



- Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.623%
- Alistipes finegoldii_nov_93.608%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.400%
- Duncaniella freteri_nov_88.978%
- Duncaniella freteri_nov_88.330%
- Marinisporobacter balticus_nov_82.692%
- Clostridium disparicum
- Eubacteriales_[G-1] bacterium_MOT-158
- Longibaculum muris_nov_86.957%
- Beduini massiliensis_nov_87.705%
- Mucispirillum schaedleri
- Flavonifractor plautii_nov_92.308%
- Lachnospiraceae_[G-3] bacterium_MOT-168_nov_94.792%
- Caproiciproducens galactitolivorans_nov_83.789%
- Clostridiales_[F-1][G-2] bacterium_HMT_402_nov_82.255%
- Muricomes intestini_nov_89.583%
- Eubacteriales_[G-4] bacterium_MOT-164
- Bacteroides acidifaciens
- Duncaniella freteri_nov_86.948%
- Bifidobacterium pseudolongum
- Eubacteriales_[G-4] bacterium_MOT-164_nov_97.655%
- Duncaniella freteri_nov_89.069%
- Enterococcus casseliflavus
- Neglectibacter timonensis_nov_95.325%
- Alistipes putredinis_nov_94.444%
- Oscillospiraceae_[G-3] bacterium_MOT-150_nov_92.931%
- Eubacteriales_[G-2] bacterium_MOT-162
- Sporobacter termitidis_nov_87.580%
- Hatthewaya proteolytica_nov_83.514%
- Enterococcus faecalis
- Acutalibacter muris
- Lachnoclostridium [Clostridium] aminophilum_nov_87.318%
- Eubacteriales_[G-1] bacterium_MOT-159
- Acetivibrio cellulolyticus_nov_83.405%
- Duncaniella freteri_nov_92.915%
- Faecalicatena multispecies_sppn9_2_nov_94.363%
- Lachnospiraceae_[G-7] bacterium_MOT-172_nov_93.279%
- Oscillospiraceae_[G-3] bacterium_MOT-150_nov_93.582%
- Eubacterium xylanophilum_nov_91.075%
- Faecalicatena multispecies_sppn7_2_nov_91.858%
- Lacrimispora xylanolytica_nov_94.363%
- Limosilactobacillus reuteri
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_90.816%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.290%
- Ileibacterium valens
- Longibaculum muris_nov_90.289%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.640%
- Turicibacter sanguinis_nov_95.923%
- Duncaniella freteri_nov_89.135%
- Duncaniella freteri_nov_89.697%
- Duncaniella freteri_nov_87.221%
- Prevotella sp._HMT_317_nov_90.244%
- Phocaeicola sartorii
- Enterobacter hormaechei
- Duncaniella freteri_nov_87.071%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.822%
- Eubacteriales_[G-4] bacterium_MOT-164_nov_97.655%
- Duncaniella freteri_nov_93.699%
- Alistipes sp._MOT-127
- Lachnospiraceae_[G-7] bacterium_MOT-172_nov_91.718%
- Lachnoclostridium [Clostridium] scindens_nov_89.027%
- Anaeroplasma abactoclasticum_nov_86.538%
- Oscillospiraceae_[G-3] bacterium_MOT-150_nov_91.134%
- Lachnospiraceae_[G-3] bacterium_MOT-168_nov_92.902%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_86.373%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_86.000%
- Duncaniella freteri_nov_92.653%
- Lactobacillus intestinalis
- Parabacteroides distasonis
- Parabacteroides distasonis_nov_97.938%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.621%
- Alistipes senegalensis_nov_93.648%
- Gluciribacter canis_nov_93.305%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_87.576%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.577%
- Muribaculum intestinale
- Erysipelatoclostridium [Clostridium] innocuum_nov_88.270%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_86.600%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.089%
- Parasutterella excrementihominis
- Duncaniella freteri_nov_87.169%
- Duncaniella freteri_nov_91.039%
- Duncaniella freteri_nov_90.184%
- Duncaniella freteri_nov_86.290%
- Alloprevotella sp._HMT_473_nov_89.366%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_85.686%
- Duncaniella freteri_nov_86.842%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_83.636%
- Prevotella shahii_nov_87.602%
- Duncaniella freteri_nov_91.853%
- Alloprevotella sp._HMT_473_nov_89.634%
- Duncaniella freteri_nov_90.612%
- Erysipelotrichaceae_[G-1] bacterium_MOT-189
- Ligilactobacillus murinus
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_84.929%
- Parabacteroides goldsteinii
- Akkermansia muciniphila
- Duncaniella freteri_nov_91.429%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_89.431%
- Muribaculaceae_[G-1] bacterium_MOT-129

Species

F15127.S25
F15127.S26
F15127.S27
F15127.S28
F15127.S29
F15127.S30

Samples