



- Lachnospiraceae_[G-11] bacterium_MOT-178
- Alistipes sp._MOT-127
- Turicimonas muris
- Bacteroides caecimuris
- Blautia caecimuris
- Prevotella sp._MOT-128
- Lactobacillus taiwanensis
- Flavonifractor plautii
- Oscillospiraceae_[G-2] bacterium_MOT-149
- Lachnospiraceae_[G-14] bacterium_MOT-183
- Robinsoniella peoriensis
- Lachnospiraceae_[G-9] bacterium_MOT-174
- Bacteroides stercorisoris
- Clostridium tertium
- Muribaculum intestinale
- Lachnospiraceae_[G-14] bacterium_MOT-184
- Parabacteroides goldsteinii
- Akkermansia muciniphila
- Kineothrix alysoides_nov_95.227%
- Anaerotruncus rubiinfantis_nov_92.760%
- Mediterraneibacter [Ruminococcus] gnavus_nov_93.424%
- Alistipes putredinis_nov_95.887%
- Oscillibacter valericigenes_nov_95.260%
- Kineothrix alysoides_nov_97.279%
- Pseudobutyrvibrio ruminis_nov_91.176%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_91.522%
- Saccharofermentans acetigenes_nov_88.764%
- Lacrimispora indolis_nov_90.724%
- Lawsonibacter asaccharolyticus_nov_97.973%
- Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.358%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_90.870%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_93.043%
- Odoribacter splanchnicus_nov_93.939%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.506%
- Ruthenibacterium lactatiformans_nov_97.045%
- Lachnospiraceae_[G-11] bacterium_MOT-178_nov_97.978%
- Lachnospiraceae_[G-9] bacterium_MOT-174_nov_96.388%
- Lachnospiraceae_[G-14] bacterium_MOT-185_nov_96.599%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_90.022%
- Prevotellamassilia timonensis_nov_94.168%
- Maiihella massiliensis_nov_92.094%
- Anaerotruncus rubiinfantis_nov_92.517%
- Neglectibacter timonensis_nov_97.727%
- Lachnospiraceae_[G-11] bacterium_MOT-176_nov_95.946%
- Acetatifactor muris_nov_96.145%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.056%
- Lachnospiraceae_[G-6] bacterium_MOT-171_nov_95.238%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_92.208%
- Alistipes putredinis_nov_96.529%
- Phocaea massiliensis_nov_95.682%
- Muribaculum intestinale_nov_93.737%
- Butyricoccus pullicaecorum_nov_94.382%
- Eubacteriales_[G-3] bacterium_MOT-163_nov_93.679%
- Lachnoclostridium [Clostridium] populeti_nov_94.331%
- Lachnospiraceae_[G-12] bacterium_MOT-179_nov_94.796%
- Lachnospiraceae_[G-14] bacterium_MOT-184_nov_95.238%
- Lachnospiraceae_[G-11] bacterium_MOT-176_nov_97.297%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.919%
- Acetatifactor muris_nov_92.551%
- Kineothrix alysoides_nov_97.059%
- Lachnoclostridium [Clostridium] populeti_nov_92.986%
- Duncaniella freteri_nov_90.456%
- Kineothrix alysoides_nov_93.682%
- Roseburia faecis_nov_97.964%
- Duncaniella freteri_nov_93.103%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_93.074%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_92.441%
- Lacrimispora xylanolytica_nov_97.285%
- Culturomica massiliensis_nov_93.709%
- Kineothrix alysoides_nov_95.928%
- Lachnospiraceae_[G-10] bacterium_MOT-175_nov_96.372%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_92.191%
- Lachnospiraceae_[G-10] bacterium_MOT-175_nov_92.174%
- Lachnoclostridium pacaense_nov_96.825%
- Eubacterium coprostanoligenes_nov_95.485%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_91.974%
- Lachnospiraceae_[G-12] bacterium_MOT-179_nov_92.534%
- Lachnospiraceae_[G-14] bacterium_MOT-184_nov_94.989%
- Pseudoflavonifractor capillosus_nov_95.721%
- Anaerotignum lactatifermentans_nov_95.270%
- Caecibacterium sporiformans_nov_95.045%
- Alistipes timonensis_nov_97.831%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.946%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_94.157%
- Alistipes senegalensis_nov_95.228%
- Oscillospiraceae_[G-4] bacterium_MOT-151_nov_91.723%
- Lachnospiraceae_[G-11] bacterium_MOT-177_nov_97.523%
- Enterobacter asburiae_cancerogenus_cloacae_hormaechei
- Lachnospiraceae_[G-12] bacterium_MOT-179_bacterium_MOT-184
- multigenus multispecies_sppn10_2_nov_95.918%
- multigenus multispecies_sppn12_2_nov_96.304%
- multigenus multispecies_sppn13_5_nov_94.570%
- multigenus multispecies_sppn17_2_nov_95.928%
- Bacteroidetes_[G-3] multispecies_sppn2_2_nov_87.554%
- Lachnospiraceae_[G-11] multispecies_sppn4_2_nov_96.847%
- multigenus multispecies_sppn5_2_nov_97.279%
- Bacteroides multispecies_sppn6_2_nov_96.312%
- multigenus multispecies_sppn7_2_nov_92.777%
- multigenus multispecies_sppn8_3_nov_95.011%
- multigenus multispecies_sppn9_2_nov_93.002%

Species

F8810.S01
F8810.S02
F8810.S03
F8810.S13
F8810.S14
F8810.S15
F8810.S07
F8810.S08
F8810.S09
F8810.S19
F8810.S20
F8810.S21

Samples