



Comparison_2
Control 1
Hamlet 1



- Lachnospiraceae_[G-11] bacterium_MOT-178
Alistipes sp._MOT-127
Bacteroides caecimuris
Peptococcaceae_[G-1] bacterium_MOT-146
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
Eubacteriales_[G-2] bacterium_MOT-162
Bacteroides stercorisoris
Lachnospiraceae_[G-14] bacterium_MOT-184
Anaerotaenia torta_nov_97.273%
Kineothrix alysoides_nov_95.227%
Anaerotruncus rubiinfantis_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%
Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.358%
Muribaculaceae_[G-2] bacterium_MOT-104_nov_91.106%
Alistipes putredinis_nov_95.879%
Odoribacter splanchnicus_nov_93.939%
Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.506%
Lachnospiraceae_[G-10] bacterium_MOT-175_nov_95.475%
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%
Butyricicoccus pullicaecorum_nov_94.820%
Lachnospiraceae_[G-11] bacterium_MOT-176_nov_95.946%
Eubacterium coprostanoligenes_nov_95.701%
Lachnospiraceae_[G-14] bacterium_MOT-184_nov_95.405%
Acetatifactor muris_nov_96.145%
Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.056%
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%
Ruminococcus albus_nov_92.500%
Alistipes putredinis_nov_96.529%
Lawsonibacter asaccharolyticus_nov_97.297%
Clostridium oryzae_nov_88.889%
Eisenbergiella massiliensis_nov_95.260%
Phocaea massiliensis_nov_95.682%
Muribaculum intestinale_nov_93.737%
Lachnospiraceae_[G-14] bacterium_MOT-184_nov_95.692%
Eubacteriales_[G-3] bacterium_MOT-163_nov_93.679%
Lachnospiraceae_[G-12] bacterium_MOT-180_nov_93.665%
Lachnoclostridium [Clostridium] populeti_nov_94.331%
Lachnospiraceae_[G-11] bacterium_MOT-176_nov_97.297%
Acetatifactor muris_nov_92.551%
Kineothrix alysoides_nov_97.059%
Lachnoclostridium [Clostridium] populeti_nov_92.986%
Duncaniella freteri_nov_90.456%
Kineothrix alysoides_nov_93.682%
Coproccoccus catus_nov_94.570%
Roseburia faecis_nov_97.964%
Eubacterium ventriosum_nov_96.825%
Muribaculaceae_[G-2] bacterium_MOT-104_nov_93.074%
Lacrimispora xylanolytica_nov_97.285%
Culturomica massiliensis_nov_93.709%
Kineothrix alysoides_nov_95.928%
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-14] bacterium_MOT-184_nov_94.989%
Pseudoflavonifractor capillosus_nov_95.721%
Anaerotignum lactatifermentans_nov_95.270%
Caecibacterium sporiformans_nov_95.045%
Alistipes timonensis_nov_97.831%
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-180
multigenus multispecies_sppn10_2_nov_95.918%
multigenus multispecies_sppn11_2_nov_95.465%
multigenus multispecies_sppn13_5_nov_94.570%
multigenus multispecies_sppn16_2_nov_96.833%
multigenus multispecies_sppn18_2_nov_92.063%
Bacteroidetes_[G-3] multispecies_sppn2_2_nov_87.554%
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.S01
F8810.S02
F8810.S03
F8810.S04
F8810.S05
F8810.S06

Samples