



- Oscillospiraceae\_[G-7] bacterium\_MOT-154
- Eubacteriales\_[G-1] bacterium\_MOT-159\_nov\_94.118%
- Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_92.625%
- multigenus multispecies\_sppn292\_3\_nov\_87.368%
- Eubacteriales\_[G-4] bacterium\_MOT-165
- Devosia insulae
- Duncaniella freteri\_nov\_92.903%
- Lachnospiraceae\_[G-10] bacterium\_MOT-175\_nov\_95.034%
- Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_92.704%
- Eubacteriales\_[G-4] bacterium\_MOT-165\_nov\_94.357%
- Lachnospiraceae\_[G-9] bacterium\_MOT-174\_nov\_95.701%
- Eubacteriales\_[G-4] bacterium\_MOT-164\_nov\_95.485%
- Eubacterium ventriosum\_nov\_96.388%
- Lachnospiraceae\_[G-12] bacterium\_MOT-179\_nov\_92.534%
- Erysipelatoclostridium [Clostridium] innocuum
- Longibaculum muris\_nov\_92.094%
- Duncaniella freteri\_nov\_88.248%
- Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_93.737%
- Eubacterium ventriosum\_nov\_96.372%
- Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_89.177%
- Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_90.693%
- Desulfovibrio fairfieldensis\_nov\_94.421%
- Oscillospiraceae\_[G-6] bacterium\_MOT-153\_nov\_94.157%
- Faecalicatena multispecies\_sppn189\_2\_nov\_97.279%
- Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_91.810%
- Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_94.577%
- Paraeggerthella hongkongensis\_nov\_92.793%
- Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_91.991%
- Bacteroides acidifaciens\_acidofaciens
- Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_89.224%
- Duncaniella freteri\_nov\_90.672%
- Bifidobacterium pseudolongum
- Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_90.929%
- Alistipes putredinis\_nov\_96.753%
- Eubacteriales\_[G-2] bacterium\_MOT-162\_nov\_95.260%
- Lachnospiraceae\_[G-12] bacterium\_MOT-179
- Kineothrix alysoides\_nov\_96.840%
- Parabacteroides merdae\_nov\_95.032%
- Anaerotaenia torta\_nov\_97.059%
- Holdemania massiliensis\_nov\_92.060%
- Butyricoccus pullicaecorum\_nov\_94.407%
- Lachnoclostridium [Clostridium] scindens
- Faecalibaculum rodentium
- Kineothrix alysoides\_nov\_97.517%
- Paraburkholderia hospita\_nov\_91.083%
- Corynebacterium mastitidis
- Ruminococcus bromii
- Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_93.074%
- Kineothrix alysoides\_nov\_93.275%
- Carnobacteriaceae\_[G-1] bacterium\_MOT-198
- Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_92.441%
- Muribaculum intestinale\_nov\_89.744%
- Adlercreutzia mucosicola
- Povalibacter uvarum\_nov\_91.702%
- Aminipila butyrica\_nov\_91.910%
- Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_92.857%
- Lactobacillus intestinalis
- Lactobacillus\_Limosilactobacillus reuteri\_reuteri\_clade\_938
- Helicobacter ganmani
- Eubacteriales\_[G-2] bacterium\_MOT-162
- Blautia henseni\_hominis\_marasmi
- Anaerostipes caccae
- Maihella massiliensis\_nov\_92.111%
- Desulfovibrio fairfieldensis\_nov\_94.647%
- Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_91.087%
- Ramlibacter monticola
- Afipia\_Bradyrhizobium archetypum\_australiense\_broomeae\_elkani
- Arthrobacter\_Paeniglutamibacter\_Pseudarthrobacter cryotolerans
- Turicimonas muris
- Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_91.323%
- Lactobacillus taiwanensis
- Rodentibacter pneumotropicus
- Ileibacterium valens
- Oscillospiraceae\_[G-3] bacterium\_MOT-150
- Muribaculaceae\_[G-1] bacterium\_MOT-129
- Alistipes sp.\_MOT-127
- Bacteroides acidifaciens
- Muribaculum intestinale
- Lachnospiraceae\_[G-11] bacterium\_MOT-178
- Lactobacillus johnsonii
- Eubacteriales\_[G-1] bacterium\_MOT-160
- Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_93.521%
- Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_91.304%
- Eubacteriales\_[G-4] bacterium\_MOT-164
- Muribaculum intestinale\_nov\_87.957%
- Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_91.757%
- Streptococcus acidominimus\_sp.\_MOT-012
- Pasteurella\_Rodentibacter caecimuris\_heylii
- Eubacteriales\_[G-1] bacterium\_MOT-159
- Clostridium disporicum\_saudiense
- Helicobacter typhlonius
- Escherichia\_Shigella coli\_fergusonii\_flexneri\_sonnei
- Parabacteroides goldsteinii
- Prevotella sp.\_MOT-128
- Ligilactobacillus animalis\_apodemi\_murinus
- Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_91.087%
- Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_91.540%
- Erysipelatoclostridium [Clostridium] cocleatum
- Erysipelotrichaceae\_[G-1] bacterium\_MOT-189
- Akkermansia muciniphila

Species

- F0000.S002
- F0000.S003
- F0000.S001
- F0000.S006
- F0000.S018
- F0000.S004
- F0000.S005
- F0000.S017
- F0000.S011
- F0000.S014
- F0000.S013
- F0000.S012
- F0000.S008
- F0000.S010
- F0000.S009
- F0000.S007
- F0000.S015
- F0000.S016

Samples