



Bacteroides thetaiotaomicron
Akkermansia muciniphila
Parasutterella excrementihominis
Blautia hominis
Eubacteriales_[G-1] bacterium_MOT-159
Lachnospiraceae_[G-11] bacterium_MOT-177
Oscillospiraceae_[G-7] bacterium_MOT-154
Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.069%
Blautia faecicola_nov_89.749%
Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.549%
Acetivibrio cellulolyticus_nov_83.851%
Lachnospiraceae_[G-10] bacterium_MOT-175_nov_91.085%
Oscillospiraceae_[G-4] bacterium_MOT-151_nov_92.899%
Lachnospiraceae_[G-7] bacterium_MOT-172_nov_87.033%
Anaerostipes caccae_nov_96.058%
Duncaniella freteri_nov_90.323%
Ruminiclostridium cellulolyticum_nov_82.704%
Oscillibacter valericigenes_nov_93.642%
Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.850%
Turicibacter sanguinis_nov_95.635%
Pseudoflavonifractor phocaeensis_nov_95.568%
Acutalibacter muris_nov_88.359%
Oscillospiraceae_[G-2] bacterium_MOT-149_nov_92.731%
Blautia hominis_nov_97.773%
Anaerotignum aminivorans_nov_92.184%
Parasutterella excrementihominis_nov_97.852%
Oscillospiraceae_[G-2] bacterium_MOT-149_nov_94.071%
Lachnospiraceae_[G-11] bacterium_MOT-178_nov_93.064%
Lachnoclostridium [Clostridium] scindens_nov_90.979%
Oscillospiraceae_[G-2] bacterium_MOT-149_nov_92.941%
Duncaniella freteri_nov_88.598%
Anaeromassilibacillus senegalensis_nov_92.460%
Lachnospiraceae_[G-14] bacterium_MOT-185_nov_93.103%
Sporobacter termitidis_nov_87.897%
Butyrivibrio pullicaecorum_nov_85.934%
Lawsonibacter asaccharolyticus_nov_90.421%
Duncaniella freteri_nov_90.262%
Lawsonibacter asaccharolyticus_nov_91.762%
Duncaniella freteri_nov_93.208%
Lachnospiraceae_[G-11] bacterium_MOT-178_nov_94.778%
Hathewayia proteolytica_nov_84.569%
Lachnospiraceae_[G-14] bacterium_MOT-185_nov_97.154%
Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.267%
Acetivibrio cellulolyticus_nov_85.921%
Lachnospiraceae_[G-11] bacterium_MOT-177_nov_93.922%
Lachnospiraceae_[G-11] bacterium_MOT-177_nov_92.549%
Phocaea massiliensis_nov_90.060%
Lachnospiraceae_[G-14] bacterium_MOT-185_nov_94.320%
Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.713%
Anaeroplasmabactoclasticum_nov_87.352%

Species

Samples