



Comparison_2
 Control 2
 Hamlet 2



- Lactobacillus taiwanensis
- Lawsonibacter asaccharolyticus_nov_97.973%
- Alistipes timonensis_nov_97.831%
- Oscillospiraceae_[G-2] bacterium_MOT-149
- Lachnospiraceae_[G-14] bacterium_MOT-184_nov_94.989%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_94.157%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_92.208%
- multigenus multispecies_sppn11_2_nov_95.465%
- Mageeibacillus indolicus_nov_87.668%
- multigenus multispecies_sppn17_2_nov_95.928%
- Alistipes putredinis_nov_95.887%
- Lachnospiraceae_[G-6] bacterium_MOT-171_nov_95.485%
- Lachnospiraceae_[G-9] bacterium_MOT-174_nov_96.388%
- Anaerospobacter mobilis_nov_95.000%
- Lachnospiraceae_[G-11] bacterium_MOT-176_nov_97.297%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.919%
- Muribaculum intestinale_nov_93.737%
- Neglectibacter timonensis_nov_97.727%
- Alistipes putredinis_nov_95.879%
- Peptococcaceae_[G-1] bacterium_MOT-146
- Ruthenibacterium lactatiformans_nov_97.045%
- Lachnospiraceae_[G-9] bacterium_MOT-174_nov_96.136%
- Alistipes senegalensis_nov_95.228%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_91.974%
- Oscillospiraceae_[G-4] bacterium_MOT-151_nov_91.723%
- Bacteroides multispecies_sppn6_2_nov_96.312%
- Butyricoccus pullicaecorum_nov_94.820%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_91.106%
- Christensenella hongkongensis_nov_86.353%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.946%
- Algmonas porphyrae_nov_83.596%
- Lachnospiraceae_[G-11] bacterium_MOT-178_nov_97.978%
- Oscillibacter valericiogenes_nov_95.260%
- Acetatifactor muris_nov_96.145%
- Lachnoclostridium [Clostridium] populeti_nov_94.331%
- Odoribacter splanchnicus_nov_93.939%
- multigenus multispecies_sppn8_3_nov_95.011%
- Bacteroides caecimuris
- Alistipes putredinis_nov_96.529%
- Duncaniella freteri_nov_90.456%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_93.074%
- Alistipes sp._MOT-127
- Lachnospiraceae_[G-14] bacterium_MOT-183
- Lachnospiraceae_[G-6] bacterium_MOT-171_nov_95.238%
- Lachnospiraceae_[G-12] bacterium_MOT-179_bacterium_MOT-183
- Pseudobutyrvibrio ruminis_nov_91.176%
- Alistipes multispecies_sppn12_2_nov_96.304%
- Lachnospiraceae_[G-10] bacterium_MOT-175_nov_95.475%
- Duncaniella freteri_nov_88.462%
- Anaerotruncus colihominis_nov_94.091%
- Clostridium oryzae_nov_88.889%
- Clostridium disporicum_saudiense
- Saccharofermentans acetigenes_nov_88.764%
- multigenus multispecies_sppn19_3_nov_96.818%
- Oscillospiraceae_[G-4] bacterium_MOT-151_nov_96.847%
- Bacteroides stercorisoris
- multigenus multispecies_sppn10_2_nov_95.918%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_90.870%
- multigenus multispecies_sppn23_2_nov_96.818%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_91.991%
- multigenus multispecies_sppn5_2_nov_97.279%
- Anaerotruncus rubiinfantis_nov_92.760%
- Culturomica massiliensis_nov_93.709%
- Lawsonibacter asaccharolyticus
- Mediterraneibacter [Ruminococcus] gnavus_nov_93.424%
- Kineothrix alysoides_nov_95.928%
- multigenus multispecies_sppn7_2_nov_92.777%
- Falcatimonas natans_nov_93.651%
- Caecibacterium sporiformans_nov_95.045%
- Lachnospiraceae_[G-11] bacterium_MOT-178
- Kineothrix alysoides_nov_97.279%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.462%
- Lachnospiraceae_[G-11] multispecies_sppn4_2_nov_96.847%
- Prevotellamassilia timonensis_nov_94.168%
- Anaerotignum lactatifermentans_nov_95.270%
- Eubacterium coprostanoligenes_nov_95.485%
- Lachnospiraceae_[G-12] bacterium_MOT-179_nov_92.534%
- Anaerotaenia torta_nov_97.273%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.506%
- Mailhella massiliensis_nov_92.094%
- Bacteroidetes_[G-3] multispecies_sppn2_2_nov_87.554%
- Lachnospiraceae_[G-9] bacterium_MOT-174_nov_96.364%
- Lachnoclostridium pacaense_nov_96.825%
- Acutalibacter muris
- Bacteroides acidifaciens_acidofaciens
- Prevotella sp._MOT-128
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_92.191%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.056%
- Lachnospiraceae_[G-9] bacterium_MOT-174
- Phocaea massiliensis_nov_95.682%
- Pseudoflavonifractor capillosus_nov_95.721%
- Lachnospiraceae_[G-14] bacterium_MOT-184
- Lacrimispora indolis_nov_90.724%
- Kineothrix alysoides_nov_93.651%
- Roseburia faecis_nov_97.964%
- Kineothrix alysoides_nov_93.682%
- Kineothrix alysoides_nov_97.059%
- multigenus multispecies_sppn9_2_nov_93.002%
- Lachnospiraceae_[G-12] bacterium_MOT-179_nov_94.796%
- Kineothrix alysoides_nov_95.227%

Species

F8810.S17
 F8810.S16
 F8810.S18
 F8810.S13
 F8810.S14
 F8810.S15

Samples