



- 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
- Finexgoldia 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 koreensis_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

O13 O15 O17 O19 O29 O31 O33 O35 O57 O59 O61 O63 O73 O75 O77 O79 O21 O23 O25 O27 O37 O39 O41 O43 O65 O67 O69 O71 O81 O83 O85 O87 N1 N3 N5 N7 N9

Samples