



- Lachnospiraceae\_[G-14] bacterium\_MOT-184
- Lachnospiraceae\_[G-1] bacterium\_MOT-166
- Clostridium disporicum
- Oscillospiraceae\_[G-6] bacterium\_MOT-153
- Lactobacillus johnsonii
- 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%
- Phoceia 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%
- Mobilitealea 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%
- Lachnospiraceae\_[G-12] bacterium\_MOT-179\_nov\_94.501%
- Eubacteriales\_[G-3] bacterium\_MOT-163\_nov\_85.944%
- Eubacteriales\_[G-4] bacterium\_MOT-164\_nov\_94.292%
- Lachnospiraceae\_[G-14] bacterium\_MOT-183\_nov\_93.348%
- Lachnospiraceae\_[G-11] bacterium\_MOT-178\_nov\_91.886%
- Lachnospiraceae\_[G-6] bacterium\_MOT-171\_nov\_95.388%
- Lachnospiraceae\_[G-6] bacterium\_MOT-171\_nov\_94.351%
- Lachnospiraceae\_[G-14] bacterium\_MOT-185\_nov\_92.719%
- Christensenella multispecies\_sppn3\_2\_nov\_86.275%

Species

F30682.S05

F30682.S04

F30682.S03

F30682.S11

F30682.S10

F30682.S09

Samples