0
20
40
60
80
100
Value
Comparison_1
Control
Hamlet
Neglectibacter timonensis_nov_97.727%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_93.919%
Muribaculum intestinale_nov_93.737%
Lawsonibacter asaccharolyticus_nov_97.973%
Alistipes putredinis_nov_95.887%
Anaerotruncus rubiinfantis_nov_92.760%
Odoribacter splanchnicus_nov_93.939%
Lachnospiraceae_[G−11] bacterium_MOT−178_nov_97.978%
Oscillospiraceae_[G−4] bacterium_MOT−151_nov_91.723%
Clostridium oryzae_nov_88.889%
Eubacteriales_[G−2] bacterium_MOT−162
multigenus multispecies_sppn16_2_nov_96.833%
Eisenbergiella massiliensis_nov_95.260%
Lachnospiraceae_[G−10] bacterium_MOT−175_nov_95.475%
Saccharofermentans acetigenes_nov_88.764%
Lachnospiraceae_[G−11] bacterium_MOT−176_nov_97.297%
Ruthenibacterium lactatiformans_nov_97.045%
Oscillospiraceae_[G−4] bacterium_MOT−151_nov_96.847%
Peptococcaceae_[G−1] bacterium_MOT−146
Alistipes putredinis_nov_95.879%
Alistipes sp._MOT−127
Bacteroides acidifaciens_acidofaciens
Lachnospiraceae_[G−9] bacterium_MOT−174_nov_96.364%
Acutalibacter muris
Alistipes multispecies_sppn12_2_nov_96.304%
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%
Kineothrix alysoides_nov_97.279%
Lachnospiraceae_[G−11] multispecies_sppn4_2_nov_96.847%
Lachnospiraceae_[G−6] bacterium_MOT−171_nov_95.238%
Lachnospiraceae_[G−14] bacterium_MOT−183
Mediterraneibacter [Ruminococcus] gnavus_nov_93.424%
Lacrimispora xylanolytica_nov_97.285%
Lachnoclostridium [Clostridium] populeti_nov_92.986%
Lachnospiraceae_[G−11] bacterium_MOT−176_nov_95.946%
multigenus multispecies_sppn13_5_nov_94.570%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.946%
Oscillibacter valericigenes_nov_95.260%
Acetatifactor muris_nov_92.551%
Eubacteriales_[G−3] bacterium_MOT−163_nov_93.679%
Bacteroides multispecies_sppn6_2_nov_96.312%
Butyricicoccus pullicaecorum_nov_94.820%
Pseudobutyrivibrio ruminis_nov_91.176%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.974%
Eubacterium coprostanoligenes_nov_95.485%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_93.074%
Duncaniella freteri_nov_90.456%
Lachnoclostridium pacaense_nov_96.825%
Prevotellamassilia timonensis_nov_94.168%
Anaerotignum lactatifermentans_nov_95.270%
Lachnospiraceae_[G−12] bacterium_MOT−179_nov_92.534%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_92.358%
Lachnospiraceae_[G−10] bacterium_MOT−175_nov_92.174%
Lachnoclostridium [Clostridium] populeti_nov_94.331%
Acetatifactor muris_nov_96.145%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_96.599%
Lachnospiraceae_[G−11] bacterium_MOT−178
Alistipes putredinis_nov_96.529%
Bacteroides caecimuris
Lactobacillus taiwanensis
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_94.157%
Lachnospiraceae_[G−14] bacterium_MOT−184_nov_94.989%
Oscillospiraceae_[G−2] bacterium_MOT−149
Prevotella sp._MOT−128
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.191%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.056%
Bacteroides stercorirosoris
Alistipes timonensis_nov_97.831%
multigenus multispecies_sppn10_2_nov_95.918%
Lachnospiraceae_[G−9] bacterium_MOT−174
Lacrimispora indolis_nov_90.724%
Culturomica massiliensis_nov_93.709%
Lachnospiraceae_[G−11] bacterium_MOT−177_nov_97.523%
Alistipes senegalensis_nov_95.228%
Lachnospiraceae_[G−12] bacterium_MOT−179_bacterium_MOT−18
multigenus multispecies_sppn8_3_nov_95.011%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.106%
Ruminococcus albus_nov_92.500%
Anaerotaenia torta_nov_97.273%
multigenus multispecies_sppn7_2_nov_92.777%
Falcatimonas natans_nov_93.651%
Lawsonibacter asaccharolyticus
Kineothrix alysoides_nov_95.928%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.462%
Kineothrix alysoides_nov_93.651%
Caecibacterium sporoformans_nov_95.045%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.506%
Mailhella massiliensis_nov_92.094%
Phocea massiliensis_nov_95.682%
Bacteroidetes_[G−3] multispecies_sppn2_2_nov_87.554%
Pseudoflavonifractor capillosus_nov_95.721%
Lachnospiraceae_[G−14] bacterium_MOT−184
Kineothrix alysoides_nov_93.682%
Roseburia faecis_nov_97.964%
multigenus multispecies_sppn9_2_nov_93.002%
Kineothrix alysoides_nov_97.059%
Lachnospiraceae_[G−12] bacterium_MOT−179_nov_94.796%
Kineothrix alysoides_nov_95.227%
Species
F8810.S01
F8810.S02
F8810.S03
F8810.S13
F8810.S14
F8810.S15
F8810.S04
F8810.S05
F8810.S06
F8810.S16
F8810.S17
F8810.S18
Samples