



Bacteroides thetaiotaomicron
Lachnospiraceae_[G-1] bacterium_MOT-166
Akkermansia muciniphila
Lachnospiraceae_[G-14] bacterium_MOT-184
Parasutterella excrementihominis
Lachnospiraceae_[G-11] bacterium_MOT-177
Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.069%
Blautia faecicola_nov_89.749%
Duncaniella freteri_nov_94.162%
Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.478%
Acetivibrio cellulolyticus_nov_83.851%
Lachnospiraceae_[G-10] bacterium_MOT-175_nov_91.085%
Sporobacter termitidis_nov_83.168%
Clostridiales_[F-1][G-1] bacterium_HMT_093_nov_84.091%
Lawsonibacter asaccharolyticus_nov_94.798%
Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.450%
Oscillospiraceae_[G-4] bacterium_MOT-151_nov_92.899%
Lachnospiraceae_[G-11] bacterium_MOT-176_nov_94.990%
Acetivibrio cellulolyticus_nov_83.090%
Duncaniella freteri_nov_90.323%
Ruminiclostridium cellulolyticum_nov_82.704%
Hydrogenoanaerobacterium saccharovorans_nov_89.942%
Kineothrix alysoides_nov_87.129%
Lachnospiraceae_[G-12] bacterium_MOT-179_nov_94.778%
Lachnoclostridium [Clostridium] polysaccharolyticum_nov_87.452%
Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.098%
Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.805%
Lachnospiraceae_[G-11] bacterium_MOT-178_nov_92.293%
Duncaniella freteri_nov_93.774%
Lachnospiraceae_[G-13] bacterium_MOT-181_nov_91.602%
Enterocloster bolteae_nov_94.951%
Lachnospiraceae_[G-9] bacterium_MOT-174_nov_90.430%
Eubacteriales_[G-3] bacterium_MOT-163_nov_89.695%
Lachnospiraceae_[G-11] bacterium_MOT-178_nov_93.064%
Lachnospiraceae_[G-14] bacterium_MOT-185_nov_93.103%
Lachnospiraceae_[G-11] bacterium_MOT-178_nov_92.293%
Lawsonibacter asaccharolyticus_nov_91.762%
Lachnoclostridium [Clostridium] polysaccharolyticum_nov_93.243%
Lachnospiraceae_[G-11] bacterium_MOT-178_nov_94.778%
Hathewayia proteolytica_nov_84.569%
Muribaculaceae_[G-1] bacterium_MOT-129_nov_88.697%
Lachnospiraceae_[G-14] bacterium_MOT-185_nov_97.154%
Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.267%
Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.886%
Lachnospiraceae_[G-11] bacterium_MOT-177_nov_93.922%
Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.590%
Lachnospiraceae_[G-11] bacterium_MOT-177_nov_92.549%
Duncaniella freteri_nov_93.208%
Phoceia massiliensis_nov_90.060%
Anaeroplasma abactoclasticum_nov_87.352%

Species

F5362.S06

F5362.S05

F5362.S12

F5362.S11

F5362.S10

Samples