



Hydrogenoanaerobacterium saccharovorans_nov_93.636%
multigenus multispecies_sppn14_2_nov_82.889%
Oscillospiraceae [G-2] bacterium_MOT-149_nov_96.171%
Flavonifractor plautii
Alistipes sp._MOT-127_nov_91.775%
Ruthenibacterium lactatiformans_nov_97.045%
Lawsonibacter asaccharolyticus_nov_95.730%
Butyrivibrio coccineus_nov_94.382%
Anaerotruncus rubifantidis_nov_92.517%
Oscillospiraceae [G-2] bacterium_MOT-149_nov_95.281%
multigenus multispecies_sppn10_2_nov_95.918%
Oscillospiraceae [G-4] bacterium_MOT-151_nov_91.723%
Muribaculaceae [G-2] bacterium_MOT-104_nov_91.106%
Anaerosporbacter mobilis_nov_95.000%
Saccharofermentans acetigenus_nov_88.764%
multigenus multispecies_sppn9_2_nov_93.002%
Bacteroidetes [G-3] multispecies_sppn2_2_nov_87.554%
Eubacteriales [G-1] bacterium_MOT-158
Faecalicatena orotica_nov_95.238%
Lachnospiraceae [G-12] bacterium_MOT-179_bacterium_MOT-1
Neglectibacter timonensis_nov_97.500%
Alistipes putredinis_nov_95.879%
Eisenbergiella massiliensis_nov_96.599%
Bacteroides acidifaciens_acidofaciens
Anaerotignum lactatifermentans_nov_97.523%
Lachnospiraceae [G-14] bacterium_MOT-184_nov_94.989%
Alistipes timonensis_nov_97.831%
Roseburia faecis
Eubacterium coprostanoligenes_nov_95.701%
Oscillospiraceae [G-2] bacterium_MOT-149_nov_95.946%
Muribaculaceae [G-2] bacterium_MOT-104_nov_91.991%
Kineothrix alysioides_nov_93.651%
Lachnospiraceae [G-14] bacterium_MOT-185_nov_96.599%
Muribaculaceae [G-2] bacterium_MOT-104_nov_90.870%
Bacteroides acidofaciens
Eubacteriales [G-1] multispecies_sppn15_2_nov_97.511%
Culturomica massiliensis_nov_93.709%
Oscillospiraceae [G-2] bacterium_MOT-149
Mediterraneibacter [Ruminococcus] gnavus_nov_93.424%
Anaerotruncus rubifantidis_nov_92.760%
Mailhella massiliensis_nov_92.094%
Oscillospiraceae [G-2] bacterium_MOT-149_nov_95.506%
Phoceea massiliensis_nov_95.682%
Anaerotignum lactatifermentans_nov_95.270%
Oscillospiraceae [G-2] bacterium_MOT-149_nov_95.056%
Muribaculaceae [G-2] bacterium_MOT-104_nov_92.191%
Bacteroides stercorisoris
Muribaculaceae [G-2] bacterium_MOT-104_nov_91.974%
Alistipes putredinis_nov_95.887%
Lachnospiraceae [G-14] bacterium_MOT-184
Lachnospiraceae [G-14] bacterium_MOT-183
Muribaculaceae [G-2] bacterium_MOT-104_nov_92.441%
Muribaculaceae [G-2] bacterium_MOT-104_nov_92.208%
Blautia hansenii_hominis_marasmi
Clostridium intestinale
Muribaculum tertium_nov_92.688%
multigenus multispecies_sppn8_3_nov_95.011%
Lachnospiraceae [G-9] bacterium_MOT-174
Oscillibacter valericigenes_nov_95.260%
Muribaculaceae [G-2] bacterium_MOT-104_nov_89.462%
Eubacterium coprostanoligenes_nov_95.485%
Lachnospiraceae [G-11] multispecies_sppn4_2_nov_96.847%
multigenus multispecies_sppn5_2_nov_97.279%
Lachnospiraceae [G-11] bacterium_MOT-178
Lachnospiraceae [G-9] bacterium_MOT-174_nov_96.388%
Lachnospiraceae [G-14] bacterium_MOT-184_nov_95.238%
multigenus multispecies_sppn7_2_nov_92.777%
Lachnospiraceae [G-12] bacterium_MOT-179_nov_92.534%
Pseudoflavonifractor capillosus_nov_95.721%
Lachnospiraceae [G-10] bacterium_MOT-175_nov_92.174%
Lachnoclostridium [Clostridium] populeti_nov_94.331%
Lachnoclostridium pacaeense_nov_96.825%
Muribaculum intestinale_nov_93.737%
Lachnospiraceae [G-10] bacterium_MOT-175_nov_96.372%
Lacrimispora xylanolytica_nov_97.285%
Bacteroides multispecies_sppn6_2_nov_96.312%
Alistipes sp._MOT-127
Parabacteroides goldsteinii
Alistipes putredinis_nov_96.529%
Muribaculaceae [G-2] bacterium_MOT-104_nov_93.043%
Muribaculaceae [G-2] bacterium_MOT-104_nov_90.022%
Alistipes multispecies_sppn12_2_nov_96.304%
Duncaniella freteri_nov_90.456%
Muribaculum intestinale
Robinsoniella periensis
Enterobacter asburiae_cancerogenus_cloacae_hormaechei
Muribaculaceae [G-2] bacterium_MOT-104_nov_93.074%
Blautia caecimuris
Acetatifactor muris_nov_92.551%
Lachnospiraceae [G-11] bacterium_MOT-176_nov_97.297%
Lachnospiraceae [G-6] bacterium_MOT-171_nov_95.238%
Lacrimispora indolis_nov_90.724%
Prevotella sp._MOT-128
Muribaculaceae [G-1] bacterium_MOT-129_nov_91.522%
Prevotellamassilia timonensis_nov_94.168%
Turicomonas muris
Kineothrix alysioides_nov_95.928%
Akkermansia muciniphila
Duncaniella freteri_nov_93.103%
Bacteroides caecimuris

Species

Samples