



- Lactobacillus johnsonii
- Bacteroides acidifaciens
- Sporosarcina sp._MOT-205
- Enterococcus hirae
- Mammalicoccus lentus
- Jeotgalicoccus halotolerans
- Sporosarcina sp._MOT-204
- Ligilactobacillus murinus
- Corynebacterium mastitidis
- Streptococcus acidominimus
- Muribacter sp._MOT-143
- Lactococcus lactis
- Streptococcus danieliae
- Staphylococcus ureilyticus
- Rodentibacter pneumotropicus
- Limosilactobacillus reuteri
- Mammaliicoccus sciuri
- Atopostipes sp._MOT-201
- Acinetobacter johnsonii
- Staphylococcus saprophyticus
- Bifidobacterium pseudolongum
- Erysipelotrichaceae_[G-1] bacterium_MOT-189
- Pediococcus pentosaceus
- Corynebacterium stationis
- Porphyromonas sp._MOT-131
- Enterococcus faecalis
- Atopostipes sp._MOT-202
- Streptococcus thoraltensis
- Helicobacter ganmani
- Faecalibaculum rodentium
- Weeksellaceae_[G-1] bacterium_MOT-126
- Muribacter muris
- Savagea faecisuis_nov_92.585%
- Clostridium collagenovorans_nov_80.258%
- Gloeobacter kilaueensis_nov_84.668%
- Sporosarcina sp._MOT-205_nov_95.565%
- Prevotella sp._HMT_820_nov_89.980%
- Kineothrix alysoides_nov_86.667%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_89.431%
- Lachnospiraceae_[G-11] bacterium_MOT-177_nov_96.066%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_83.636%
- Lachnospiraceae_[G-12] bacterium_MOT-180_nov_89.613%
- Prevotella shahii_nov_86.667%
- Lachnospiraceae_[G-7] bacterium_MOT-172_nov_93.075%
- Caproicibacter fermentans_nov_89.662%
- Prevotella sp._HMT_317_nov_90.041%
- Staphylococcus edaphicus_saprophyticus
- Staphylococcus saprophyticus_xylosus
- Pasteurella_Rodentibacter caecimuris_heylii
- Gloeobacter multispecies_sppn4_2_nov_83.144%

F31878.S10

F31878.S09

F31878.S08

F31878.S07

F31878.S06

F31878.S21

F31878.S20

F31878.S19

F31878.S18

F31878.S17

Samples

Species