



- Streptococcus danieliae
- Akkermansia muciniphila
- Clostridium disporicum
- Mammaliococcus lentus
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_91.870%
- Enterococcus faecalis
- Salmonella enterica
- Citrobacter koseri
- Limosilactobacillus fermentum
- Staphylococcus argenteus_aureus_roterodami
- Duncaniella freteri_nov_88.577%
- Prevotella shahii_nov_87.903%
- Staphylococcus saprophyticus_xylosus
- Streptococcus mitis
- Burkholderia aenigmatica_cepacia_contaminans_lata_multivorans
- Burkholderia_Caenibaculum aenigmatica_baiyandianus_cepacia_cc
- Staphylococcus hominis
- Mesorhizobium huakuii
- Bacillus halotolerans_spizizenii
- Listeria monocytogenes
- Shigella flexneri
- Cutibacterium granulosum
- Fingoldia magna
- Cutibacterium acnes_nov_96.815%
- Cutibacterium granulosum_nov_96.788%
- Anaerococcus octavius_nov_94.583%
- Streptococcus danieliae_nov_93.387%
- Bosea eneeae_vestrisii
- Duncaniella freteri_nov_91.751%
- Corynebacterium pilbarensis
- Thermus thermophilus
- Segetibacter korensis_nov_89.733%
- Corynebacterium mucifaciens
- Streptococcus mitis_pneumoniae_pseudopneumoniae
- Staphylococcus capitis_epidermidis
- Corynebacterium tuberculostearicum
- Enterobacter sp._MOT-050
- Peptoniphilus lacydonensis
- Staphylococcus ureilyticus
- Streptococcus danieliae_nov_95.582%
- Escherichia_Shigella albertii_sonnei
- Escherichia fergusonii
- Agrobacterium_Rhizobium lusitanum_rhizogenes
- Streptococcus chosunense
- Streptococcus danieliae_nov_96.787%
- Streptococcus danieliae_nov_93.952%
- Mammaliococcus sciuri
- Shigella sonnei
- Shigella boydii
- Streptococcus danieliae_nov_97.379%
- Ligilactobacillus murinus
- Burkholderia contaminans_lata_multivorans
- Escherichia coli
- Cutibacterium acnes
- Phyllobacterium myrsinacearum
- Streptococcus sp._MOT-012

Species

Samples