



- Eubacteriales_[G-3] bacterium_MOT-163
- Oscillospiraceae_[G-4] bacterium_MOT-151
- Lachnospiraceae_[G-14] bacterium_MOT-184
- Mollicutes_[G-1] bacterium_MOT-186
- Lachnospiraceae_[G-1] bacterium_MOT-166
- Oscillospiraceae_[G-6] bacterium_MOT-153
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_85.887%
- Lachnospiraceae_[G-11] bacterium_MOT-177_nov_96.066%
- Acetivibrio cellulolyticus_nov_83.153%
- Anaerotignum faecicola_nov_85.287%
- Lachnoclostridium phytofermentans_nov_91.458%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.466%
- Agrococcus versicolor_nov_83.227%
- Anaerotignum lactatifermentans_nov_84.040%
- Clostridiales_[F-1][G-1] bacterium_HMT_093_nov_91.775%
- Acetivibrio cellulolyticus_nov_82.189%
- Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.489%
- Turicibacter sanguinis_nov_95.923%
- Lachnospiraceae_[G-14] bacterium_MOT-185_nov_93.348%
- Acetivibrio cellulolyticus_nov_85.560%
- Lachnospiraceae_[G-14] bacterium_MOT-182_nov_90.042%
- Phoceea massiliensis_nov_89.722%
- Eubacterium ventriosum_nov_93.320%
- Acetivibrio cellulolyticus_nov_83.190%
- Acetivibrio cellulolyticus_nov_83.761%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.407%
- Lachnoclostridium [Clostridium] herbivorans_nov_93.125%
- Lachnospiraceae_[G-14] bacterium_MOT-185_nov_97.210%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.125%
- Mobilitalea sibirica_nov_90.583%
- Hathewayia proteolytica_nov_83.297%
- Marvinbryantia formatexigens_nov_91.435%
- Pseudoflavonifractor capillosus_nov_89.897%
- Lachnospiraceae_[G-14] bacterium_MOT-184_nov_92.584%
- Acetivibrio cellulolyticus_nov_83.801%
- Lachnospiraceae_[G-12] bacterium_MOT-180_nov_89.613%
- Butyrivibrio fibrisolvens_nov_86.831%
- Blautia faecicola_nov_91.170%
- Oscillospiraceae_[G-4] bacterium_MOT-151_nov_95.634%
- Christensenella hongkongensis_nov_85.294%
- Oscillospiraceae_[G-3] bacterium_MOT-150_nov_91.736%
- Hathewayia proteolytica_nov_84.233%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.041%
- Eubacteriales_[G-1] bacterium_MOT-159_nov_94.055%
- Lachnoclostridium [Clostridium] scindens_nov_89.648%
- Fusicatenibacter saccharivorans_nov_90.041%
- Lachnospiraceae_[G-6] bacterium_MOT-171_nov_93.096%
- Lachnospiraceae_[G-14] bacterium_MOT-183_nov_93.348%
- Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.719%
- Christensenella multispecies_sppn3_2_nov_86.275%

Species

F30682.S02

F30682.S01

F30682.S08

F30682.S07

F30682.S06

Samples