



- Staphylococcus capitis
- Brachybacterium paraconglomeratum
- Granulicatella adiacens
- Leifsonia aquatica
- Mammaliicoccus sciuri
- Klebsiella grimontii
- Staphylococcus ureilyticus
- Mammaliicoccus lentus
- Streptococcus agalactiae
- Acinetobacter courvalinii
- Brachybacterium conglomeratum
- Corynebacterium stationis
- Clostridium disporicum
- Enterococcus faecalis
- Streptococcus acidominimus
- Enterococcus gallinarum
- Corynebacterium lowii
- Haemophilus parainfluenzae
- Aerococcus urinaeequi
- Jeotgalicoccus halotolerans
- Ligilactobacillus murinus
- Klebsiella oxytoca
- Enterococcus haemoperoxidus
- Acinetobacter bereziniae
- Romboutsia ilealis
- Gordonia polyisoprenivorans
- Streptococcus parasanguinis
- Streptococcus mitis
- Rothia nasimurium
- Streptococcus sp._MOT-045
- Streptococcus mutans
- Proteus mirabilis
- Citrobacter koseri
- Vallitalea pronyensis_nov_88.227%
- Streptococcus sp._MOT-045_nov_97.609%
- Globicatella sanguinis_nov_93.157%
- Turicibacter sanguinis_nov_97.503%
- Sporosarcina sp._MOT-205_nov_97.019%
- Blautia schinkii_nov_95.261%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_91.473%
- Sporosarcina sp._MOT-204_nov_95.599%
- Lederbergia lenta_nov_95.190%
- Ileibacterium valens_nov_94.552%
- Anaerostipes caccae_nov_97.708%
- Staphylococcus capitis_epidermidis
- Escherichia_Shigella fergusonii_flexneri
- Staphylococcus argenteus_aureus_roterodami
- Shigella flexneri_sonnei
- Staphylococcus saprophyticus_xylosus
- Facklamia_Ruoffia tabacinasalis

Species

- F24059.S06
- F24059.S05
- F24059.S04
- F24059.S03
- F24059.S02
- F24059.S01
- F24059.S17
- F24059.S16
- F24059.S15
- F24059.S14
- F24059.S13

Samples