0 20 40 60 80 100
Value
Group1
Brain Perio
Brain Non−perio
Fusobacterium nucleatum_subsp._vincentii
Acinetobacter radioresistens
Streptococcus oralis_subsp._tigurinus_clade_071
Lachnospiraceae_[G−14] bacterium_MOT−185
Citrobacter koseri
Campylobacter showae
Enterobacter cloacae
Cetobacterium somerae
Treponema denticola
Lactobacillus johnsonii
Enterobacter hormaechei
Streptococcus gordonii
Streptococcus oralis
Shigella flexneri
Fusobacterium periodonticum
Bifidobacterium longum
Agromyces mediolanus
Alloprevotella tannerae
Peptostreptococcaceae_[XI][G−4] bacterium_HMT_369
Bradyrhizobium pachyrhizi
Peptidiphaga sp._HMT_183
Leucobacter chromiiresistens
Acinetobacter lwoffii
Deinococcus geothermalis
Selenomonas flueggei
Catonella sp._HMT_164
Acinetobacter johnsonii
Methylobacterium brachiatum
Atlantibacter hermannii
Streptococcus oralis_subsp._dentisani_clade_058
Aggregatibacter aphrophilus
Streptococcus oralis_subsp._tigurinus_clade_070
Streptococcus mutans
Aggregatibacter sp._HMT_512
Johnsonella sp._HMT_166
Janibacter melonis
Prevotella melaninogenica
Dialister pneumosintes
Corynebacterium matruchotii
Schaalia sp._HMT_178
Thermicanus aegyptius
Staphylococcus warneri
Rothia dentocariosa
Actinomyces sp._HMT_448
Leptotrichia buccalis
Actinomyces sp._HMT_414
Alishewanella agri
Kosakonia sacchari
Porphyromonas endodontalis
Cutibacterium acnes
Klebsiella pneumoniae
Enterococcus faecalis
Proteus mirabilis
Mogibacterium timidum
Sphingobacterium multivorum
Bacteroidetes_[G−5] bacterium_HMT_511
Shigella sonnei
Fusobacterium nucleatum
Pseudomonas putida
Prevotella buccae
Staphylococcus capitis
Chryseobacterium gambrini
Enemella evansiae
Enterobacter asburiae
Dialister invisus
Microbacterium maritypicum
Staphylococcus ureilyticus
Campylobacter concisus
Gemella morbillorum
Rhodococcus qingshengii
Bifidobacterium pseudolongum
Phocaeicola sartorii
Lawsonella clevelandensis
Campylobacter sp._HMT_044
Acidovorax ebreus
Leptotrichia sp._HMT_219
Enterococcus casseliflavus
Streptococcus sanguinis
Duncaniella freteri_nov_90.184%
Leptotrichia hofstadii_nov_96.970%
Duncaniella freteri_nov_90.612%
Duncaniella freteri_nov_93.699%
Muribaculaceae_[G−1] bacterium_MOT−129_nov_90.816%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.198%
Phocea massiliensis_nov_90.129%
Duncaniella freteri_nov_93.712%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_93.319%
Duncaniella freteri_nov_89.919%
Actinomyces sp._HMT_169_nov_97.992%
Natroniella acetigena_nov_80.992%
Chryseobacterium yeoncheonense_nov_97.484%
Pyrinomonas methylaliphatogenes_nov_96.809%
Lachnoclostridium [Clostridium] polysaccharolyticum_nov_92.083%
Duncaniella freteri_nov_91.039%
Veillonella dispar_parvula
Bacillus albus_cereus_luti_nitratireducens_paramycoides_tro ...(6 sp
Staphylococcus argenteus_aureus_roterodami
Streptococcus infantis_infantis_clade_638
Fusobacterium nucleatum_nucleatum_subsp._animalis
Staphylococcus capitis_epidermidis
Species
F15127.S13
F15127.S14
F15127.S15
F15127.S16
F15127.S17
F15127.S18
F15127.S07
F15127.S08
F15127.S09
F15127.S10
F15127.S11
F15127.S12
Samples