0
20
40
60
80
100
Value
Group
Rat dental plaque
Rat feces
Bacteroides intestinalis
Bacteroides thetaiotaomicron
Atopostipes sp._MOT−200
Haemophilus parainfluenzae
Clostridium disporicum
Microbacterium hydrocarbonoxydans
Corynebacterium amycolatum
Odoribacter laneus
Collinsella aerofaciens
Erysipelatoclostridium [Clostridium] saccharogumia
Phocaeicola vulgatus
Shigella flexneri
Escherichia coli
Ligilactobacillus murinus
Blautia glucerasea
Cutibacterium acnes
Granulicatella adiacens
Bacteroides xylanisolvens
Bifidobacterium pseudolongum
Longicatena caecimuris
Lactobacillus intestinalis
Shigella dysenteriae
Rothia aerolata
Eisenbergiella massiliensis
Triticum aestivum
Staphylococcus ureilyticus
Actinidia eriantha
Fusicatenibacter saccharivorans
Parasutterella excrementihominis
Streptococcus agalactiae
Veillonella parvula
Streptococcus sp._HMT_064
Streptococcus mutans
Mammaliicoccus sciuri
Alistipes finegoldii
Romboutsia ilealis
Parabacteroides goldsteinii
Akkermansia muciniphila
Adlercreutzia equolifaciens
Monoglobus pectinilyticus
Bacteroides uniformis
Lachnospiraceae_[G−3] bacterium_MOT−168_nov_94.154%
Niveispirillum fermenti_nov_85.242%
Rothia nasimurium_nov_97.228%
Streptococcus sp._MOT−045_nov_97.764%
Anaerotignum aminivorans_nov_92.812%
Kiloniella majae_nov_85.144%
Blautia luti_nov_96.008%
Frisingicoccus caecimuris_nov_96.451%
Blautia glucerasea_nov_97.899%
Eubacterium xylanophilum_nov_91.313%
Turicibacter sanguinis_nov_96.137%
Lachnospiraceae_[G−6] bacterium_MOT−171_nov_94.130%
Eubacterium coprostanoligenes_nov_91.379%
Coprococcus eutactus_nov_92.704%
Agathobaculum desmolans_nov_86.345%
Butyricicoccus pullicaecorum_nov_87.216%
Oscillospiraceae_[G−6] bacterium_MOT−153_nov_97.831%
Longibaculum muris_nov_92.754%
Gluceribacter canis_nov_95.388%
Blautia faecicola_nov_93.111%
Oscillospiraceae_[G−4] bacterium_MOT−151_nov_90.496%
Lachnoclostridium [Clostridium] herbivorans_nov_92.723%
Lachnospiraceae_[G−3] bacterium_MOT−168_nov_93.750%
Ruminococcus albus_nov_91.045%
Lawsonibacter asaccharolyticus_nov_90.909%
Acutalibacter muris_nov_95.661%
Barnesiella viscericola_nov_90.447%
Fusicatenibacter saccharivorans_nov_89.474%
Oscillospiraceae_[G−1] bacterium_MOT−147_nov_95.218%
Lachnospiraceae_[G−9] bacterium_MOT−174_nov_96.855%
Erysipelatoclostridium [Clostridium] saccharogumia_nov_96.042%
Anaerostipes hadrus_nov_92.754%
Fusicatenibacter saccharivorans_nov_97.479%
Faecalimonas umbilicata_nov_96.033%
Lachnospiraceae_[G−11] bacterium_MOT−178_nov_94.929%
Oscillospiraceae_[G−4] bacterium_MOT−151_nov_92.946%
Anaerotruncus rubiinfantis_nov_88.820%
Oscillospiraceae_[G−3] bacterium_MOT−150_nov_91.097%
Pseudoflavonifractor phocaeensis_nov_94.191%
Pseudoflavonifractor phocaeensis_nov_92.340%
Lachnospiraceae_[G−3] bacterium_MOT−168_nov_94.154%
Oscillospiraceae_[G−3] bacterium_MOT−150_nov_90.702%
Coprococcus eutactus_nov_95.085%
Lachnospiraceae_[G−4] bacterium_MOT−169_nov_97.065%
Arthrospira platensis_nov_90.531%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_86.228%
Blautia schinkii_nov_94.340%
Duncaniella freteri_nov_89.024%
Alistipes finegoldii_nov_94.835%
Blautia producta_nov_96.017%
Ruminococcus albus_nov_92.225%
Chryseobacterium culicis_nov_97.280%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_96.473%
Blautia producta_nov_95.178%
Spongiibacter marinus_nov_80.142%
Staphylococcus argenteus_aureus_roterodami
Streptococcus parasanguinis_parasanguinis_clade_721
Veillonella dispar_parvula
Kordiimonas multispecies_sppn3_2_nov_79.121%
Species
F19489.S07
F19489.S08
F19489.S09
F19489.S10
F19489.S11
F19489.S12
F19489.S13
F19489.S14
F19489.S15
F19489.S16
F19489.S17
F19489.S18
Samples