



- Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_86.089%
- Haemophilus haemolyticus
- Pseudomonas putida
- Lachnospiraceae\_[G-14] bacterium\_MOT-182\_nov\_90.254%
- Actinomyces sp.\_HMT\_896
- Acidovorax ebreus
- Staphylococcus pasteurii
- Sphingobacterium multivorum
- Aggregatibacter seignis
- Lachnospiraceae\_[G-6] bacterium\_MOT-171\_nov\_94.572%
- Butyrivibrio proteoclasticus\_nov\_83.594%
- Duncaniella freteri\_nov\_89.135%
- Sphingomonas zeae
- Streptococcus sanguinis
- Staphylococcus argenteus\_aureus\_roterodami
- Corynebacterium aurimucosum
- Selenomonas sp.\_HMT\_137\_nov\_97.228%
- Gemella morbillorum
- Streptococcus thoraltensis
- Streptococcus cristatus\_clade\_578
- Lachnospiraceae\_[G-3] bacterium\_MOT-168\_nov\_94.792%
- Streptococcus chosunense
- Streptococcus mitis
- Escherichia fergusonii
- Filifactor alocis
- Porphyromonas gingivalis
- Acidovorax temperans
- Ileibacterium valens
- Lachnospiraceae\_[G-5] bacterium\_MOT-170
- Parasutterella excrementihominis
- Acutalibacter muris
- Bacteroides acidifaciens
- Streptococcus oralis\_subsp\_tigurinus\_clade\_071
- Lachnospiraceae\_[G-14] bacterium\_MOT-185
- Lactobacillus intestinalis
- Campylobacter showae
- Cetobacterium somerae
- Streptococcus oralis
- Eubacteriales\_[G-1] bacterium\_MOT-158
- Stenotrophomonas [Pseudomonas] hibiscicola
- Sphingomonas aquatilis\_melonis
- Desulfobulbus sp.\_HMT\_041
- Duncaniella freteri\_nov\_93.712%
- Psychrobacter arcticus\_fjordensis
- Streptococcus salivarius
- Enterococcus faecalis
- Agromyces mediolanus
- Porphyromonas pasteri
- Akkermansia muciniphila
- Prevotella nigrescens
- Acinetobacter lwoffii
- Leptotrichia buccalis
- Alishewanella agri
- Lactobacillus johnsonii
- Leucobacter chromiirestiens
- Citrobacter koseri
- Staphylococcus capitis
- Porphyromonas endodontalis
- Fusobacterium periodonticum
- Enterobacter cancerogenus
- Enterobacter cloacae
- Phocaeicola sartorii
- Leptotrichia hofstadii\_nov\_96.970%
- Schaalia sp.\_HMT\_178
- Staphylococcus saprophyticus\_xylosum
- Duncaniella freteri\_nov\_90.612%
- Kosakonia sacchari
- Fusobacterium nucleatum\_subsp\_vincentii
- Methylobacterium brachiatum
- Cutibacterium acnes
- Campylobacter concisus
- Streptococcus gordonii
- Enterococcus casseliflavus
- Duncaniella freteri\_nov\_89.919%
- Mogibacterium timidum
- Acinetobacter radioresistens
- Treponema denticola
- Bifidobacterium pseudolongum
- Dialister invisus
- Enterobacter asburiae
- Deinococcus geothermalis
- Shigella sonnei
- Aggregatibacter sp.\_HMT\_512
- Pyrinomonas methylaliphatogenes\_nov\_96.809%
- Shigella flexneri
- Fusobacterium nucleatum
- Chryseobacterium yeoncheonense\_nov\_97.484%
- Staphylococcus capitis\_epidermidis
- Enterobacter hormaechei
- Microbacterium maritypicum
- Peptidiphaga sp.\_HMT\_183
- Atlantibacter hermannii
- Chryseobacterium gambrini
- Alloprevotella tannerae
- Oscillospiraceae\_[G-2] bacterium\_MOT-149\_nov\_95.198%
- Rhodococcus qingshengii
- Bacillus albus\_cereus\_luti\_nitratireducens\_paramycoides\_tro ... (6 sp)
- Acinetobacter johnsonii
- Klebsiella pneumoniae
- Proteus mirabilis

Species



F15127.S07 F15127.S08 F15127.S09 F15127.S10 F15127.S11 F15127.S12

Samples

