0
20
40
60
80
100
Value
Group2
Ligature Old Female
Ligature Old Male
Ligature Young Female
Ligature Young Male
Brain Old Perio Female
Brain Old Perio Male
Brain Young Perio Female
Brain Young Perio Male
Stool Old Perio Female
Stool Old Perio Male
Stool Young Perio Female
Stool Young Perio Male
Muribaculaceae_[G−1] bacterium_MOT−104_nov_89.000%
Duncaniella freteri_nov_89.290%
Eubacterium xylanophilum_nov_89.940%
Duncaniella freteri_nov_89.699%
Desulfovibrio fairfieldensis_nov_96.349%
Erysipelatoclostridium_ramosum_innocuum_nov_88.270%
Acetivibrio cellulolyticus_nov_83.405%
Parasutterella excrementihominis_nov_97.972%
Ileibacterium valens
Duncaniella freteri_nov_86.290%
Longibaculum muris_nov_86.957%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_86.640%
Alistipes putredinis_nov_94.444%
Phocaeicola sartorii
Leptotrichia hofstadii_nov_96.970%
Streptococcus acidominimus
Oscillospiraceae_[G−6] bacterium_MOT−153_nov_91.631%
Lachnospiraceae_[G−14] bacterium_MOT−183_nov_97.854%
Hathewaya proteolytica_nov_83.297%
Duncaniella freteri_nov_86.948%
Lachnospiraceae_[G−2] bacterium_MOT−167_nov_93.096%
Acutalibacter muris
Muribaculaceae_[G−1] bacterium_MOT−129_nov_87.576%
Duncaniella freteri_nov_86.842%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_86.000%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_86.373%
Limosilactobacillus reuteri
Enterococcus casseliflavus
Microbacterium maritypicum
Streptococcus gordonii
Leucobacter chromiiresistens
Duncaniella freteri_nov_89.919%
Comamonas testosteroni
Prevotella shahii_nov_87.602%
Duncaniella freteri_nov_92.653%
Duncaniella freteri_nov_87.424%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_83.636%
Bacteroides acidifaciens
Erysipelotrichaceae_[G−1] bacterium_MOT−189
Gluceribacter canis_nov_93.305%
Parabacteroides goldsteinii
Muribaculaceae_[G−2] bacterium_MOT−104_nov_88.577%
Duncaniella freteri_nov_90.184%
Muribaculum intestinale
Alistipes senegalensis_nov_93.673%
Duncaniella freteri_nov_90.612%
Parasutterella excrementihominis
Duncaniella freteri_nov_93.712%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_88.129%
Turicibacter sanguinis_nov_95.923%
Alistipes sp._MOT−127
Alistipes senegalensis_nov_93.686%
Bacteroides thetaiotaomicron
Bifidobacterium pseudolongum
Mammaliicoccus lentus
Staphylococcus ureilyticus
Dialister invisus
Enterobacter asburiae
Deinococcus geothermalis
Aggregatibacter sp._HMT_512
Pyrinomonas methylaliphatogenes_nov_96.809%
Shigella sonnei
Shigella flexneri
Fusobacterium nucleatum
Muribaculaceae_[G−1] bacterium_MOT−129_nov_86.089%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_84.929%
Fusobacterium nucleatum_subsp._vincentii
Methylobacterium brachiatum
Cutibacterium acnes
Chryseobacterium yeoncheonense_nov_97.484%
Staphylococcus capitis_epidermidis
Klebsiella quasipneumoniae
Enterobacter kobei
Alistipes senegalensis_nov_93.648%
Duncaniella freteri_nov_91.429%
Akkermansia muciniphila
Rhodococcus qingshengii
Bacillus albus_cereus_luti_nitratireducens_paramycoides_tro ...(6 sp
Neisseriaceae_[G−1] bacterium_MOT−031
Pasteurella_Rodentibacter caecimuris_heylii
Streptococcus thoraltensis
Streptococcus azizii_nov_95.171%
Citrobacter koseri
Chryseobacterium gambrini
Acinetobacter johnsonii
Atlantibacter hermannii
Klebsiella pneumoniae
Kosakonia sacchari
Enterobacter mori_nov_97.951%
Enterobacter mori
Muribaculaceae_[G−1] bacterium_MOT−129_nov_89.431%
Muribaculaceae_[G−1] bacterium_MOT−129
Ligilactobacillus murinus
Lactobacillus johnsonii
Proteus mirabilis
Enterococcus gallinarum
Staphylococcus argenteus_aureus_roterodami
Enterococcus faecalis
Enterobacter hormaechei
Staphylococcus saprophyticus_xylosus
Species
F15127.S07
F15127.S08
F15127.S09
F15127.S10
F15127.S11
F15127.S12
F13506.S09
F13506.S10
F13506.S11
F13506.S12
F15127.S01
F15127.S02
F15127.S03
F15127.S04
F15127.S05
F15127.S06
F13506.S01
F13506.S02
F13506.S03
F13506.S04
F15127.S19
F15127.S20
F15127.S21
F15127.S22
F15127.S23
F15127.S24
F4401.S20
F4401.S21
F4401.S22
F4401.S23
F4401.S24
F4401.S25
Samples