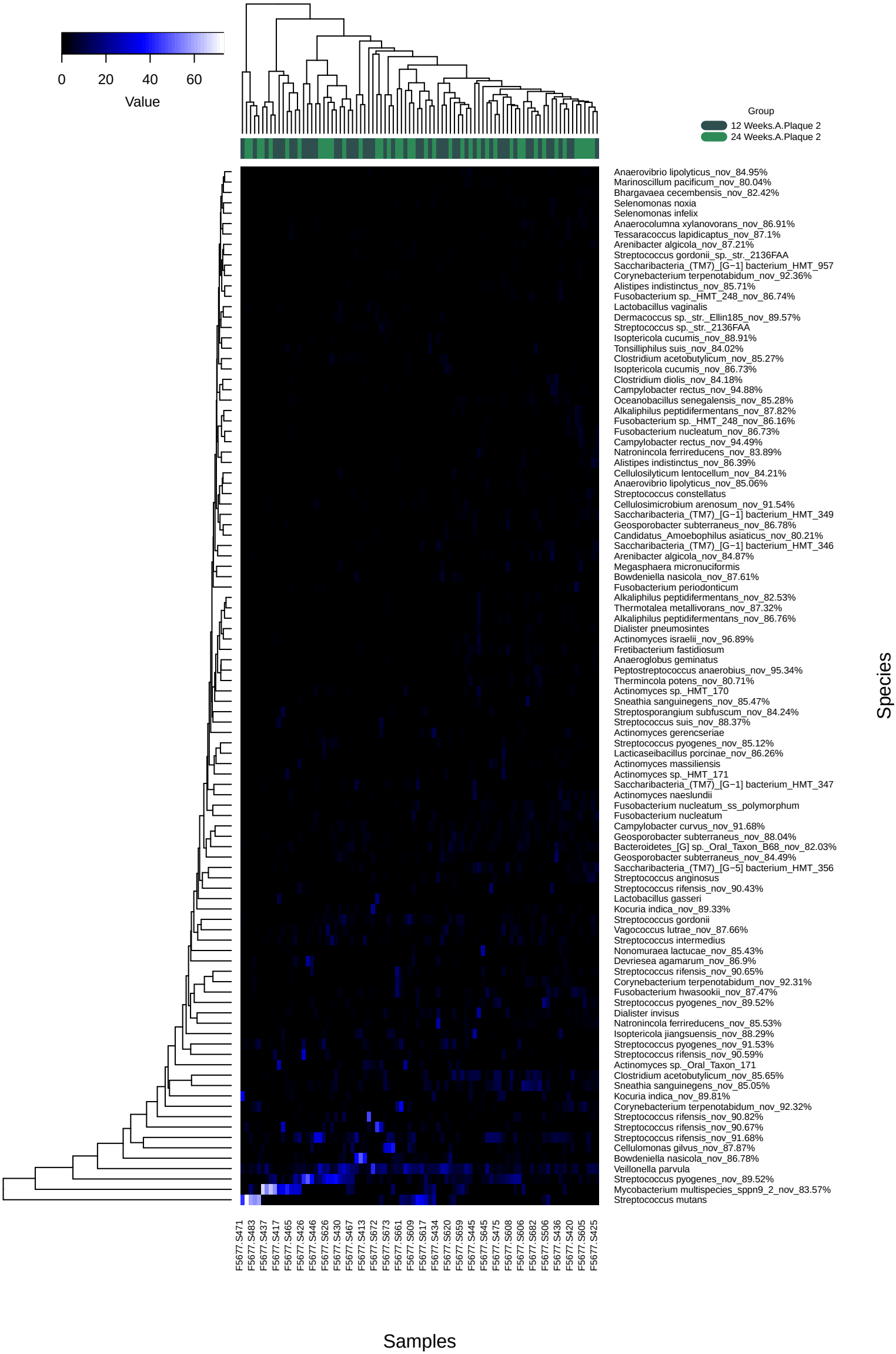
0
20
40
60
Value
Group
12 Weeks.A.Plaque 2
24 Weeks.A.Plaque 2
Anaerovibrio lipolyticus_nov_84.95%
Marinoscillum pacificum_nov_80.04%
Bhargavaea cecembensis_nov_82.42%
Selenomonas noxia
Selenomonas infelix
Anaerocolumna xylanovorans_nov_86.91%
Tessaracoccus lapidicaptus_nov_87.1%
Arenibacter algicola_nov_87.21%
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._Ellin185_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%
Lacticaseibacillus 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.S608
F5677.S606
F5677.S682
F5677.S506
F5677.S436
F5677.S420
F5677.S605
F5677.S425
Samples