0
20
40
60
80
100
Value
Group
Feces Group 8
Feces Group 9
Bifidobacterium pseudolongum
Eisenbergiella massiliensis
Oscillospiraceae_[G−6] bacterium_MOT−153
Lactobacillus intestinalis
Clostridium disporicum
Blautia luti
Phocaeicola vulgatus
Bacteroides uniformis
Parasutterella excrementihominis
Longicatena caecimuris
Collinsella aerofaciens
Akkermansia muciniphila
Bacteroides intestinalis
Odoribacter laneus
Parabacteroides goldsteinii
Erysipelatoclostridium [Clostridium] saccharogumia
Bacteroides xylanisolvens
Romboutsia ilealis
Eubacteriales_[G−1] bacterium_MOT−158
Bacteroides thetaiotaomicron
Blautia glucerasea
Fusicatenibacter saccharivorans
Adlercreutzia equolifaciens
Ligilactobacillus murinus
Alistipes finegoldii
Eubacterium coprostanoligenes_nov_91.379%
Blautia schinkii_nov_94.340%
Lachnospiraceae_[G−14] bacterium_MOT−183_nov_92.060%
Faecalicatena fissicatena_nov_93.946%
Kiloniella majae_nov_85.144%
Lachnospiraceae_[G−11] bacterium_MOT−177_nov_95.851%
Mediterraneibacter [Ruminococcus] torques_nov_93.515%
Blautia faecicola_nov_93.319%
Ruminococcus champanellensis_nov_90.713%
Roseburia faecis_nov_97.917%
Dysosmobacter welbionis_nov_86.902%
Sporobacter termitidis_nov_88.223%
Duncaniella freteri_nov_89.024%
Gluceribacter canis_nov_94.561%
Blautia producta_nov_95.178%
Spongiibacter marinus_nov_80.142%
Lactobacillus rogosae_nov_94.990%
Kineothrix alysoides_nov_91.441%
Coprococcus eutactus_nov_95.085%
Lachnoclostridium [Clostridium] herbivorans_nov_92.723%
Lachnoclostridium [Clostridium] herbivorans_nov_92.917%
Erysipelatoclostridium [Clostridium] saccharogumia_nov_96.042%
Agathobaculum desmolans_nov_86.345%
Ruminococcus flavefaciens_nov_94.760%
Anaerocolumna cellulosilytica_nov_88.655%
Oscillospiraceae_[G−4] bacterium_MOT−151_nov_90.496%
Turicibacter sanguinis_nov_96.137%
Oscillospiraceae_[G−4] bacterium_MOT−151_nov_93.347%
Eubacterium coprostanoligenes_nov_91.379%
Eubacterium ventriosum_nov_93.320%
Longibaculum muris_nov_89.506%
Pseudoflavonifractor phocaeensis_nov_91.511%
Cuneatibacter caecimuris_nov_92.308%
Eubacterium coprostanoligenes_nov_90.129%
Rothia nasimurium_nov_97.228%
Paludicola psychrotolerans_nov_88.115%
Gluceribacter canis_nov_95.388%
Lachnoclostridium [Clostridium] polysaccharolyticum_nov_90.437%
Eubacterium xylanophilum_nov_90.505%
Lawsonibacter asaccharolyticus_nov_90.909%
Blautia faecicola_nov_93.111%
Sporobacter termitidis_nov_87.124%
Raoultibacter timonensis_nov_90.476%
Butyricicoccus pullicaecorum_nov_87.216%
Gluceribacter canis_nov_93.946%
Eubacteriales_[G−1] bacterium_MOT−158_nov_91.507%
Lawsonibacter asaccharolyticus_nov_90.083%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_93.555%
Frisingicoccus caecimuris_nov_96.451%
Lachnospira multipara_nov_84.298%
Oscillospiraceae_[G−2] bacterium_MOT−149_nov_87.885%
Coprococcus eutactus_nov_92.704%
Barnesiella viscericola_nov_90.447%
Clostridium disporicum_nov_97.619%
Mediterraneibacter [Ruminococcus] gnavus_nov_95.188%
Blautia schinkii_nov_94.130%
Lachnospiraceae_[G−12] bacterium_MOT−180_nov_90.184%
Ruminococcus champanellensis_nov_91.810%
Ruminococcus albus_nov_91.045%
Streptococcus sp._MOT−045_nov_97.764%
Oscillospiraceae_[G−6] bacterium_MOT−153_nov_97.831%
Blautia luti_nov_96.429%
Muribaculaceae_[G−2] bacterium_MOT−104_nov_86.228%
Fusicatenibacter saccharivorans_nov_97.689%
Alistipes finegoldii_nov_94.835%
Lachnospiraceae_[G−4] bacterium_MOT−169_nov_97.484%
Blautia luti_nov_97.899%
Blautia producta_nov_96.226%
Lacrimispora saccharolytica_nov_90.625%
Oscillospiraceae_[G−1] bacterium_MOT−147_nov_90.760%
Oscillospiraceae_[G−1] bacterium_MOT−147_nov_93.595%
Eubacterium xylanophilum_nov_91.313%
Eisenbergiella massiliensis_nov_86.486%
Oscillospiraceae_[G−7] bacterium_MOT−154_nov_92.918%
Blautia multispecies_sppn28_2_nov_95.198%
Species
F18249.S33
F18249.S34
F18249.S35
F18249.S36
F18249.S37
F18249.S38
F18249.S39
F18249.S40
F18249.S41
F18249.S42
F18249.S43
F18249.S44
F18249.S45
F18249.S46
F18249.S47
F18249.S48
Samples