0
20
40
60
80
100
Value
Group
KO5Dfec
KO5Dlig
Oscillospiraceae_[G−2] bacterium_MOT−149
Oscillospiraceae_[G−3] bacterium_MOT−150
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.625%
Lawsonibacter asaccharolyticus_nov_97.303%
Oscillospiraceae_[G−6] bacterium_MOT−153
Muribaculum intestinale_nov_92.258%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_93.074%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_90.929%
Lactobacillus johnsonii
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.457%
Clostridium disporicum_saudiense
Lactobacillus intestinalis
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.067%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.441%
Lactobacillus_Limosilactobacillus reuteri_reuteri_clade_938
multigenus multispecies_sppn626_2_nov_95.701%
Peptococcaceae_[G−1] bacterium_MOT−146
Mailhella massiliensis_nov_92.111%
Erysipelatoclostridium ramosum
Triticum aestivum
Alistipes timonensis
Lachnospiraceae_[G−3] bacterium_MOT−168
Ileibacterium valens
Parabacteroides goldsteinii
Enterococcus faecalis
Bifidobacterium pseudolongum
Robinsoniella peoriensis
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.641%
Anaerotignum lactatifermentans_nov_97.297%
Eisenbergiella massiliensis_nov_95.928%
Bacteroidetes_[G−3] multispecies_sppn45_2_nov_87.686%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_87.636%
Muribaculum intestinale_nov_93.534%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.810%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.304%
Prevotella sp._MOT−128_nov_92.191%
Adlercreutzia mucosicola
Staphylococcus saprophyticus_xylosus
Corynebacterium mastitidis
Desulfovibrio fairfieldensis
Alistipes putredinis_nov_96.104%
Gemella sp._MOT−043
Ruthenibacterium lactatiformans
Muribaculaceae_[G−2] bacterium_MOT−104_nov_91.991%
Prevotella sp._MOT−128_nov_91.595%
Prevotellamassilia timonensis_nov_93.534%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.704%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.540%
Turicimonas muris
Oscillospiraceae_[G−7] bacterium_MOT−154
Muribaculaceae_[G−2] bacterium_MOT−104_nov_94.181%
Faecalicatena orotica_nov_97.291%
Prevotella sp._MOT−128_nov_91.416%
Prevotella sp._MOT−128
Bacteroides acidifaciens
Escherichia_Shigella coli_fergusonii_flexneri_sonnei
Kineothrix alysoides_nov_93.275%
Enterocloster aldenensis
Eubacteriales_[G−2] bacterium_MOT−162
Lachnospiraceae_[G−11] bacterium_MOT−178
Lawsonibacter asaccharolyticus_nov_94.843%
Phocea massiliensis_nov_92.099%
Duncaniella freteri_nov_92.903%
Muribaculum intestinale_nov_89.744%
Lachnospiraceae_[G−12] bacterium_MOT−179_bacterium_MOT−18
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.540%
Lachnoclostridium pacaense_nov_96.163%
Eubacteriales_[G−4] bacterium_MOT−164
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.991%
Helicobacter typhlonius
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.473%
Lachnospiraceae_[G−10] bacterium_MOT−175_nov_92.191%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.323%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.087%
Bacteroides multispecies_sppn9_2_nov_96.095%
Carnobacteriaceae_[G−1] bacterium_MOT−198
multigenus multispecies_sppn87_2_nov_92.225%
Prevotella sp._MOT−128_nov_92.009%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_93.521%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_92.625%
Helicobacter ganmani
Alistipes sp._MOT−127
Erysipelatoclostridium [Clostridium] cocleatum
Muribaculaceae_[G−2] bacterium_MOT−104_nov_94.577%
Lachnospiraceae_[G−14] bacterium_MOT−185
Muribaculum intestinale
Alistipes senegalensis_nov_95.680%
Lachnospiraceae_[G−14] bacterium_MOT−184
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.757%
Phocaeicola sartorii
Streptococcus danieliae
Prevotellamassilia timonensis_nov_94.194%
Muribaculaceae_[G−1] bacterium_MOT−129
Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.087%
Ligilactobacillus animalis_apodemi_murinus
Erysipelotrichaceae_[G−1] bacterium_MOT−189
Akkermansia muciniphila
Pasteurella_Rodentibacter caecimuris_heylii
Streptococcus acidominimus_sp._MOT−012
Rodentibacter pneumotropicus
Species
F0000.S053
F0000.S052
F0000.S051
F0000.S050
F0000.S047
F0000.S048
F0000.S046
F0000.S049
F0000.S080
F0000.S087
F0000.S083
F0000.S082
F0000.S086
F0000.S084
F0000.S081
F0000.S085
Samples