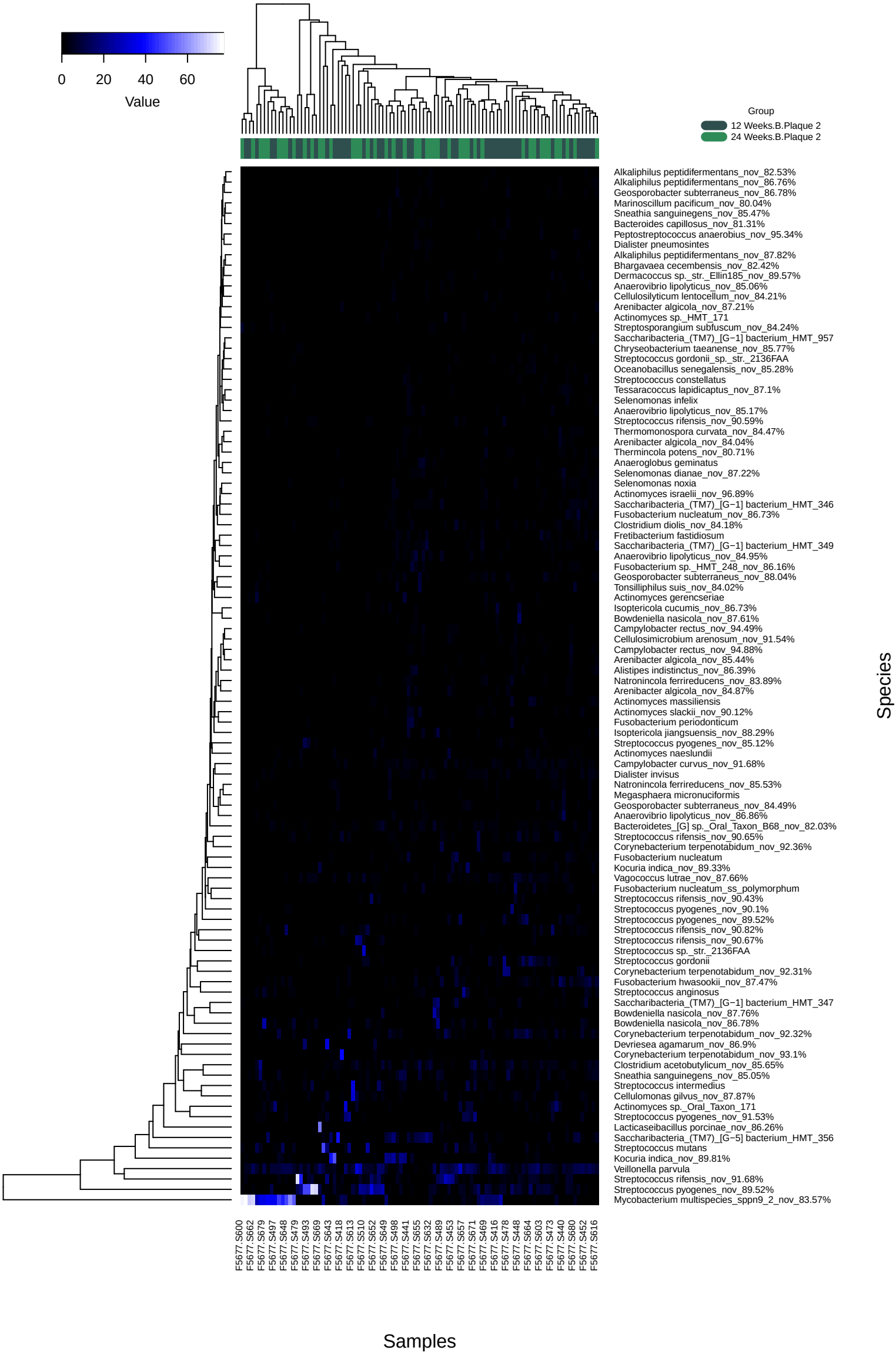
0
20
40
60
Value
Group
12 Weeks.B.Plaque 2
24 Weeks.B.Plaque 2
Alkaliphilus peptidifermentans_nov_82.53%
Alkaliphilus peptidifermentans_nov_86.76%
Geosporobacter subterraneus_nov_86.78%
Marinoscillum pacificum_nov_80.04%
Sneathia sanguinegens_nov_85.47%
Bacteroides capillosus_nov_81.31%
Peptostreptococcus anaerobius_nov_95.34%
Dialister pneumosintes
Alkaliphilus peptidifermentans_nov_87.82%
Bhargavaea cecembensis_nov_82.42%
Dermacoccus sp._str._Ellin185_nov_89.57%
Anaerovibrio lipolyticus_nov_85.06%
Cellulosilyticum lentocellum_nov_84.21%
Arenibacter algicola_nov_87.21%
Actinomyces sp._HMT_171
Streptosporangium subfuscum_nov_84.24%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_957
Chryseobacterium taeanense_nov_85.77%
Streptococcus gordonii_sp._str._2136FAA
Oceanobacillus senegalensis_nov_85.28%
Streptococcus constellatus
Tessaracoccus lapidicaptus_nov_87.1%
Selenomonas infelix
Anaerovibrio lipolyticus_nov_85.17%
Streptococcus rifensis_nov_90.59%
Thermomonospora curvata_nov_84.47%
Arenibacter algicola_nov_84.04%
Thermincola potens_nov_80.71%
Anaeroglobus geminatus
Selenomonas dianae_nov_87.22%
Selenomonas noxia
Actinomyces israelii_nov_96.89%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Fusobacterium nucleatum_nov_86.73%
Clostridium diolis_nov_84.18%
Fretibacterium fastidiosum
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Anaerovibrio lipolyticus_nov_84.95%
Fusobacterium sp._HMT_248_nov_86.16%
Geosporobacter subterraneus_nov_88.04%
Tonsilliphilus suis_nov_84.02%
Actinomyces gerencseriae
Isoptericola cucumis_nov_86.73%
Bowdeniella nasicola_nov_87.61%
Campylobacter rectus_nov_94.49%
Cellulosimicrobium arenosum_nov_91.54%
Campylobacter rectus_nov_94.88%
Arenibacter algicola_nov_85.44%
Alistipes indistinctus_nov_86.39%
Natronincola ferrireducens_nov_83.89%
Arenibacter algicola_nov_84.87%
Actinomyces massiliensis
Actinomyces slackii_nov_90.12%
Fusobacterium periodonticum
Isoptericola jiangsuensis_nov_88.29%
Streptococcus pyogenes_nov_85.12%
Actinomyces naeslundii
Campylobacter curvus_nov_91.68%
Dialister invisus
Natronincola ferrireducens_nov_85.53%
Megasphaera micronuciformis
Geosporobacter subterraneus_nov_84.49%
Anaerovibrio lipolyticus_nov_86.86%
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Streptococcus rifensis_nov_90.65%
Corynebacterium terpenotabidum_nov_92.36%
Fusobacterium nucleatum
Kocuria indica_nov_89.33%
Vagococcus lutrae_nov_87.66%
Fusobacterium nucleatum_ss_polymorphum
Streptococcus rifensis_nov_90.43%
Streptococcus pyogenes_nov_90.1%
Streptococcus pyogenes_nov_89.52%
Streptococcus rifensis_nov_90.82%
Streptococcus rifensis_nov_90.67%
Streptococcus sp._str._2136FAA
Streptococcus gordonii
Corynebacterium terpenotabidum_nov_92.31%
Fusobacterium hwasookii_nov_87.47%
Streptococcus anginosus
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Bowdeniella nasicola_nov_87.76%
Bowdeniella nasicola_nov_86.78%
Corynebacterium terpenotabidum_nov_92.32%
Devriesea agamarum_nov_86.9%
Corynebacterium terpenotabidum_nov_93.1%
Clostridium acetobutylicum_nov_85.65%
Sneathia sanguinegens_nov_85.05%
Streptococcus intermedius
Cellulomonas gilvus_nov_87.87%
Actinomyces sp._Oral_Taxon_171
Streptococcus pyogenes_nov_91.53%
Lacticaseibacillus porcinae_nov_86.26%
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Streptococcus mutans
Kocuria indica_nov_89.81%
Veillonella parvula
Streptococcus rifensis_nov_91.68%
Streptococcus pyogenes_nov_89.52%
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S600
F5677.S662
F5677.S679
F5677.S497
F5677.S648
F5677.S479
F5677.S493
F5677.S669
F5677.S643
F5677.S418
F5677.S613
F5677.S510
F5677.S652
F5677.S649
F5677.S498
F5677.S441
F5677.S655
F5677.S632
F5677.S489
F5677.S453
F5677.S657
F5677.S671
F5677.S469
F5677.S416
F5677.S478
F5677.S448
F5677.S664
F5677.S603
F5677.S473
F5677.S440
F5677.S680
F5677.S452
F5677.S616
Samples