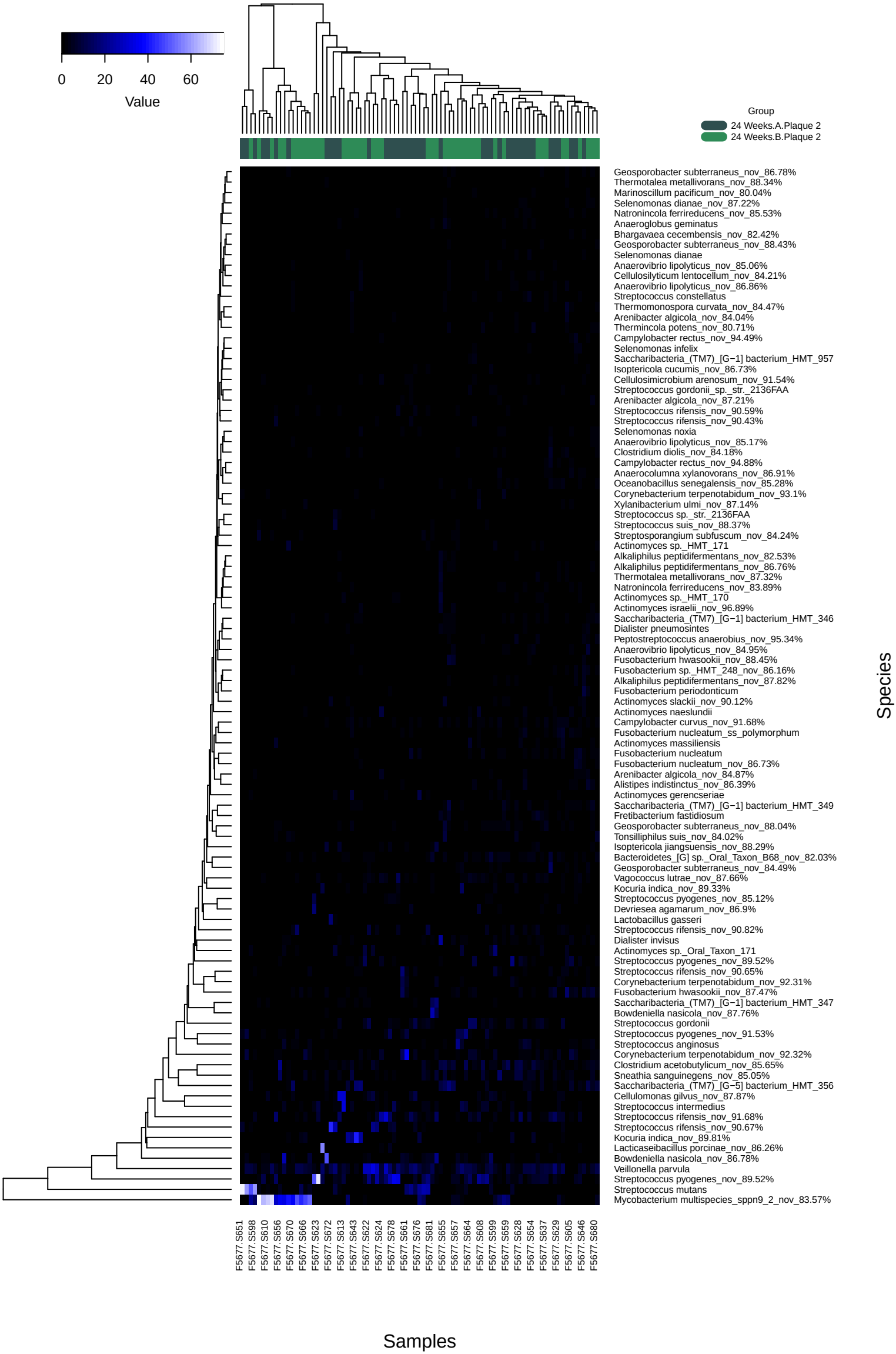
0
20
40
60
Value
Group
24 Weeks.A.Plaque 2
24 Weeks.B.Plaque 2
Geosporobacter subterraneus_nov_86.78%
Thermotalea metallivorans_nov_88.34%
Marinoscillum pacificum_nov_80.04%
Selenomonas dianae_nov_87.22%
Natronincola ferrireducens_nov_85.53%
Anaeroglobus geminatus
Bhargavaea cecembensis_nov_82.42%
Geosporobacter subterraneus_nov_88.43%
Selenomonas dianae
Anaerovibrio lipolyticus_nov_85.06%
Cellulosilyticum lentocellum_nov_84.21%
Anaerovibrio lipolyticus_nov_86.86%
Streptococcus constellatus
Thermomonospora curvata_nov_84.47%
Arenibacter algicola_nov_84.04%
Thermincola potens_nov_80.71%
Campylobacter rectus_nov_94.49%
Selenomonas infelix
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_957
Isoptericola cucumis_nov_86.73%
Cellulosimicrobium arenosum_nov_91.54%
Streptococcus gordonii_sp._str._2136FAA
Arenibacter algicola_nov_87.21%
Streptococcus rifensis_nov_90.59%
Streptococcus rifensis_nov_90.43%
Selenomonas noxia
Anaerovibrio lipolyticus_nov_85.17%
Clostridium diolis_nov_84.18%
Campylobacter rectus_nov_94.88%
Anaerocolumna xylanovorans_nov_86.91%
Oceanobacillus senegalensis_nov_85.28%
Corynebacterium terpenotabidum_nov_93.1%
Xylanibacterium ulmi_nov_87.14%
Streptococcus sp._str._2136FAA
Streptococcus suis_nov_88.37%
Streptosporangium subfuscum_nov_84.24%
Actinomyces sp._HMT_171
Alkaliphilus peptidifermentans_nov_82.53%
Alkaliphilus peptidifermentans_nov_86.76%
Thermotalea metallivorans_nov_87.32%
Natronincola ferrireducens_nov_83.89%
Actinomyces sp._HMT_170
Actinomyces israelii_nov_96.89%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Dialister pneumosintes
Peptostreptococcus anaerobius_nov_95.34%
Anaerovibrio lipolyticus_nov_84.95%
Fusobacterium hwasookii_nov_88.45%
Fusobacterium sp._HMT_248_nov_86.16%
Alkaliphilus peptidifermentans_nov_87.82%
Fusobacterium periodonticum
Actinomyces slackii_nov_90.12%
Actinomyces naeslundii
Campylobacter curvus_nov_91.68%
Fusobacterium nucleatum_ss_polymorphum
Actinomyces massiliensis
Fusobacterium nucleatum
Fusobacterium nucleatum_nov_86.73%
Arenibacter algicola_nov_84.87%
Alistipes indistinctus_nov_86.39%
Actinomyces gerencseriae
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Fretibacterium fastidiosum
Geosporobacter subterraneus_nov_88.04%
Tonsilliphilus suis_nov_84.02%
Isoptericola jiangsuensis_nov_88.29%
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Geosporobacter subterraneus_nov_84.49%
Vagococcus lutrae_nov_87.66%
Kocuria indica_nov_89.33%
Streptococcus pyogenes_nov_85.12%
Devriesea agamarum_nov_86.9%
Lactobacillus gasseri
Streptococcus rifensis_nov_90.82%
Dialister invisus
Actinomyces sp._Oral_Taxon_171
Streptococcus pyogenes_nov_89.52%
Streptococcus rifensis_nov_90.65%
Corynebacterium terpenotabidum_nov_92.31%
Fusobacterium hwasookii_nov_87.47%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Bowdeniella nasicola_nov_87.76%
Streptococcus gordonii
Streptococcus pyogenes_nov_91.53%
Streptococcus anginosus
Corynebacterium terpenotabidum_nov_92.32%
Clostridium acetobutylicum_nov_85.65%
Sneathia sanguinegens_nov_85.05%
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Cellulomonas gilvus_nov_87.87%
Streptococcus intermedius
Streptococcus rifensis_nov_91.68%
Streptococcus rifensis_nov_90.67%
Kocuria indica_nov_89.81%
Lacticaseibacillus porcinae_nov_86.26%
Bowdeniella nasicola_nov_86.78%
Veillonella parvula
Streptococcus pyogenes_nov_89.52%
Streptococcus mutans
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S651
F5677.S598
F5677.S610
F5677.S656
F5677.S670
F5677.S666
F5677.S623
F5677.S672
F5677.S613
F5677.S643
F5677.S622
F5677.S624
F5677.S678
F5677.S661
F5677.S676
F5677.S681
F5677.S655
F5677.S657
F5677.S664
F5677.S608
F5677.S599
F5677.S659
F5677.S628
F5677.S654
F5677.S637
F5677.S629
F5677.S605
F5677.S646
F5677.S680
Samples