



- Species
- Staphylococcus saprophyticus
 - Lachnospiraceae_[G-14] bacterium_MOT-185
 - Enterococcus faecalis
 - Corynebacterium ammoniagenes
 - Staphylococcus equorum
 - Mammalicoccus lentus
 - Bradyrhizobium pachyrhizi
 - Triticum aestivum
 - Moraxella osloensis
 - Eubacteriales_[G-4] bacterium_MOT-164
 - Jeotgalicoccus halotolerans
 - Lachnospiraceae_[G-11] bacterium_MOT-177
 - Actinidia eriantha
 - Limosilactobacillus reuteri
 - Clostridium disporicum
 - Enterococcus gallinarum
 - Cutibacterium acnes
 - Mollicutes_[G-1] bacterium_MOT-186
 - Corynebacterium stationis
 - Staphylococcus ureilyticus
 - Psychrobacter alimentarius
 - Ligilactobacillus murinus
 - Streptococcus thermophilus
 - Stenotrophomonas maltophilia
 - Lactocaseibacillus rhamnosus
 - Atopostipes sp._MOT-201
 - Massilia aurea
 - Akkermansia muciniphila
 - Pelomonas saccharophila
 - Ralstonia sp._HMT_406
 - Pseudomonas helleri
 - Cutibacterium granulosum
 - Delftia acidovorans
 - Ligilactobacillus animalis
 - Lactobacillus johnsonii
 - Stenotrophomonas [Pseudomonas] hibiscicola
 - Eubacteriales_[G-2] bacterium_MOT-162
 - Leptothrix sp._HMT_025
 - Dubosiella newyorkensis
 - Streptomyces aculeolatus
 - Secundilactobacillus paracollinoides
 - Bifidobacterium pseudolongum
 - Anaerococcus sp._HMT_290
 - 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%
 - Duncaniella freteri_nov_88.934%
 - Yaniella halotolerans_nov_97.040%
 - 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_85.887%
 - Muribaculaceae_[G-1] bacterium_MOT-129_nov_88.105%
 - Parafannyhessea umbonata_nov_92.161%
 - Oribacterium parvum_nov_89.770%
 - Turicibacter sanguinis_nov_95.923%
 - Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.198%
 - Staphylococcus saprophyticus_xylosus
 - Sphingomonas aquatilis_melonis
 - Faecalicatena multispecies_sppn3_2_nov_92.067%

Samples