



- Species
- Enterococcus faecalis
 - Acinetobacter johnsonii
 - Lawsonella clevelandensis
 - Eubacteriales_[G-1] bacterium_MOT-159
 - Acinetobacter lwoffii
 - Eubacteriales_[G-2] bacterium_MOT-162
 - Pseudomonas putida
 - Streptococcus acidominimus
 - Comamonas testosteroni
 - Acidovorax monticola
 - Streptococcus mutans
 - Citrobacter koseri
 - Enterobacter hormaechei
 - Mammaliococcus sciuri
 - Mammaliococcus lentus
 - Bacteroides acidifaciens
 - Lachnospiraceae_[G-14] bacterium_MOT-185
 - Escherichia fergusonii
 - Actinomyces sp._HMT_448
 - Corynebacterium matruchotii
 - Neisseriaceae_[G-1] bacterium_MOT-031
 - Fusobacterium nucleatum_subsp._vincentii
 - Streptococcus sp._MOT-012
 - Fusobacterium nucleatum
 - Porphyromonas endodontalis
 - Erysipelotrichaceae_[G-1] bacterium_MOT-189
 - Methylobacterium goeisingense
 - Magnetospirillum magnetotacticum
 - Parabacteroides distasonis
 - Klebsiella pneumoniae
 - Proteus mirabilis
 - Lysinibacillus sphaericus
 - Enterobacter mori
 - Bacillus subtilis
 - Lachnospiraceae_[G-11] bacterium_MOT-177
 - Hungatella hathewayi
 - Muribaculaceae_[G-2] bacterium_MOT-104
 - Microbacterium maritipicum
 - Enterobacter asburiae
 - Sphingobacterium multivorum
 - Kosakonia sacchari
 - Citrobacter amalonaticus
 - Leucobacter chromiirestiens
 - Acidovorax ebreus
 - Shigella flexneri
 - Streptococcus oralis_subsp._tigurinus_clade_071
 - Lactobacillus johnsonii
 - Muribaculaceae_[G-1] bacterium_MOT-129
 - Aggregatibacter sp._HMT_512
 - Dialister invisus
 - Chryseobacterium gambrini
 - Bifidobacterium pseudolongum
 - Staphylococcus warneri
 - Shigella sonnei
 - Atlantibacter hermannii
 - Deinococcus geothermalis
 - Limosilactobacillus reuteri
 - Klebsiella quasipneumoniae
 - Enterobacter kobei
 - Akkermansia muciniphila
 - Enemella evansiae
 - Staphylococcus ureilyticus
 - Streptococcus danieliae
 - Enterococcus casseliflavus
 - Janibacter melonis
 - Phocaeicola sartorii
 - Streptococcus gordonii
 - Streptococcus thoraltensis
 - Peptostreptococcaceae_[X1][G-4] bacterium_HMT_369
 - Ligilactobacillus murinus
 - Cutibacterium acnes
 - Methylobacterium brachiatum
 - Cetobacterium somerae
 - Sphingomonas yabuuchiiae
 - Acinetobacter radioresistens
 - Acidovorax temperans
 - Rhodococcus qingshengii
 - Streptococcus oralis
 - Enterococcus gallinarum
 - Chryseobacterium yeoncheonense_nov_97.484%
 - Duncaniella freteri_nov_90.184%
 - Duncaniella freteri_nov_93.712%
 - Duncaniella freteri_nov_93.699%
 - Muribaculaceae_[G-1] bacterium_MOT-129_nov_90.816%
 - Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.089%
 - Enterobacter mori_nov_97.951%
 - Duncaniella freteri_nov_91.039%
 - Leptotrichia hofstadii_nov_96.970%
 - Lachnospiraceae_[G-3] bacterium_MOT-168_nov_94.792%
 - Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.319%
 - Acinetobacter johnsonii_nov_97.737%
 - Duncaniella freteri_nov_89.919%
 - Pyrinomonas methylaliphatogenes_nov_96.809%
 - Duncaniella freteri_nov_89.135%
 - Staphylococcus saprophyticus_xylosum
 - Bacillus albus_cereus_luti_nitratireducens_paramycoideus_tro...(6 sp)
 - Bradyrhizobium archetypum_australiense_cajani_japonicum_liaoningense
 - Staphylococcus capitis_epidermidis
 - Staphylococcus argenteus_aureus_roterodami
 - Sphingomonas aquatilis_melonis

F13506.S05
F13506.S06
F13506.S07
F13506.S08
F13506.S09
F13506.S10
F13506.S11
F13506.S12
F15127.S07
F15127.S08
F15127.S09
F15127.S10
F15127.S11
F15127.S12
F15127.S13
F15127.S14
F15127.S15
F15127.S16
F15127.S17
F15127.S18

Samples