



- Triticum aestivum
- Delftia acidovorans
- Enterococcus gallinarum
- Staphylococcus saprophyticus
- Ligilactobacillus murinus
- Pseudomonas helleri
- Eubacteriales_[G-4] bacterium_MOT-164
- Stenotrophomonas maltophilia
- Psychrobacter alimentarius
- Eubacteriales_[G-2] bacterium_MOT-162
- Clostridium disporicum
- Stenotrophomonas [Pseudomonas] hibiscicola
- Cutibacterium acnes
- Lachnospiraceae_[G-11] bacterium_MOT-177
- Cutibacterium granulosum
- Corynebacterium stationis
- Streptococcus thermophilus
- Staphylococcus equorum
- Massilia aurea
- Jeotgalicoccus halotolerans
- Mammaliococcus lentus
- Lachnospiraceae_[G-14] bacterium_MOT-185
- Dubosiella newyorkensis
- Corynebacterium ammoniagenes
- Enterococcus faecalis
- Actinidia eriantha
- Atopostipes sp._MOT-201
- Mollicutes_[G-1] bacterium_MOT-186
- Ralstonia sp._HMT_406
- Anaerococcus sp._HMT_290
- Streptomyces aculeolatus
- Lactobacillus johnsonii
- Bifidobacterium pseudolongum
- Staphylococcus ureilyticus
- Lacticaseibacillus rhamnosus
- Akkermansia muciniphila
- Leptothrix sp._HMT_025
- Limosilactobacillus reuteri
- Moraxella osloensis
- Secundilactobacillus paracollinoides
- Pelomonas saccharophila
- Ligilactobacillus animalis
- Bradyrhizobium pachyrhizi
- Fusobacterium varium_nov_96.696%
- Duncaniella freteri_nov_93.293%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.867%
- Enterococcus faecalis_nov_95.825%
- Fusicatenibacter saccharivorans_nov_90.526%
- Duncaniella freteri_nov_89.775%
- Alistipes senegalensis_nov_93.443%
- Peptococcus sp._HMT_168_nov_84.866%
- Duncaniella freteri_nov_89.718%
- Oscillospiraceae_[G-6] bacterium_MOT-153_nov_91.631%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.000%
- Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.719%
- Fusobacterium perfoetens_nov_91.126%
- Actinidia eriantha_nov_97.011%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.423%
- Yaniella halotolerans_nov_97.040%
- Duncaniella freteri_nov_88.934%
- Oscillospiraceae_[G-1] bacterium_MOT-147_nov_96.674%
- Lachnoclostridium [Clostridium] aminophilum_nov_89.792%
- Duncaniella freteri_nov_93.699%
- Oscillospiraceae_[G-3] bacterium_MOT-150_nov_92.917%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_88.105%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_85.887%
- Parafannyhessea umbonata_nov_92.161%
- Turicibacter sanguinis_nov_95.923%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.198%
- Oribacterium parvum_nov_89.770%
- Staphylococcus saprophyticus_xylosus
- Sphingomonas aquatilis_melonis
- Faecalicatena multispecies_sppn3_2_nov_92.067%

Species

F17722.S01

F17722.S02

F17722.S03

F17722.S04

F17722.S05

F17722.S06

F17722.S07

F17722.S08

F17722.S09

F17722.S10

Samples