0
20
40
60
80
100
Value
Group
Rat saliva
Rat feces
Pseudoflavonifractor phocaeensis_nov_94.191%
Lachnospiraceae_[G−3] bacterium_MOT−168_nov_94.154%
Lachnospiraceae_[G−6] bacterium_MOT−171_nov_94.340%
Lachnospiraceae_[G−11] bacterium_MOT−178_nov_94.929%
Anaerotruncus rubiinfantis_nov_88.820%
Oscillospiraceae_[G−3] bacterium_MOT−150_nov_90.702%
Longibaculum muris_nov_92.754%
Lawsonibacter asaccharolyticus_nov_90.909%
Acutalibacter muris_nov_95.661%
Oscillospiraceae_[G−1] bacterium_MOT−147_nov_95.218%
Oscillospiraceae_[G−4] bacterium_MOT−151_nov_92.946%
Barnesiella viscericola_nov_90.447%
Oscillospiraceae_[G−6] bacterium_MOT−153_nov_97.831%
Oscillospiraceae_[G−4] bacterium_MOT−151_nov_90.496%
Lachnoclostridium [Clostridium] herbivorans_nov_92.723%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_96.473%
Anaerotignum aminivorans_nov_92.812%
Lachnospiraceae_[G−6] bacterium_MOT−171_nov_94.130%
Eubacterium xylanophilum_nov_91.313%
Lachnospiraceae_[G−3] bacterium_MOT−168_nov_93.750%
Fusicatenibacter saccharivorans_nov_89.474%
Lachnospiraceae_[G−9] bacterium_MOT−174_nov_96.855%
Adlercreutzia equolifaciens
Erysipelatoclostridium [Clostridium] saccharogumia_nov_96.042%
Frisingicoccus caecimuris_nov_96.451%
Kiloniella majae_nov_85.144%
Blautia producta_nov_95.178%
Blautia glucerasea_nov_97.899%
Monoglobus pectinilyticus
Faecalimonas umbilicata_nov_96.033%
Zhongshania marina_nov_79.765%
Eubacterium coprostanoligenes_nov_91.379%
Staphylococcus haemolyticus
Kosakonia sacchari
Shigella sonnei
Corynebacterium amycolatum
Escherichia coli
Shigella dysenteriae
Enterococcus faecalis
Spongiibacter marinus_nov_80.142%
Coprococcus eutactus_nov_92.704%
Triticum aestivum
Veillonella parvula
Mammaliicoccus sciuri
Staphylococcus ureilyticus
Enterobacter cancerogenus
Streptococcus agalactiae
Granulicatella adiacens
Streptococcus sp._HMT_064
Enterococcus avium
Bacteroides thetaiotaomicron
Alistipes finegoldii
Gluceribacter canis_nov_95.388%
Niveispirillum fermenti_nov_85.242%
Blautia schinkii_nov_94.340%
Coprococcus eutactus_nov_95.085%
Kordiimonas multispecies_sppn3_2_nov_79.121%
Agathobaculum desmolans_nov_86.345%
Ruminococcus albus_nov_91.045%
Eisenbergiella massiliensis
Blautia producta_nov_96.017%
Bacteroides xylanisolvens
Anaerostipes hadrus_nov_92.754%
Odoribacter laneus
Bacteroides intestinalis
Bacteroides uniformis
Butyricicoccus pullicaecorum_nov_87.216%
Turicibacter sanguinis_nov_96.137%
Clostridium disporicum
Erysipelatoclostridium [Clostridium] saccharogumia
Pseudoflavonifractor phocaeensis_nov_92.340%
Ruminococcus albus_nov_92.225%
Lachnospiraceae_[G−3] bacterium_MOT−168_nov_94.154%
Oscillospiraceae_[G−3] bacterium_MOT−150_nov_91.097%
Parabacteroides goldsteinii
Alistipes finegoldii_nov_94.835%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_86.228%
Duncaniella freteri_nov_89.024%
Parasutterella excrementihominis
Veillonella dispar_parvula
Streptococcus parasanguinis_parasanguinis_clade_721
Staphylococcus argenteus_aureus_roterodami
Ligilactobacillus murinus
Akkermansia muciniphila
Lachnospiraceae_[G−4] bacterium_MOT−169_nov_97.065%
Longicatena caecimuris
Phocaeicola vulgatus
Fusicatenibacter saccharivorans_nov_97.479%
Fusicatenibacter saccharivorans
Romboutsia ilealis
Blautia luti_nov_96.008%
Haemophilus parainfluenzae
Blautia glucerasea
Blautia faecicola_nov_93.111%
Lactobacillus intestinalis
Bifidobacterium pseudolongum
Collinsella aerofaciens
Streptococcus mutans
Streptococcus sp._MOT−045_nov_97.764%
Rothia nasimurium_nov_97.228%
Species
F19489.S01
F19489.S02
F19489.S03
F19489.S04
F19489.S05
F19489.S06
F19489.S13
F19489.S14
F19489.S15
F19489.S16
F19489.S17
F19489.S18
Samples