0
20
40
60
80
100
Value
Group
Brain WT + Ligature
Brain PSAPP + Ligature
Brain WT − Ligature
Brain PSAPP − Ligature
Enterococcus faecalis
Staphylococcus hominis
Klebsiella aerogenes
Veillonella sp._HMT_780
Parasutterella excrementihominis
Corynebacterium coyleae
Glutamicibacter soli
Tepidiphilus succinatimandens
Cutibacterium avidum
Corynebacterium matruchotii
Neisseria mucosa
Enterobacter hormaechei
Bradyrhizobium valentinum
Ligilactobacillus murinus
Janibacter cremeus
Staphylococcus caprae
Klebsiella pneumoniae
Streptococcus danieliae
Varibaculum cambriense
Staphylococcus capitis
Corynebacterium tuberculostearicum
Fusobacterium nucleatum
Facklamia languida
Rothia mucilaginosa
Actinomyces sp._HMT_448
Pantoea allii
Streptococcus parasanguinis_clade_411
Streptococcus sp._HMT_066
Methylophilus leisingeri
Neisseriaceae_[G−1] bacterium_HMT_174
Cutibacterium acnes
Weizmannia ginsengihumi
Bradyrhizobium lupini
Enterococcus gallinarum
Acinetobacter lwoffii
Enterobacter cloacae
Streptococcus mitis
Corynebacterium durum
Bifidobacterium pseudolongum
Cutibacterium namnetense
Microbacterium maritypicum
Corynebacterium ureicelerivorans
Corynebacterium aurimucosum
Jeotgalicoccus halotolerans
Corynebacterium jeikeium
Triticum aestivum
Bifidobacterium longum
Faecalibaculum rodentium
Bacillus subtilis
Corynebacterium afermentans
Salmonella enterica
Peribacillus frigoritolerans
Rhodococcus qingshengii
Streptococcus lactarius
Acinetobacter pittii
Actinidia eriantha
Staphylococcus warneri
Bradyrhizobium japonicum
Staphylococcus pasteuri
Acinetobacter johnsonii
Tannerella serpentiformis
Mammaliicoccus sciuri
Shigella flexneri
Corynebacterium kroppenstedtii
Corynebacterium pilbarense
Mammaliicoccus lentus
Mediterraneibacter faecis
Escherichia fergusonii
Bifidobacterium bifidum
Streptococcus chosunense
Massilia arenae
Lawsonella clevelandensis
Sphingomonas leidyi
Streptococcus oralis
Streptococcus mutans
Acinetobacter septicus
Bacillus halotolerans
Abiotrophia defectiva
Pseudoramibacter alactolyticus
Corynebacterium pseudodiphtheriticum
Corynebacterium falsenii
Lactiplantibacillus plantarum
Prevotella sp._HMT_314
Kocuria indica_nov_94.872%
Corynebacterium argentoratense_nov_95.850%
Daejeonella oryzae_nov_85.551%
Thermodesulfobium acidiphilum_nov_81.729%
Arthrospira platensis_nov_88.987%
Sporosarcina sp._MOT−205_nov_96.360%
Brevibacterium paucivorans_nov_97.368%
Clostridium saccharoperbutylacetonicum_nov_95.800%
Microbacterium saccharophilum_nov_76.981%
Duncaniella freteri_nov_90.075%
Ralstonia solanacearum_nov_95.922%
Frankia discariae_nov_81.481%
Staphylococcus argenteus_aureus_roterodami
Streptococcus infantis_infantis_clade_638
Bacillus albus_cereus_luti_nitratireducens_paramycoides_tro ...(6 sp
Terrimonas multispecies_sppn3_2_nov_90.822%
Gloeobacter multispecies_sppn4_2_nov_84.254%
Species
F9929.S13
F9929.S14
F9929.S15
F9929.S16
F9929.S17
F9929.S18
F9929.S19
F9929.S20
F9929.S21
F9929.S22
F9929.S23
F9929.S24
F9929.S25
F9929.S26
F9929.S27
F9929.S28
F9929.S29
F9929.S30
F9929.S31
F9929.S32
F9929.S33
F9929.S34
F9929.S35
Samples