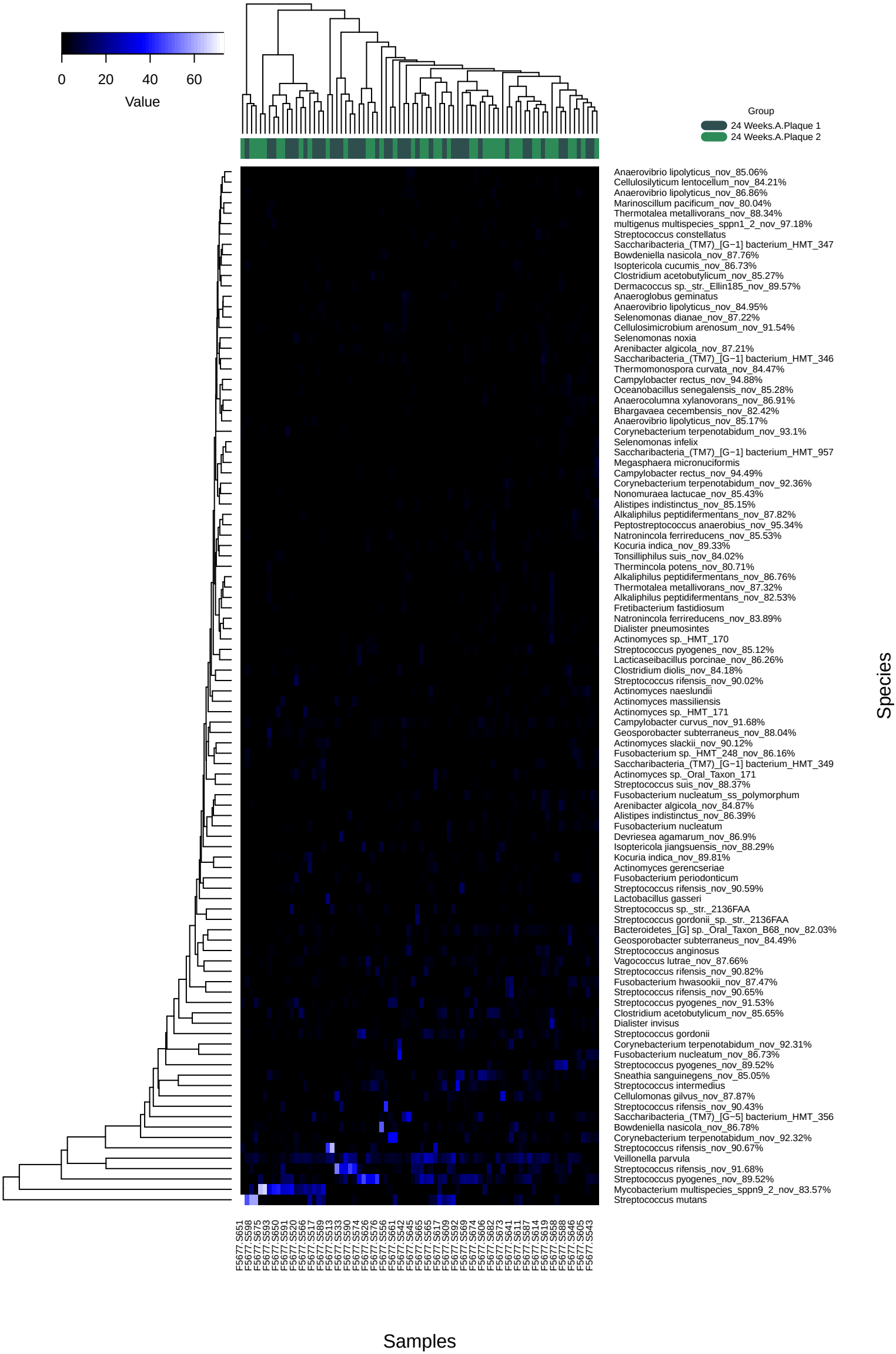
0
20
40
60
Value
Group
24 Weeks.A.Plaque 1
24 Weeks.A.Plaque 2
Anaerovibrio lipolyticus_nov_85.06%
Cellulosilyticum lentocellum_nov_84.21%
Anaerovibrio lipolyticus_nov_86.86%
Marinoscillum pacificum_nov_80.04%
Thermotalea metallivorans_nov_88.34%
multigenus multispecies_sppn1_2_nov_97.18%
Streptococcus constellatus
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Bowdeniella nasicola_nov_87.76%
Isoptericola cucumis_nov_86.73%
Clostridium acetobutylicum_nov_85.27%
Dermacoccus sp._str._Ellin185_nov_89.57%
Anaeroglobus geminatus
Anaerovibrio lipolyticus_nov_84.95%
Selenomonas dianae_nov_87.22%
Cellulosimicrobium arenosum_nov_91.54%
Selenomonas noxia
Arenibacter algicola_nov_87.21%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Thermomonospora curvata_nov_84.47%
Campylobacter rectus_nov_94.88%
Oceanobacillus senegalensis_nov_85.28%
Anaerocolumna xylanovorans_nov_86.91%
Bhargavaea cecembensis_nov_82.42%
Anaerovibrio lipolyticus_nov_85.17%
Corynebacterium terpenotabidum_nov_93.1%
Selenomonas infelix
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_957
Megasphaera micronuciformis
Campylobacter rectus_nov_94.49%
Corynebacterium terpenotabidum_nov_92.36%
Nonomuraea lactucae_nov_85.43%
Alistipes indistinctus_nov_85.15%
Alkaliphilus peptidifermentans_nov_87.82%
Peptostreptococcus anaerobius_nov_95.34%
Natronincola ferrireducens_nov_85.53%
Kocuria indica_nov_89.33%
Tonsilliphilus suis_nov_84.02%
Thermincola potens_nov_80.71%
Alkaliphilus peptidifermentans_nov_86.76%
Thermotalea metallivorans_nov_87.32%
Alkaliphilus peptidifermentans_nov_82.53%
Fretibacterium fastidiosum
Natronincola ferrireducens_nov_83.89%
Dialister pneumosintes
Actinomyces sp._HMT_170
Streptococcus pyogenes_nov_85.12%
Lacticaseibacillus porcinae_nov_86.26%
Clostridium diolis_nov_84.18%
Streptococcus rifensis_nov_90.02%
Actinomyces naeslundii
Actinomyces massiliensis
Actinomyces sp._HMT_171
Campylobacter curvus_nov_91.68%
Geosporobacter subterraneus_nov_88.04%
Actinomyces slackii_nov_90.12%
Fusobacterium sp._HMT_248_nov_86.16%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Actinomyces sp._Oral_Taxon_171
Streptococcus suis_nov_88.37%
Fusobacterium nucleatum_ss_polymorphum
Arenibacter algicola_nov_84.87%
Alistipes indistinctus_nov_86.39%
Fusobacterium nucleatum
Devriesea agamarum_nov_86.9%
Isoptericola jiangsuensis_nov_88.29%
Kocuria indica_nov_89.81%
Actinomyces gerencseriae
Fusobacterium periodonticum
Streptococcus rifensis_nov_90.59%
Lactobacillus gasseri
Streptococcus sp._str._2136FAA
Streptococcus gordonii_sp._str._2136FAA
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Geosporobacter subterraneus_nov_84.49%
Streptococcus anginosus
Vagococcus lutrae_nov_87.66%
Streptococcus rifensis_nov_90.82%
Fusobacterium hwasookii_nov_87.47%
Streptococcus rifensis_nov_90.65%
Streptococcus pyogenes_nov_91.53%
Clostridium acetobutylicum_nov_85.65%
Dialister invisus
Streptococcus gordonii
Corynebacterium terpenotabidum_nov_92.31%
Fusobacterium nucleatum_nov_86.73%
Streptococcus pyogenes_nov_89.52%
Sneathia sanguinegens_nov_85.05%
Streptococcus intermedius
Cellulomonas gilvus_nov_87.87%
Streptococcus rifensis_nov_90.43%
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Bowdeniella nasicola_nov_86.78%
Corynebacterium terpenotabidum_nov_92.32%
Streptococcus rifensis_nov_90.67%
Veillonella parvula
Streptococcus rifensis_nov_91.68%
Streptococcus pyogenes_nov_89.52%
Mycobacterium multispecies_sppn9_2_nov_83.57%
Streptococcus mutans
Species
F5677.S651
F5677.S598
F5677.S675
F5677.S593
F5677.S650
F5677.S591
F5677.S520
F5677.S566
F5677.S517
F5677.S589
F5677.S513
F5677.S533
F5677.S590
F5677.S574
F5677.S626
F5677.S576
F5677.S556
F5677.S661
F5677.S542
F5677.S645
F5677.S665
F5677.S565
F5677.S617
F5677.S609
F5677.S592
F5677.S569
F5677.S674
F5677.S606
F5677.S682
F5677.S673
F5677.S641
F5677.S611
F5677.S587
F5677.S614
F5677.S619
F5677.S658
F5677.S588
F5677.S646
F5677.S605
F5677.S543
Samples