0
20
40
60
Value
Group
24 Weeks.A.Plaque 2
24 Weeks.B.Plaque 2
Streptococcus gordonii
Actinomyces sp._HMT_170
Fretibacterium fastidiosum
Selenomonas dianae
Streptococcus intermedius
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Fusobacterium nucleatum
Selenomonas infelix
Dialister invisus
Selenomonas noxia
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_957
Streptococcus anginosus
Streptococcus mutans
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
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
Lactobacillus gasseri
Anaeroglobus geminatus
Actinomyces naeslundii
Streptococcus sp._str._2136FAA
Streptococcus constellatus
Dialister pneumosintes
Veillonella parvula
Streptococcus pyogenes_nov_91.53%
Geosporobacter subterraneus_nov_88.43%
Anaerocolumna xylanovorans_nov_86.91%
Fusobacterium hwasookii_nov_87.47%
Thermotalea metallivorans_nov_88.34%
Fusobacterium hwasookii_nov_88.45%
Vagococcus lutrae_nov_87.66%
Arenibacter algicola_nov_84.04%
Bowdeniella nasicola_nov_86.78%
Thermotalea metallivorans_nov_87.32%
Xylanibacterium ulmi_nov_87.14%
Streptococcus rifensis_nov_90.82%
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Campylobacter curvus_nov_91.68%
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%
Isoptericola jiangsuensis_nov_88.29%
Arenibacter algicola_nov_84.87%
Cellulosilyticum lentocellum_nov_84.21%
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%
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%
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%
Actinomyces israelii_nov_96.89%
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%
Arenibacter algicola_nov_87.21%
Campylobacter rectus_nov_94.49%
Corynebacterium terpenotabidum_nov_93.1%
Corynebacterium terpenotabidum_nov_92.32%
Bowdeniella nasicola_nov_87.76%
Bhargavaea cecembensis_nov_82.42%
Alistipes indistinctus_nov_86.39%
Anaerovibrio lipolyticus_nov_86.86%
Alkaliphilus peptidifermentans_nov_86.76%
Fusobacterium sp._HMT_248_nov_86.16%
Selenomonas dianae_nov_87.22%
Geosporobacter subterraneus_nov_86.78%
Cellulomonas gilvus_nov_87.87%
Streptococcus gordonii_sp._str._2136FAA
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S598
F5677.S606
F5677.S609
F5677.S614
F5677.S619
F5677.S626
F5677.S629
F5677.S641
F5677.S647
F5677.S654
F5677.S661
F5677.S673
F5677.S676
F5677.S682
F5677.S601
F5677.S612
F5677.S616
F5677.S624
F5677.S632
F5677.S635
F5677.S638
F5677.S643
F5677.S649
F5677.S655
F5677.S660
F5677.S664
F5677.S668
F5677.S671
F5677.S681
Samples