



Comparison_2
Hamlet + Amoxicillin 1
Amoxicillin 1



- Lachnospiraceae_[G-11] bacterium_MOT-178
- Alistipes sp._MOT-127
- Turicimonas muris
- Bacteroides acidofaciens
- Bacteroides caecimuris
- Blautia caecimuris
- Prevotella sp._MOT-128
- Acutalibacter muris
- Roseburia faecis
- Flavonifractor plautii
- Oscillospiraceae_[G-2] bacterium_MOT-149
- Lachnospiraceae_[G-14] bacterium_MOT-183
- Robinsoniella peoriensis
- Lachnospiraceae_[G-9] bacterium_MOT-174
- Bacteroides stercorisoris
- Muribaculum intestinale
- Lachnospiraceae_[G-14] bacterium_MOT-184
- Parabacteroides goldsteinii
- Akkermansia muciniphila
- Parabacteroides goldsteinii_nov_97.614%
- Anaerotruncus rubiinfantis_nov_92.760%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_91.991%
- Mediterraneibacter [Ruminococcus] gnavus_nov_93.424%
- Oscillibacter valericigenes_nov_95.260%
- Saccharofermentans acetigenes_nov_88.764%
- Lacrimispora indolis_nov_90.724%
- Kineothrix alysoides_nov_93.651%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_90.870%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_91.106%
- 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-9] bacterium_MOT-174_nov_96.388%
- Lachnospiraceae_[G-14] bacterium_MOT-185_nov_96.599%
- Enterocloster asparagiformis_nov_94.344%
- Prevotellamassilia timonensis_nov_94.168%
- Eisenbergiella massiliensis_nov_96.599%
- Maihella massiliensis_nov_92.094%
- Anaerotruncus rubiinfantis_nov_92.517%
- Eubacterium coprostanoligenes_nov_95.701%
- Lachnospiraceae_[G-14] bacterium_MOT-184_nov_95.405%
- 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%
- Lachnospiraceae_[G-6] bacterium_MOT-171_nov_95.238%
- Anaerosporeobacter mobilis_nov_95.000%
- Alistipes putredinis_nov_96.529%
- Lawsonibacter asaccharolyticus_nov_97.297%
- Lawsonibacter asaccharolyticus_nov_95.730%
- Anaerotignum lactatifermentans_nov_97.523%
- Phocaea massiliensis_nov_95.682%
- Muribaculum intestinale_nov_93.737%
- Clostridiales_[F-1][G-1] bacterium_HMT_093_nov_90.337%
- Butyrivibrio pullicaecorum_nov_94.382%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.396%
- Lachnoclostridium [Clostridium] populeti_nov_94.331%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.270%
- Lachnospiraceae_[G-14] bacterium_MOT-184_nov_95.238%
- Neglectibacter timonensis_nov_97.959%
- Lachnospiraceae_[G-11] bacterium_MOT-176_nov_97.297%
- Acetatifactor muris_nov_92.551%
- Alistipes sp._MOT-127_nov_91.775%
- Blautia caecimuris_nov_96.825%
- Duncaniella freteri_nov_90.456%
- Coprococcus catus_nov_94.570%
- 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%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_92.191%
- 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%
- Alistipes timonensis_nov_97.831%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.946%
- Oscillospiraceae_[G-4] bacterium_MOT-151_nov_91.723%
- Enterobacter asburiae_cancerogenus_cloacae_hormaechei
- Bacteroides acidifaciens_acidofaciens
- multigenus multispecies_sppn10_2_nov_95.918%
- Alistipes multispecies_sppn12_2_nov_96.304%
- Eubacteriales_[G-1] multispecies_sppn15_2_nov_97.511%
- multigenus multispecies_sppn17_2_nov_95.928%
- multigenus multispecies_sppn20_3_nov_95.455%
- Roseburia multispecies_sppn21_3_nov_95.711%
- multigenus multispecies_sppn22_2_nov_95.465%
- Lachnospiraceae_[G-11] multispecies_sppn4_2_nov_96.847%
- multigenus multispecies_sppn5_2_nov_97.279%
- 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.S07
F8810.S08
F8810.S09

Samples