



- Porphyromonas sp.\_MOT-131  
Pediococcus pentosaceus  
Streptococcus acidominimus  
Streptococcus danieliae  
Eubacteriales\_[G-4] bacterium\_MOT-164  
Enterococcus faecalis  
Corynebacterium stationis  
Mammaliicoccus lentus  
Bifidobacterium pseudolongum  
Mollicutes\_[G-1] bacterium\_MOT-186  
Lactococcus lactis  
Lactobacillus johnsonii  
Streptococcus thoraltensis  
Stenotrophomonas [Pseudomonas] hibiscicola  
Staphylococcus ureilyticus  
Weissella cibaria  
Staphylococcus saprophyticus  
Anaerococcus sp.\_HMT\_290  
Triticum aestivum  
Muribacter sp.\_MOT-143  
Rodentibacter pneumotropicus  
Streptococcus thermophilus  
Moraxella osloensis  
Enterococcus gallinarum  
Limosilactobacillus reuteri  
Staphylococcus equorum  
Massilia aurea  
Psychrobacter alimentarius  
Dubosiella newyorkensis  
Atopostipes sp.\_MOT-201  
Ligilactobacillus animalis  
Lachnospiraceae\_[G-14] bacterium\_MOT-185  
Ligilactobacillus murinus  
Eubacteriales\_[G-2] bacterium\_MOT-162  
Cutibacterium granulosum  
Corynebacterium mastitidis  
Cutibacterium acnes  
Clostridium disporicum  
Stenotrophomonas maltophilia  
Streptomyces aculeolatus  
Lachnospiraceae\_[G-11] bacterium\_MOT-177  
Pseudomonas helleri  
Lactiseibacillus rhamnosus  
Jeotgalicoccus halotolerans  
Levilactobacillus brevis  
Gemella sp.\_MOT-033  
Corynebacterium ammoniagenes  
Secundilactobacillus paracollinoides  
Leptothrix sp.\_HMT\_025  
Weissella paramesenteroides  
Delftia acidovorans  
Carnobacteriaceae\_[G-1] bacterium\_MOT-197  
Rothia nasimurium  
Pelomonas saccharophila  
Ralstonia sp.\_HMT\_406  
Actinidia eriantha  
Akkermansia muciniphila  
Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_88.105%  
Duncaniella freteri\_nov\_93.293%  
Turicibacter sanguinis\_nov\_95.923%  
Enterococcus faecalis\_nov\_95.825%  
Fusobacterium varium\_nov\_96.696%  
Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_88.867%  
Oribacterium parvum\_nov\_89.770%  
Duncaniella freteri\_nov\_89.775%  
Alistipes senegalensis\_nov\_93.443%  
Erysipelotrichaceae\_[G-1] bacterium\_MOT-189\_nov\_88.798%  
Peptococcus sp.\_HMT\_168\_nov\_84.866%  
Duncaniella freteri\_nov\_89.718%  
Fusicatenibacter saccharivorans\_nov\_90.526%  
Oscillospiraceae\_[G-6] bacterium\_MOT-153\_nov\_91.631%  
Actinidia eriantha\_nov\_97.011%  
Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_89.000%  
Lachnospiraceae\_[G-14] bacterium\_MOT-185\_nov\_92.719%  
Fusobacterium perfoetens\_nov\_91.126%  
Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_88.423%  
Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_85.887%  
Duncaniella freteri\_nov\_88.934%  
Duncaniella freteri\_nov\_93.699%  
Latilactobacillus curvatus\_graminis  
Sphingomonas aquatilis\_melonis  
Staphylococcus saprophyticus\_xylosus

Species

F17722.S06  
F17722.S07  
F17722.S08  
F17722.S09  
F17722.S10  
F18971.S06  
F18971.S07  
F18971.S08  
F18971.S09  
F18971.S10

Samples