0
20
40
60
80
100
Value
Group
WT5Dfec
WT5Dlig
Lactobacillus_Limosilactobacillus reuteri_reuteri_clade_938
Lactobacillus intestinalis
Peptococcaceae_[G−1] bacterium_MOT−146
Clostridiales_[F−1][G−1] bacterium_HMT_093_nov_90.337%
Acutalibacter muris
Duncaniella freteri
Anaeroplasma abactoclasticum_nov_92.394%
Parabacteroides distasonis_nov_97.397%
Bacteroidetes_[G−3] multispecies_sppn13_2_nov_86.422%
Prevotellamassilia timonensis_nov_93.521%
Lachnospiraceae_[G−2] bacterium_MOT−167
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.208%
Culturomica massiliensis_nov_91.126%
Millionella massiliensis_nov_92.408%
Lachnospiraceae_[G−1] bacterium_MOT−166
Oscillospiraceae_[G−6] bacterium_MOT−153
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.842%
Bacteroidetes_[G−3] multispecies_sppn670_2_nov_87.768%
Prevotella sp._MOT−128_nov_91.540%
Lachnospiraceae_[G−12] bacterium_MOT−179
Clostridium disporicum_saudiense
Lachnospiraceae_[G−11] bacterium_MOT−177
Muribaculaceae_[G−1] bacterium_MOT−129_nov_88.069%
multigenus multispecies_sppn559_2_nov_92.457%
Duncaniella freteri_nov_90.476%
Lachnoclostridium [Clostridium] populeti_nov_96.145%
Lachnospiraceae_[G−14] bacterium_MOT−182_nov_97.732%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_93.018%
multigenus multispecies_sppn546_2_nov_92.657%
Lachnospiraceae_[G−9] bacterium_MOT−174
Oscillospiraceae_[G−2] bacterium_MOT−149
Parabacteroides merdae_nov_95.032%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.441%
Oscillospiraceae_[G−7] bacterium_MOT−154
Muribaculaceae_[G−1] bacterium_MOT−129
Muribaculaceae_[G−2] bacterium_MOT−104_nov_90.929%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_93.722%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.067%
Muribaculum intestinale_nov_92.258%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.625%
Lawsonibacter asaccharolyticus_nov_96.396%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.757%
Lachnospiraceae_[G−3] bacterium_MOT−168
Parabacteroides goldsteinii
Corynebacterium mastitidis
Ileibacterium valens
Faecalibaculum rodentium
Erysipelatoclostridium [Clostridium] cocleatum
Terrisporobacter petrolearius
Clostridium subterminale_sulfidigenes_thiosulfatireducens
Escherichia_Shigella coli_fergusonii_flexneri_sonnei
Enterococcus faecalis
Mailhella massiliensis_nov_92.111%
Alistipes sp._MOT−127
Eubacteriales_[G−2] bacterium_MOT−162
Bacteroides acidifaciens
Mediterraneibacter [Ruminococcus] gnavus_nov_93.213%
Anaerotaenia torta_nov_96.833%
Acetatifactor muris_nov_95.023%
Muribaculum intestinale_nov_92.208%
Lactobacillus johnsonii
Lawsonibacter asaccharolyticus_nov_95.281%
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_95.937%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.087%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.991%
Lachnospiraceae_[G−14] bacterium_MOT−185
Acetatifactor muris_nov_92.777%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_94.589%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.540%
Lachnospiraceae_[G−14] bacterium_MOT−184_nov_95.023%
Phocaeicola sartorii
Bifidobacterium pseudolongum
Streptococcus danieliae
Carnobacteriaceae_[G−1] bacterium_MOT−198
Alistipes timonensis
Enterocloster aldenensis
Caecibacterium sporoformans_nov_94.831%
Muribaculum intestinale
Prevotella sp._MOT−128
Alistipes senegalensis_nov_95.680%
Lachnospiraceae_[G−11] bacterium_MOT−178
Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.677%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_93.074%
Lactobacillus taiwanensis
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.323%
Helicobacter typhlonius
Lachnospiraceae_[G−11] bacterium_MOT−178_nov_96.847%
Lachnospiraceae_[G−14] bacterium_MOT−184
Helicobacter ganmani
Erysipelatoclostridium ramosum
Eubacterium coprostanoligenes_nov_95.937%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_93.521%
Ligilactobacillus animalis_apodemi_murinus
Akkermansia muciniphila
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.087%
Erysipelotrichaceae_[G−1] bacterium_MOT−189
Streptococcus acidominimus_sp._MOT−012
Kineothrix alysoides_nov_93.275%
Rodentibacter pneumotropicus
Pasteurella_Rodentibacter caecimuris_heylii
Species
F0000.S043
F0000.S037
F0000.S044
F0000.S038
F0000.S045
F0000.S042
F0000.S040
F0000.S041
F0000.S039
F0000.S075
F0000.S079
F0000.S078
F0000.S071
F0000.S073
F0000.S077
F0000.S072
F0000.S074
F0000.S076
Samples