nov_84.866%
- Duncaniella freteri_nov_89.718%
- Fusicatenibacter saccharivorans_nov_90.526%
- Oscillospiraceae_[G-6] bacterium_MOT-153_nov_91.631%
- Actinidia eriantha_nov_97.011%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.000%
- Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.719%
- Fusobacterium perfoetens_nov_91.126%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.423%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_85.887%
- Duncaniella freteri_nov_88.934%
- Duncaniella freteri_nov_93.699%
- Latilactobacillus curvatus_graminis
- Sphingomonas aquatilis_melonis
- Staphylococcus saprophyticus_xylosus

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Samples

Species