



- Species
- 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

N2 O61 N9 N8 N3 O88 O75 O29 O13 O44 O60 O64 O58 O22 O38 O14 O82 O65 O62 O32 O70 O34 O40 O36 O41 O78 O28 O17 O43 O15 O30 O21 O76 O79 O86 O74 O68

Samples