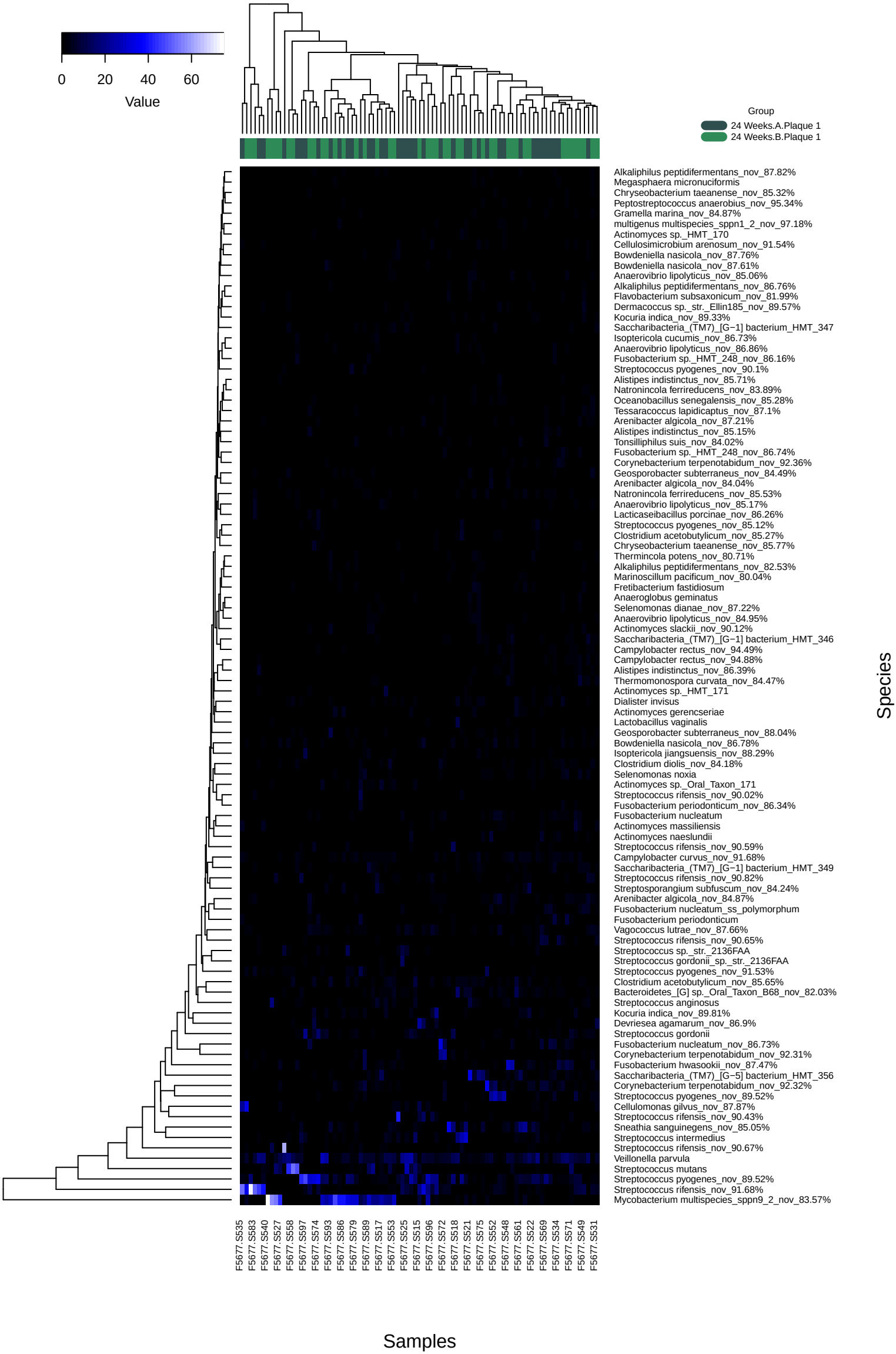
0
20
40
60
Value
Group
24 Weeks.A.Plaque 1
24 Weeks.B.Plaque 1
Alkaliphilus peptidifermentans_nov_87.82%
Megasphaera micronuciformis
Chryseobacterium taeanense_nov_85.32%
Peptostreptococcus anaerobius_nov_95.34%
Gramella marina_nov_84.87%
multigenus multispecies_sppn1_2_nov_97.18%
Actinomyces sp._HMT_170
Cellulosimicrobium arenosum_nov_91.54%
Bowdeniella nasicola_nov_87.76%
Bowdeniella nasicola_nov_87.61%
Anaerovibrio lipolyticus_nov_85.06%
Alkaliphilus peptidifermentans_nov_86.76%
Flavobacterium subsaxonicum_nov_81.99%
Dermacoccus sp._str._Ellin185_nov_89.57%
Kocuria indica_nov_89.33%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Isoptericola cucumis_nov_86.73%
Anaerovibrio lipolyticus_nov_86.86%
Fusobacterium sp._HMT_248_nov_86.16%
Streptococcus pyogenes_nov_90.1%
Alistipes indistinctus_nov_85.71%
Natronincola ferrireducens_nov_83.89%
Oceanobacillus senegalensis_nov_85.28%
Tessaracoccus lapidicaptus_nov_87.1%
Arenibacter algicola_nov_87.21%
Alistipes indistinctus_nov_85.15%
Tonsilliphilus suis_nov_84.02%
Fusobacterium sp._HMT_248_nov_86.74%
Corynebacterium terpenotabidum_nov_92.36%
Geosporobacter subterraneus_nov_84.49%
Arenibacter algicola_nov_84.04%
Natronincola ferrireducens_nov_85.53%
Anaerovibrio lipolyticus_nov_85.17%
Lacticaseibacillus porcinae_nov_86.26%
Streptococcus pyogenes_nov_85.12%
Clostridium acetobutylicum_nov_85.27%
Chryseobacterium taeanense_nov_85.77%
Thermincola potens_nov_80.71%
Alkaliphilus peptidifermentans_nov_82.53%
Marinoscillum pacificum_nov_80.04%
Fretibacterium fastidiosum
Anaeroglobus geminatus
Selenomonas dianae_nov_87.22%
Anaerovibrio lipolyticus_nov_84.95%
Actinomyces slackii_nov_90.12%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Campylobacter rectus_nov_94.49%
Campylobacter rectus_nov_94.88%
Alistipes indistinctus_nov_86.39%
Thermomonospora curvata_nov_84.47%
Actinomyces sp._HMT_171
Dialister invisus
Actinomyces gerencseriae
Lactobacillus vaginalis
Geosporobacter subterraneus_nov_88.04%
Bowdeniella nasicola_nov_86.78%
Isoptericola jiangsuensis_nov_88.29%
Clostridium diolis_nov_84.18%
Selenomonas noxia
Actinomyces sp._Oral_Taxon_171
Streptococcus rifensis_nov_90.02%
Fusobacterium periodonticum_nov_86.34%
Fusobacterium nucleatum
Actinomyces massiliensis
Actinomyces naeslundii
Streptococcus rifensis_nov_90.59%
Campylobacter curvus_nov_91.68%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Streptococcus rifensis_nov_90.82%
Streptosporangium subfuscum_nov_84.24%
Arenibacter algicola_nov_84.87%
Fusobacterium nucleatum_ss_polymorphum
Fusobacterium periodonticum
Vagococcus lutrae_nov_87.66%
Streptococcus rifensis_nov_90.65%
Streptococcus sp._str._2136FAA
Streptococcus gordonii_sp._str._2136FAA
Streptococcus pyogenes_nov_91.53%
Clostridium acetobutylicum_nov_85.65%
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Streptococcus anginosus
Kocuria indica_nov_89.81%
Devriesea agamarum_nov_86.9%
Streptococcus gordonii
Fusobacterium nucleatum_nov_86.73%
Corynebacterium terpenotabidum_nov_92.31%
Fusobacterium hwasookii_nov_87.47%
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Corynebacterium terpenotabidum_nov_92.32%
Streptococcus pyogenes_nov_89.52%
Cellulomonas gilvus_nov_87.87%
Streptococcus rifensis_nov_90.43%
Sneathia sanguinegens_nov_85.05%
Streptococcus intermedius
Streptococcus rifensis_nov_90.67%
Veillonella parvula
Streptococcus mutans
Streptococcus pyogenes_nov_89.52%
Streptococcus rifensis_nov_91.68%
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S535
F5677.S583
F5677.S540
F5677.S527
F5677.S558
F5677.S597
F5677.S574
F5677.S593
F5677.S586
F5677.S579
F5677.S589
F5677.S517
F5677.S553
F5677.S525
F5677.S515
F5677.S596
F5677.S572
F5677.S518
F5677.S521
F5677.S575
F5677.S552
F5677.S548
F5677.S561
F5677.S522
F5677.S569
F5677.S534
F5677.S571
F5677.S549
F5677.S531
Samples