0
20
40
60
Value
Group
Baseline.B.Plaque 1
24 Weeks.B.Plaque 1
Streptococcus gordonii
Lactobacillus vaginalis
Fretibacterium fastidiosum
Streptococcus intermedius
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Fusobacterium nucleatum
Peptococcus sp._Oral_Taxon_167
Dialister invisus
Veillonella atypica
Selenomonas noxia
Streptococcus anginosus
Streptococcus mutans
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Megasphaera micronuciformis
Actinomyces sp._HMT_171
Actinomyces massiliensis
Actinomyces gerencseriae
Fusobacterium periodonticum
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Actinomyces sp._Oral_Taxon_171
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Fusobacterium nucleatum_ss_polymorphum
Anaeroglobus geminatus
Actinomyces naeslundii
Dialister pneumosintes
Veillonella parvula
Streptococcus pyogenes_nov_91.53%
Fusobacterium hwasookii_nov_87.47%
Streptococcus rifensis_nov_90.02%
Winogradskyella eckloniae_nov_83.62%
Vagococcus lutrae_nov_87.66%
Fusobacterium periodonticum_nov_86.34%
Arenibacter algicola_nov_84.04%
Fusobacterium sp._HMT_248_nov_86.74%
Bowdeniella nasicola_nov_86.78%
Tessaracoccus lapidicaptus_nov_87.1%
Streptococcus rifensis_nov_90.82%
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Campylobacter curvus_nov_91.68%
Coprococcus eutactus_nov_89.48%
Streptococcus rifensis_nov_90.65%
Streptococcus pyogenes_nov_89.52%
Corynebacterium terpenotabidum_nov_92.31%
Geosporobacter subterraneus_nov_88.04%
Streptococcus pyogenes_nov_89.52%
Clostridium diolis_nov_84.18%
Kocuria indica_nov_89.33%
Streptococcus rifensis_nov_91.68%
Natronincola ferrireducens_nov_85.53%
Streptococcus rifensis_nov_90.59%
Streptococcus rifensis_nov_90.67%
Thermomonospora curvata_nov_84.68%
Isoptericola jiangsuensis_nov_88.29%
Arenibacter algicola_nov_84.87%
Geosporobacter subterraneus_nov_84.49%
Tonsilliphilus suis_nov_84.02%
Streptococcus rifensis_nov_90.43%
Streptococcus pyogenes_nov_85.12%
Devriesea agamarum_nov_86.9%
Clostridium acetobutylicum_nov_85.27%
Oceanobacillus senegalensis_nov_85.28%
Anaerovibrio lipolyticus_nov_84.95%
Sneathia sanguinegens_nov_85.05%
Lacticaseibacillus porcinae_nov_86.26%
Fusobacterium nucleatum_nov_86.73%
Streptosporangium subfuscum_nov_84.24%
Bowdeniella nasicola_nov_87.61%
Leptotrichia sp._HMT_498_nov_84.5%
Anaerovibrio lipolyticus_nov_85.06%
Streptococcus suis_nov_88.37%
Marinoscillum pacificum_nov_80.04%
Kocuria indica_nov_89.81%
Peptostreptococcus anaerobius_nov_95.34%
Natronincola ferrireducens_nov_83.89%
Thermomonospora curvata_nov_84.47%
Actinomyces slackii_nov_90.12%
Thermincola potens_nov_80.71%
Alkaliphilus peptidifermentans_nov_82.53%
Isoptericola cucumis_nov_86.73%
Clostridium acetobutylicum_nov_85.65%
Cellulosimicrobium arenosum_nov_91.54%
Campylobacter rectus_nov_94.88%
Alkaliphilus peptidifermentans_nov_87.82%
Anaerovibrio lipolyticus_nov_85.17%
Chryseobacterium taeanense_nov_85.77%
Corynebacterium terpenotabidum_nov_92.36%
Flavobacterium subsaxonicum_nov_81.99%
Arenibacter algicola_nov_87.21%
Campylobacter rectus_nov_94.49%
Corynebacterium terpenotabidum_nov_92.32%
Bowdeniella nasicola_nov_87.76%
Bhargavaea cecembensis_nov_82.42%
Dermacoccus sp._str._Ellin185_nov_89.57%
Alistipes indistinctus_nov_86.39%
Anaerovibrio lipolyticus_nov_86.86%
Streptococcus pyogenes_nov_90.1%
Alkaliphilus peptidifermentans_nov_86.76%
Alistipes indistinctus_nov_85.71%
Cellulomonas gilvus_nov_87.87%
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S003
F5677.S006
F5677.S021
F5677.S029
F5677.S044
F5677.S048
F5677.S058
F5677.S065
F5677.S070
F5677.S074
F5677.S080
F5677.S091
F5677.S094
F5677.S101
F5677.S109
F5677.S116
F5677.S126
F5677.S131
F5677.S134
F5677.S144
F5677.S150
F5677.S154
F5677.S516
F5677.S527
F5677.S531
F5677.S539
F5677.S547
F5677.S550
F5677.S553
F5677.S558
F5677.S564
F5677.S570
F5677.S575
F5677.S579
F5677.S583
F5677.S586
F5677.S596
Samples