0
20
40
60
80
100
Value
Group
WT−F17722
WT−F18971
Actinidia eriantha
Rothia nasimurium
Weissella cibaria
Erysipelotrichaceae_[G−1] bacterium_MOT−189_nov_88.798%
Latilactobacillus curvatus_graminis
Lactococcus lactis
Corynebacterium mastitidis
Ralstonia sp._HMT_406
Enterococcus faecalis_nov_95.825%
Streptococcus thermophilus
Stenotrophomonas maltophilia
Fusobacterium varium_nov_96.696%
Cutibacterium acnes
Leptothrix sp._HMT_025
Pseudomonas helleri
Stenotrophomonas [Pseudomonas] hibiscicola
Moraxella osloensis
Anaerococcus sp._HMT_290
Massilia aurea
Psychrobacter alimentarius
Cutibacterium granulosum
Streptomyces aculeolatus
Secundilactobacillus paracollinoides
Delftia acidovorans
Pelomonas saccharophila
Actinidia eriantha_nov_97.011%
Fusobacterium perfoetens_nov_91.126%
Lacticaseibacillus rhamnosus
Enterococcus gallinarum
Enterococcus faecalis
Lachnospiraceae_[G−14] bacterium_MOT−185_nov_92.719%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_85.887%
Limosilactobacillus reuteri
Atopostipes sp._MOT−201
Sphingomonas aquatilis_melonis
Levilactobacillus brevis
Streptococcus thoraltensis
Gemella sp._MOT−033
Ligilactobacillus murinus
Lactobacillus johnsonii
Staphylococcus saprophyticus
Mollicutes_[G−1] bacterium_MOT−186
Eubacteriales_[G−2] bacterium_MOT−162
Oribacterium parvum_nov_89.770%
Fusicatenibacter saccharivorans_nov_90.526%
Oscillospiraceae_[G−6] bacterium_MOT−153_nov_91.631%
Eubacteriales_[G−4] bacterium_MOT−164
Lachnospiraceae_[G−11] bacterium_MOT−177
Muribaculaceae_[G−2] bacterium_MOT−104_nov_88.423%
Duncaniella freteri_nov_89.718%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.000%
Clostridium disporicum
Alistipes senegalensis_nov_93.443%
Peptococcus sp._HMT_168_nov_84.866%
Lachnospiraceae_[G−14] bacterium_MOT−185
Weissella paramesenteroides
Streptococcus acidominimus
Pediococcus pentosaceus
Akkermansia muciniphila
Muribaculaceae_[G−1] bacterium_MOT−129_nov_88.105%
Duncaniella freteri_nov_88.934%
Duncaniella freteri_nov_89.775%
Turicibacter sanguinis_nov_95.923%
Duncaniella freteri_nov_93.293%
Bifidobacterium pseudolongum
Dubosiella newyorkensis
Triticum aestivum
Corynebacterium stationis
Corynebacterium ammoniagenes
Staphylococcus saprophyticus_xylosus
Duncaniella freteri_nov_93.699%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_88.867%
Porphyromonas sp._MOT−131
Carnobacteriaceae_[G−1] bacterium_MOT−197
Staphylococcus equorum
Jeotgalicoccus halotolerans
Staphylococcus ureilyticus
Mammaliicoccus lentus
Rodentibacter pneumotropicus
Muribacter sp._MOT−143
Streptococcus danieliae
Ligilactobacillus animalis
Species
F17722.S06
F17722.S08
F17722.S10
F17722.S07
F17722.S09
F18971.S01
F18971.S02
F18971.S04
F18971.S03
F18971.S05
Samples