



- 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_cepacia
- 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

NZ
O61
N9
N8
N3
O88
O75
O29
O13
O44
O60
O64
O58
O22
O38
O14
O82
O65
O65
O71
O70
O66
O80
O36
O41
O78
O28
O17
O43
O15
O30
O21
O76
O79
O86
O74
O68

Species

Samples