



- Brachybacterium paraconglomeratum
- Enterobacter mori
- Granulicatella adiacens
- Mammaliococcus sciuri
- Klebsiella grimontii
- Neisseria elongata
- Staphylococcus ureilyticus
- Mammaliococcus lentus
- Streptococcus agalactiae
- Lactobacillus intestinalis
- Corynebacterium stationis
- Akkermansia muciniphila
- Clostridium disporicum
- Enterococcus faecalis
- Streptococcus acidominimus
- Lactobacillus johnsonii
- Enterococcus gallinarum
- Corynebacterium lowii
- Haemophilus parainfluenzae
- Aerococcus urinaeequi
- Jeotgalicoccus halotolerans
- Ligilactobacillus murinus
- Streptococcus oralis
- Clostridium saudiense
- Romboutsia ilealis
- Streptococcus parasanguinis
- Streptococcus mitis
- Rothia nasimurium
- Streptococcus sp.\_MOT-045
- Streptococcus mutans
- Bifidobacterium pseudolongum
- Escherichia coli
- Limosilactobacillus reuteri
- Citrobacter koseri
- Rodentibacter rattii
- Staphylococcus equorum
- Veillonella parvula
- Streptococcus sp.\_MOT-045\_nov\_97.609%
- Globicatella sanguinis\_nov\_93.157%
- Turicibacter sanguinis\_nov\_97.503%
- Lachnospiraceae\_[G-5] bacterium\_MOT-170\_nov\_92.852%
- Ruminococcus albus\_nov\_94.425%
- Blautia schinkii\_nov\_95.261%
- Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_91.102%
- Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_91.473%
- Escherichia\_Shigella fergusonii\_flexneri
- Staphylococcus argenteus\_aureus\_roterodami
- Shigella flexneri\_sonnei
- Staphylococcus saprophyticus\_xylosus
- Facklamia\_Ruoffia tabacinasalis

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Samples

Species