



- Akkermansia muciniphila
- Mammaliococcus lentus
- Lactobacillus johnsonii
- Acinetobacter radioresistens
- Eubacteriales_[G-3] bacterium_MOT-163
- Oscillospiraceae_[G-7] bacterium_MOT-154
- Klebsiella pneumoniae
- Oscillospiraceae_[G-4] bacterium_MOT-151
- Acutalibacter muris
- Eubacteriales_[G-1] bacterium_MOT-159
- Streptococcus thoraltensis
- Lachnospiraceae_[G-14] bacterium_MOT-185
- Bifidobacterium pseudolongum
- Eubacteriales_[G-2] bacterium_MOT-162
- Faecalibaculum rodentium
- Enterococcus faecalis
- Eubacteriales_[G-4] bacterium_MOT-164
- Mollicutes_[G-2] bacterium_MOT-187_nov_90.607%
- Phocaea massiliensis_nov_91.471%
- Oscillospiraceae_[G-4] bacterium_MOT-151_nov_95.842%
- Duncanella freteri_nov_93.712%
- Adlercreutzia equolifaciens_nov_94.577%
- Paludicola psychrotolerans_nov_87.759%
- Lachnospiraceae_[G-7] bacterium_MOT-172_nov_91.097%
- Lachnospiraceae_[G-9] bacterium_MOT-174_nov_89.918%
- Acutalibacter muris_nov_94.227%
- Lachnospiraceae_[G-11] bacterium_MOT-178_nov_95.519%
- Streptococcus azizii_nov_95.171%
- Lachnoclostridium [Clostridium] polysaccharolyticum_nov_90.437%
- Lawsonibacter asaccharolyticus_nov_93.996%
- Lawsonibacter asaccharolyticus_nov_91.116%
- Ihubacter massiliensis_nov_94.572%
- Lachnospiraceae_[G-2] bacterium_HMT_096_nov_91.632%
- Eubacterium xylanophilum_nov_91.075%
- Mollicutes_[G-2] bacterium_MOT-187_nov_94.892%
- Lacrimispora xylanolytica_nov_94.363%
- Muricomes intestini_nov_89.375%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_88.577%
- Lachnospiraceae_[G-5] bacterium_MOT-170_nov_97.904%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_87.576%
- Duncanella freteri_nov_93.699%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.319%
- Oscillospiraceae_[G-3] bacterium_MOT-150_nov_91.511%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.010%
- Lachnospiraceae_[G-7] bacterium_MOT-172_nov_91.097%
- Faecalicatena fissicatena_nov_95.407%
- Neglectibacter timonensis_nov_95.325%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_87.174%
- Eubacteriales_[G-3] bacterium_MOT-163_nov_85.944%
- Duncanella freteri_nov_87.424%
- Lachnospiraceae_[G-11] bacterium_MOT-176_nov_89.858%
- Christensenella hongkongensis_nov_85.122%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_85.887%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.022%
- Oscillospiraceae_[G-3] bacterium_MOT-150_nov_93.582%
- Anaerotignum faecicola_nov_85.287%
- Prevotella shahii_nov_87.602%
- Oscillospiraceae_[G-1] bacterium_MOT-147_nov_96.674%
- Turicibacter sanguinis_nov_95.923%
- Acetivibrio cellulolyticus_nov_83.801%
- Duncanella freteri_nov_89.775%
- Lachnospiraceae_[G-10] bacterium_MOT-175_nov_90.369%
- Lachnospiraceae_[G-2] bacterium_MOT-167_nov_88.773%
- Breznakia pachnodae_nov_82.824%
- Acetivibrio cellulolyticus_nov_83.153%
- Phocaea massiliensis_nov_89.914%
- Anaerotruncus rubiinfantis_nov_92.708%
- Adlercreutzia caecimuris_nov_92.291%
- Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.979%
- Marinisporobacter balticus_nov_82.692%
- Oscillospiraceae_[G-4] bacterium_MOT-151_nov_94.179%
- Eubacteriales_[G-4] bacterium_MOT-164_nov_97.228%
- Flavonifractor plautii_nov_92.308%
- Sporobacter termitidis_nov_87.580%
- Eubacteriales_[G-4] bacterium_MOT-164_nov_97.228%
- Longibaculum muris_nov_93.361%
- Faecalicoccus acidiformans_nov_89.600%
- Lachnospiraceae_[G-6] bacterium_MOT-171_nov_93.305%
- Christensenella massiliensis_nov_88.041%
- Adlercreutzia caecimuris_nov_89.009%
- Butyricoccus pullicaecorum_nov_85.093%
- Roseburia hominis_nov_91.476%
- Eubacteriales_[G-1] bacterium_MOT-159_nov_92.161%
- Hydrogenoanaerobacterium saccharovorans_nov_88.589%
- Agrococcus versicolor_nov_83.227%
- Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.561%
- Clostridiales_[F-1][G-1] bacterium_HMT_093_nov_84.086%
- Eubacteriales_[G-3] bacterium_MOT-163_nov_87.952%
- Flavonifractor plautii_nov_93.555%
- Ihubacter massiliensis_nov_90.644%
- Adlercreutzia muris_nov_88.961%
- Neglectibacter timonensis_nov_94.490%
- Oscillospiraceae_[G-4] bacterium_MOT-151_nov_96.881%
- Hydrogenoanaerobacterium saccharovorans_nov_87.759%
- Eubacteriales_[G-3] bacterium_MOT-163_nov_85.825%
- Anaeromassilibacillus senegalensis_nov_92.489%
- Butyricoccus pullicaecorum_nov_85.644%
- Acetivibrio cellulolyticus_nov_85.776%
- Staphylococcus saprophyticus_xylosus
- Faecalicatena multispecies_sppn8_2_nov_92.067%

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Samples

Species