



F8810.S01
F8810.S02
F8810.S03
F8810.S13
F8810.S14
F8810.S15

Samples

- Species
- Alistipes putredinis_nov_96.529%
 - Bacteroides caecimuris
 - Neglectibacter timonensis_nov_97.727%
 - Oscillospiraceae_[G-2] bacterium_MOT-149_nov_94.157%
 - Lachnospiraceae_[G-14] bacterium_MOT-184_nov_94.989%
 - Muribaculum intestinale_nov_93.737%
 - Lawsonibacter asaccharolyticus_nov_97.973%
 - Lachnospiraceae_[G-9] bacterium_MOT-174_nov_96.388%
 - Anaerosporebacter mobilis_nov_95.000%
 - Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.919%
 - multigenus multispecies_sppn17_2_nov_95.928%
 - Anaerotaenia torta_nov_97.273%
 - Lachnospiraceae_[G-12] bacterium_MOT-179_nov_92.534%
 - Coprococcus catus_nov_94.570%
 - Bacteroides acidifaciens_acidofaciens
 - Lachnospiraceae_[G-14] bacterium_MOT-184_nov_95.692%
 - multigenus multispecies_sppn18_2_nov_92.063%
 - Anaerotignum lactatifermentans_nov_95.270%
 - Petrocella atlantisensis_nov_87.810%
 - Lacrimispora xylanolytica_nov_97.285%
 - Lachnospiraceae_[G-14] bacterium_MOT-184_nov_95.405%
 - Lachnospiraceae_[G-14] bacterium_MOT-183
 - Alistipes putredinis_nov_95.887%
 - Duncaniella freteri_nov_90.456%
 - Anaerotruncus rubiinfantis_nov_92.760%
 - Muribaculaceae_[G-2] bacterium_MOT-104_nov_90.870%
 - multigenus multispecies_sppn23_2_nov_96.818%
 - Muribaculaceae_[G-2] bacterium_MOT-104_nov_91.991%
 - Bacteroides stercorisoris
 - Prevotellamassilia timonensis_nov_94.168%
 - Alistipes sp._MOT-127
 - Eubacteriales_[G-2] bacterium_MOT-162_nov_95.260%
 - Peptococcaceae_[G-1] bacterium_MOT-146
 - Clostridium oryzae_nov_88.889%
 - Lawsonibacter asaccharolyticus
 - Saccharofermentans acetigenes_nov_88.764%
 - Clostridium disporicum_saudiense
 - Ruthenibacterium lactatiformans_nov_97.045%
 - Alistipes putredinis_nov_95.879%
 - Butyricoccus pullicaecorum_nov_94.820%
 - Neglectibacter timonensis_nov_97.500%
 - Alistipes multispecies_sppn12_2_nov_96.304%
 - Kineothrix alysoides_nov_95.928%
 - Alistipes timonensis
 - Bacteroides multispecies_sppn6_2_nov_96.312%
 - Oscillospiraceae_[G-4] bacterium_MOT-151_nov_91.723%
 - Lachnospiraceae_[G-11] bacterium_MOT-178_nov_97.978%
 - Odoribacter splanchnicus_nov_93.939%
 - multigenus multispecies_sppn8_3_nov_95.011%
 - Lacrimispora indolis_nov_90.724%
 - Muribaculaceae_[G-2] bacterium_MOT-104_nov_93.074%
 - Oscillibacter valericiigenes_nov_95.260%
 - Culturomica massiliensis_nov_93.709%
 - Eubacterium coprostanoligenes_nov_95.485%
 - Prevotella sp._MOT-128
 - Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.056%
 - Maiihella massiliensis_nov_92.094%
 - Alistipes timonensis_nov_97.831%
 - multigenus multispecies_sppn5_2_nov_97.279%
 - Muribaculaceae_[G-2] bacterium_MOT-104_nov_92.191%
 - multigenus multispecies_sppn10_2_nov_95.918%
 - Lachnospiraceae_[G-9] bacterium_MOT-174
 - Lachnospiraceae_[G-6] bacterium_MOT-171_nov_95.238%
 - Mediterraneibacter [Ruminococcus] gnavus_nov_93.424%
 - Kineothrix alysoides_nov_97.279%
 - Lachnoclostridium [Clostridium] populeti_nov_92.986%
 - Lachnospiraceae_[G-11] multispecies_sppn4_2_nov_96.847%
 - Lachnospiraceae_[G-11] bacterium_MOT-176_nov_95.946%
 - multigenus multispecies_sppn7_2_nov_92.777%
 - Lachnospiraceae_[G-11] bacterium_MOT-176_nov_97.297%
 - multigenus multispecies_sppn13_5_nov_94.570%
 - Caecibacterium sporiformans_nov_95.045%
 - Lachnospiraceae_[G-10] bacterium_MOT-175_nov_92.174%
 - Lachnospiraceae_[G-14] bacterium_MOT-185_nov_96.599%
 - Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.358%
 - Lachnoclostridium [Clostridium] populeti_nov_94.331%
 - Acetatifactor muris_nov_96.145%
 - Kineothrix alysoides_nov_95.227%
 - Lachnospiraceae_[G-11] bacterium_MOT-178
 - Pseudoflavonifractor capillosus_nov_95.721%
 - Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.506%
 - Lachnospiraceae_[G-14] bacterium_MOT-184
 - Phoceia massiliensis_nov_95.682%
 - Roseburia faecis_nov_97.964%
 - Lactobacillus taiwanensis
 - Oscillospiraceae_[G-2] bacterium_MOT-149
 - Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.946%
 - Lachnospiraceae_[G-11] bacterium_MOT-177_nov_97.523%
 - Acetatifactor muris_nov_92.551%
 - Eubacteriales_[G-3] bacterium_MOT-163_nov_93.679%
 - Pseudobutyrvibrio ruminis_nov_91.176%
 - Lachnospiraceae_[G-12] bacterium_MOT-179_bacterium_MOT-18
 - Lachnoclostridium pacaense_nov_96.825%
 - Alistipes senegalensis_nov_95.228%
 - Muribaculaceae_[G-2] bacterium_MOT-104_nov_91.974%
 - Bacteroidetes_[G-3] multispecies_sppn2_2_nov_87.554%
 - Kineothrix alysoides_nov_93.682%
 - multigenus multispecies_sppn9_2_nov_93.002%
 - Kineothrix alysoides_nov_97.059%
 - Lachnospiraceae_[G-12] bacterium_MOT-179_nov_94.796%