



- 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

NZ  
 O61 N9 N8 N3 O88 O75 O29 O13 O44 O60 O64 O58 O22 O14 O82 O65 O62 O32 O70 O34 O80 O36 O41 O78 O28 O17 O43 O15 O87 O72 O76 O79 O86 O74 O68

Samples