



Faecalibaculum rodentium
Clostridium disporicum
Akkermansia muciniphila
Lactococcus lactis
Eubacteriales_[G-1] bacterium_MOT-159
Erysipelatoclostridium [Clostridium] cocleatum
Eubacteriales_[G-3] bacterium_MOT-163
Eubacteriales_[G-2] bacterium_MOT-162
Streptococcus danieliae
Lactococcus cremoris
Bacteroides acidifaciens
Lactobacillus gasseri
Ligilactobacillus murinus
Lachnospiraceae_[G-14] bacterium_MOT-185
Eubacteriales_[G-4] bacterium_MOT-164
Phocaeicola vulgatus
Adlercreutzia mucosicola
Romboutsia ilealis
Oscillospiraceae_[G-6] bacterium_MOT-153
Adlercreutzia muris
Adlercreutzia caecimuris
Eubacterium ramulus_nov_91.296%
Lachnospiraceae_[G-3] bacterium_MOT-168_nov_94.653%
Lawsonibacter asaccharolyticus_nov_91.571%
Lancefieldella parvula_nov_91.552%
Lachnospiraceae_[G-9] bacterium_MOT-174_nov_90.234%
Faecalibaculum rodentium_nov_96.571%
Sporobacter termitidis_nov_82.970%
Leifsonia kaffniensis_nov_84.158%
Lachnospiraceae_[G-7] bacterium_MOT-172_nov_91.945%
Phocaea massiliensis_nov_90.297%
Pseudoflavonifractor phocaeensis_nov_95.761%
Anaerotignum aminivorans_nov_93.173%
Lachnospiraceae_[G-9] bacterium_MOT-174_nov_86.957%
Tyzzereella [Clostridium] colinum_nov_88.494%
Lachnospiraceae_[G-7] bacterium_MOT-172_nov_86.508%
Eubacterium coprostanoligenes_nov_91.892%
Lachnospiraceae_[G-7] bacterium_MOT-172_nov_86.047%
Mediterraneibacter [Ruminococcus] torques_nov_95.200%
Lachnoclostridium [Clostridium] polysaccharolyticum_nov_92.664%
Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.590%
Lachnospiraceae_[G-2] bacterium_MOT-167_nov_89.349%
Lachnoclostridium [Clostridium] scindens_nov_89.827%
Eubacterium oxidoreducens_nov_88.846%
Acutalibacter muris_nov_94.264%
Phocaea massiliensis_nov_90.239%
Lachnoclostridium [Clostridium] polysaccharolyticum_nov_88.466%
Clostridium collagenovorans_nov_83.466%
Oscillibacter valericigenes_nov_95.164%
Faecalicatena fissicatena_nov_94.521%
Breznakia pachnodae_nov_83.181%
Hathewayia proteolytica_nov_84.970%
Lachnoclostridium [Clostridium] polysaccharolyticum_nov_90.751%
Eubacterium ramulus_nov_89.362%
Eisenbergiella massiliensis_nov_86.127%
Lachnospiraceae_[G-3] bacterium_MOT-168_nov_96.252%
Lachnospiraceae_[G-9] bacterium_MOT-174_nov_92.216%
Lachnospiraceae_[G-11] bacterium_MOT-178_nov_95.358%
Faecalicatena fissicatena_nov_95.499%
Pseudoflavonifractor phocaeensis_nov_86.756%
Lachnoclostridium phytofermentans_nov_89.057%
Eubacterium coprostanoligenes_nov_90.476%
Clostridiales_[F-1][G-1] bacterium_HMT_093_nov_84.298%
Oscillospiraceae_[G-4] bacterium_MOT-151_nov_94.477%
Clostridiales_[F-1][G-1] bacterium_HMT_093_nov_91.892%
Eisenbergiella massiliensis_nov_87.669%
Lacrimispora xylanolytica_nov_93.969%
Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.276%
Faecalicatena fissicatena_nov_93.580%
Lachnospiraceae_[G-11] bacterium_MOT-178_nov_96.332%
Lachnospiraceae_[G-11] bacterium_MOT-178_nov_93.064%
Lachnospiraceae_[G-6] bacterium_MOT-171_nov_93.307%
Lachnoclostridium [Clostridium] aminophilum_nov_87.476%
Blautia faecicola_nov_89.709%
Lawsonibacter asaccharolyticus_nov_90.421%
Anaeroplasma abactoclasticum_nov_87.352%
Lachnoclostridium [Clostridium] aminophilum_nov_89.961%
Hydrogenoanaerobacterium saccharovorans_nov_88.395%
Eisenbergiella massiliensis_nov_90.805%
Lacrimispora xylanolytica_nov_94.314%
Lachnospiraceae_[G-11] bacterium_MOT-177_nov_95.874%
Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.643%
Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.050%
Eisenbergiella massiliensis_nov_88.697%
Bacteroidetes_[G-3] bacterium_HMT_436_nov_85.575%
Lachnoclostridium [Clostridium] scindens_nov_88.247%
Lachnospiraceae_[G-11] bacterium_MOT-176_nov_95.155%
Lachnospiraceae_[G-6] bacterium_MOT-171_nov_95.050%
Lachnoclostridium [Clostridium] polysaccharolyticum_nov_87.308%
Olsenella phocaeensis_nov_92.172%
Parabacteroides distasonis_nov_97.514%
Faecalicatena orotica_nov_92.218%
Longibaculum muris_nov_93.910%
Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.105%
Lachnospiraceae_[G-6] bacterium_MOT-171_nov_95.644%
Turicibacter sanguinis_nov_95.817%
Fusicatenibacter saccharivorans_nov_91.018%
Desulfovibrio fairfieldensis_nov_89.168%
Lachnospiraceae_[G-14] bacterium_MOT-182_nov_89.784%

F10949.S01

F10949.S02

F10949.S03

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F10949.S05

F10949.S06

Samples

Species