



Comparison\_2

- Control 1
- Amoxicillin 1



- Lachnospiraceae\_[G-11] bacterium\_MOT-178
- Alistipes sp.\_MOT-127
- Bacteroides acidofaciens
- Bacteroides caecimuris
- Blautia caecimuris
- Prevotella sp.\_MOT-128
- Lactobacillus taiwanensis
- Oscillospiraceae\_[G-2] bacterium\_MOT-149
- Lachnospiraceae\_[G-14] bacterium\_MOT-183
- Lachnospiraceae\_[G-9] bacterium\_MOT-174
- Alistipes timonensis
- Bacteroides stercorisoris
- Muribaculum intestinale
- Lachnospiraceae\_[G-14] bacterium\_MOT-184
- Parabacteroides goldsteinii
- Akkermansia muciniphila
- Kineothrix alysoides\_nov\_95.227%
- Anaerotruncus rubifaciens\_nov\_92.760%
- Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_91.991%
- Mediterraneibacter [Ruminococcus] gnavus\_nov\_93.424%
- Alistipes putredinis\_nov\_95.887%
- Oscillibacter valericigenes\_nov\_95.260%
- Kineothrix alysoides\_nov\_97.279%
- Pseudobutyrvibrio ruminis\_nov\_91.176%
- Saccharofermentans acetigenes\_nov\_88.764%
- Lacrimispora indolis\_nov\_90.724%
- Kineothrix alysoides\_nov\_93.651%
- Lachnospiraceae\_[G-14] bacterium\_MOT-185\_nov\_92.358%
- Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_90.870%
- Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_91.106%
- Odoribacter splanchnicus\_nov\_93.939%
- Oscillospiraceae\_[G-2] bacterium\_MOT-149\_nov\_95.506%
- Ruthenibacterium lactatiformans\_nov\_97.045%
- Lachnospiraceae\_[G-11] bacterium\_MOT-178\_nov\_97.978%
- Lachnospiraceae\_[G-14] bacterium\_MOT-185\_nov\_96.599%
- Prevotellamassilia timonensis\_nov\_94.168%
- Neglectibacter timonensis\_nov\_97.500%
- Maihella massiliensis\_nov\_92.094%
- Neglectibacter timonensis\_nov\_97.727%
- Butyricoccus pullicaecorum\_nov\_94.820%
- Lachnospiraceae\_[G-11] bacterium\_MOT-176\_nov\_95.946%
- Lachnospiraceae\_[G-14] bacterium\_MOT-184\_nov\_95.405%
- Acetatifactor muris\_nov\_96.145%
- Oscillospiraceae\_[G-2] bacterium\_MOT-149\_nov\_95.056%
- Lachnospiraceae\_[G-14] bacterium\_MOT-183\_nov\_95.425%
- Oscillospiraceae\_[G-2] bacterium\_MOT-149\_nov\_95.281%
- Petrocella atlantisensis\_nov\_87.810%
- Oscillospiraceae\_[G-3] bacterium\_MOT-150\_nov\_96.396%
- Lachnospiraceae\_[G-6] bacterium\_MOT-171\_nov\_95.238%
- Eubacteriales\_[G-2] bacterium\_MOT-162\_nov\_95.260%
- Anaerosporeobacter mobilis\_nov\_95.000%
- Alistipes putredinis\_nov\_96.529%
- Phoceia massiliensis\_nov\_95.682%
- Muribaculum intestinale\_nov\_93.737%
- Clostridiales\_[F-1][G-1] bacterium\_HMT\_093\_nov\_90.337%
- Lachnospiraceae\_[G-14] bacterium\_MOT-184\_nov\_95.692%
- Eubacteriales\_[G-3] bacterium\_MOT-163\_nov\_93.679%
- Lachnoclostridium [Clostridium] populeti\_nov\_94.331%
- Lachnospiraceae\_[G-11] bacterium\_MOT-176\_nov\_97.297%
- Acetatifactor muris\_nov\_92.551%
- Lachnoclostridium [Clostridium] populeti\_nov\_92.986%
- Blautia caecimuris\_nov\_96.825%
- Duncaniella freteri\_nov\_90.456%
- Kineothrix alysoides\_nov\_93.682%
- Coproccoccus catus\_nov\_94.570%
- Roseburia faecis\_nov\_97.964%
- Duncaniella freteri\_nov\_93.103%
- Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_93.074%
- Lacrimispora xylanolytica\_nov\_97.285%
- Culturomica massiliensis\_nov\_93.709%
- Kineothrix alysoides\_nov\_95.928%
- Lachnospiraceae\_[G-10] bacterium\_MOT-175\_nov\_96.372%
- Lachnospiraceae\_[G-10] bacterium\_MOT-175\_nov\_92.174%
- Lachnoclostridium pacaense\_nov\_96.825%
- Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_89.462%
- Eubacterium coprostanoligenes\_nov\_95.485%
- Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_91.974%
- Lachnospiraceae\_[G-12] bacterium\_MOT-179\_nov\_92.534%
- Lachnospiraceae\_[G-14] bacterium\_MOT-184\_nov\_94.989%
- Pseudoflavonifractor capillosus\_nov\_95.721%
- Anaerotignum lactatifermentans\_nov\_95.270%
- Caecibacterium sporofaciens\_nov\_95.045%
- Oscillospiraceae\_[G-2] bacterium\_MOT-149\_nov\_95.946%
- Oscillospiraceae\_[G-2] bacterium\_MOT-149\_nov\_94.157%
- Alistipes senegalensis\_nov\_95.228%
- Oscillospiraceae\_[G-4] bacterium\_MOT-151\_nov\_91.723%
- Lachnospiraceae\_[G-11] bacterium\_MOT-177\_nov\_97.523%
- Bacteroides acidifaciens\_acidofaciens
- Lachnospiraceae\_[G-12] bacterium\_MOT-179\_bacterium\_MOT-18
- multigenus multispecies\_sppn10\_2\_nov\_95.918%
- Alistipes multispecies\_sppn12\_2\_nov\_96.304%
- multigenus multispecies\_sppn13\_5\_nov\_94.570%
- multigenus multispecies\_sppn18\_2\_nov\_92.063%
- Bacteroidetes\_[G-3] multispecies\_sppn2\_2\_nov\_87.554%
- Roseburia multispecies\_sppn21\_3\_nov\_95.711%
- Lachnospiraceae\_[G-11] multispecies\_sppn4\_2\_nov\_96.847%
- Bacteroides multispecies\_sppn6\_2\_nov\_96.312%
- multigenus multispecies\_sppn7\_2\_nov\_92.777%
- multigenus multispecies\_sppn8\_3\_nov\_95.011%
- multigenus multispecies\_sppn9\_2\_nov\_93.002%

Species

F8810.S10

F8810.S11

F8810.S12

F8810.S01

F8810.S02

F8810.S03

Samples