



Group
 12 Weeks.A.Plaque 2
 24 Weeks.A.Plaque 2

- Anaerovibrio lipolyticus_nov_84.95%
- Marinoscillum pacificum_nov_80.04%
- Bhargavaea cecembensis_nov_82.42%
- Selenomonas noxia
- Selenomonas infelix
- Anaerocolumna xylanovorans_nov_86.91%
- Tessaracoccus lapidicaptus_nov_87.1%
- Arenibacter algicola_nov_87.21%
- Streptococcus gordonii_sp._str._2136FAA
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957
- Corynebacterium terpenotabidum_nov_92.36%
- Alistipes indistinctus_nov_85.71%
- Fusobacterium sp._HMT_248_nov_86.74%
- Lactobacillus vaginalis
- Dermacoccus sp._str._Elin185_nov_89.57%
- Streptococcus sp._str._2136FAA
- Isoptericola cucumis_nov_88.91%
- Tonsilliphilus suis_nov_84.02%
- Clostridium acetobutylicum_nov_85.27%
- Isoptericola cucumis_nov_86.73%
- Clostridium diolis_nov_84.18%
- Campylobacter rectus_nov_94.88%
- Oceanobacillus senegalensis_nov_85.28%
- Alkaliphilus peptidifermentans_nov_87.82%
- Fusobacterium sp._HMT_248_nov_86.16%
- Fusobacterium nucleatum_nov_86.73%
- Campylobacter rectus_nov_94.49%
- Natronincola ferrireducens_nov_83.89%
- Alistipes indistinctus_nov_86.39%
- Cellulosilyticum lentocellum_nov_84.21%
- Anaerovibrio lipolyticus_nov_85.06%
- Streptococcus constellatus
- Cellulosimicrobium arenosum_nov_91.54%
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349
- Geosporobacter subterraneus_nov_86.78%
- Candidatus_Amoebophilus asiaticus_nov_80.21%
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346
- Arenibacter algicola_nov_84.87%
- Megasphaera micronuciformis
- Bowdeniella nasicola_nov_87.61%
- Fusobacterium periodonticum
- Alkaliphilus peptidifermentans_nov_82.53%
- Thermotalea metallivorans_nov_87.32%
- Alkaliphilus peptidifermentans_nov_86.76%
- Dialister pneumosintes
- Actinomyces israelii_nov_96.89%
- Fretibacterium fastidiosum
- Anaeroglobus geminatus
- Peptostreptococcus anaerobius_nov_95.34%
- Thermincola potens_nov_80.71%
- Actinomyces sp._HMT_170
- Sneathia sanguinegens_nov_85.47%
- Streptosporangium subfuscum_nov_84.24%
- Streptococcus suis_nov_88.37%
- Actinomyces gerencseriae
- Streptococcus pyogenes_nov_85.12%
- Lactocaseibacillus porcinae_nov_86.26%
- Actinomyces massiliensis
- Actinomyces sp._HMT_171
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_347
- Actinomyces naeslundii
- Fusobacterium nucleatum_ss_polymorphum
- Fusobacterium nucleatum
- Campylobacter curvus_nov_91.68%
- Geosporobacter subterraneus_nov_88.04%
- Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
- Geosporobacter subterraneus_nov_84.49%
- Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356
- Streptococcus anginosus
- Streptococcus rifensis_nov_90.43%
- Lactobacillus gasseri
- Kocuria indica_nov_89.33%
- Streptococcus gordonii
- Vagococcus lutrae_nov_87.66%
- Streptococcus intermedius
- Nonomuraea lactucae_nov_85.43%
- Devriesea agamarum_nov_86.9%
- Streptococcus rifensis_nov_90.65%
- Corynebacterium terpenotabidum_nov_92.31%
- Fusobacterium hwasookii_nov_87.47%
- Streptococcus pyogenes_nov_89.52%
- Dialister invisus
- Natronincola ferrireducens_nov_85.53%
- Isoptericola jiangsuensis_nov_88.29%
- Streptococcus pyogenes_nov_91.53%
- Streptococcus rifensis_nov_90.59%
- Actinomyces sp._Oral_Taxon_171
- Clostridium acetobutylicum_nov_85.65%
- Sneathia sanguinegens_nov_85.05%
- Kocuria indica_nov_89.81%
- Corynebacterium terpenotabidum_nov_92.32%
- Streptococcus rifensis_nov_90.82%
- Streptococcus rifensis_nov_90.67%
- Streptococcus rifensis_nov_91.68%
- Cellulomonas gilvus_nov_87.87%
- Bowdeniella nasicola_nov_86.78%
- Veillonella parvula
- Streptococcus pyogenes_nov_89.52%
- Mycobacterium multispecies_sppn9_2_nov_83.57%
- Streptococcus mutans

Species

- F5677_S471
- F5677_S483
- F5677_S437
- F5677_S417
- F5677_S465
- F5677_S426
- F5677_S446
- F5677_S626
- F5677_S430
- F5677_S467
- F5677_S413
- F5677_S672
- F5677_S673
- F5677_S661
- F5677_S609
- F5677_S617
- F5677_S434
- F5677_S620
- F5677_S659
- F5677_S445
- F5677_S645
- F5677_S475
- F5677_S682
- F5677_S606
- F5677_S506
- F5677_S436
- F5677_S420
- F5677_S605
- F5677_S425

Samples