0
20
40
60
80
100
Value
Group
KO10Dfec
KO10Dlig
Kineothrix alysoides_nov_95.937%
Alistipes multispecies_sppn426_2_nov_96.320%
Neglectibacter timonensis_nov_97.511%
Lawsonibacter asaccharolyticus_nov_96.847%
Lachnospiraceae_[G−6] bacterium_MOT−171_nov_94.570%
Aminipila butyrica_nov_91.910%
Muribaculum intestinale_nov_93.548%
Kineothrix alysoides_nov_97.517%
Alistipes putredinis_nov_95.269%
Eubacteriales_[G−2] bacterium_MOT−162_nov_95.034%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.888%
Saccharibacteria_(TM7)_[G−3] bacterium_HMT_351_nov_96.380%
Faecalibaculum rodentium
Oscillospiraceae_[G−3] bacterium_MOT−150
Alistipes putredinis_nov_96.753%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.677%
Helicobacter ganmani
Oscillospiraceae_[G−7] bacterium_MOT−154
Lactobacillus_Limosilactobacillus reuteri_reuteri_clade_938
Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.991%
Anaerotaenia torta_nov_96.833%
Roseburia faecis_nov_97.743%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_87.284%
Mucispirillum schaedleri_nov_96.802%
Eubacteriales_[G−2] bacterium_MOT−162_nov_95.260%
Lactobacillus johnsonii
Prevotella sp._MOT−128
Ileibacterium valens
Muribaculum intestinale_nov_92.424%
Culturomica massiliensis_nov_90.929%
Lachnospiraceae_[G−9] bacterium_MOT−174_nov_95.485%
Ruminococcus bromii
Eubacteriales_[G−2] bacterium_MOT−162
Bacteroides acidifaciens
Turicimonas muris
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.857%
Muribaculum intestinale_nov_92.026%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.991%
Bacteroides acidifaciens_acidofaciens
Arthrobacter_Paeniglutamicibacter_Pseudarthrobacter cryotolerans_
Bifidobacterium pseudolongum
Muribaculaceae_[G−2] bacterium_MOT−104_nov_90.693%
Lactobacillus iners
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.087%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.304%
Lactobacillus rogosae
Bacteroides fragilis
Fusicatenibacter saccharivorans
Roseburia faecis_nov_97.072%
Flavonifractor plautii
Ruminococcus callidus
Carnobacteriaceae_[G−1] bacterium_MOT−198
Lachnospira pectinoschiza_nov_97.506%
Blautia faecis
Olsenella multispecies_sppn107_2_nov_95.982%
Mediterraneibacter [Ruminococcus] gnavus
Corynebacterium mastitidis
Blautia wexlerae
Bacteroides multispecies_sppn9_2_nov_96.095%
Bacteroides uniformis
Klebsiella pneumoniae
Streptococcus thoraltensis
Blautia hansenii_hominis_marasmi
Erysipelatoclostridium [Clostridium] innocuum
Parabacteroides distasonis_nov_96.753%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_93.521%
Streptococcus danieliae
Phocaeicola sartorii
Eubacteriales_[G−1] multispecies_sppn229_2_nov_97.517%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.323%
Lactobacillus taiwanensis
Muribaculaceae_[G−2] bacterium_MOT−104_nov_90.929%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.540%
Eubacteriales_[G−4] bacterium_MOT−164
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.991%
Muribaculum intestinale
Lachnospiraceae_[G−11] bacterium_MOT−178
Eubacteriales_[G−1] bacterium_MOT−160
Alistipes sp._MOT−127
Muribaculaceae_[G−1] bacterium_MOT−129
Faecalibacterium prausnitzii
Roseburia inulinivorans
Ruminococcus multispecies_sppn677_2_nov_94.808%
Enterobacter asburiae_cancerogenus_cloacae_hormaechei
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.540%
Escherichia_Shigella coli_fergusonii_flexneri_sonnei
Anaerostipes caccae
Parabacteroides goldsteinii
Helicobacter typhlonius
Eubacteriales_[G−1] bacterium_MOT−159
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.087%
Akkermansia muciniphila
Actinidia eriantha_nov_97.065%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.757%
Ligilactobacillus animalis_apodemi_murinus
Streptococcus acidominimus_sp._MOT−012
Rodentibacter pneumotropicus
Erysipelatoclostridium [Clostridium] cocleatum
Erysipelotrichaceae_[G−1] bacterium_MOT−189
Pasteurella_Rodentibacter caecimuris_heylii
Species
F0000.S097
F0000.S096
F0000.S024
F0000.S019
F0000.S020
F0000.S023
F0000.S021
F0000.S027
F0000.S022
F0000.S026
F0000.S028
F0000.S025
F0000.S088
F0000.S093
F0000.S089
F0000.S091
F0000.S090
F0000.S094
F0000.S092
F0000.S095
Samples