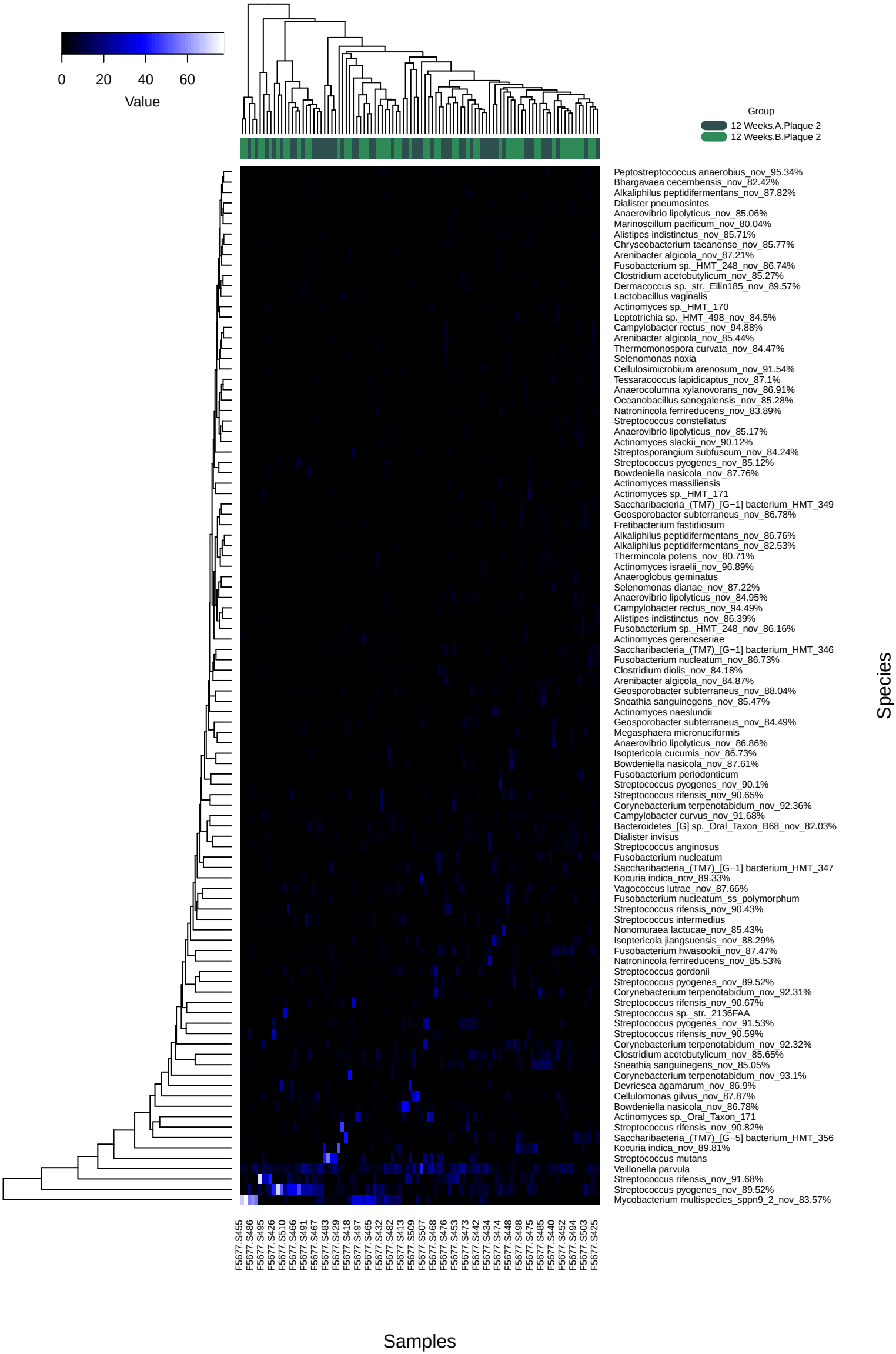
0
20
40
60
Value
Group
12 Weeks.A.Plaque 2
12 Weeks.B.Plaque 2
Peptostreptococcus anaerobius_nov_95.34%
Bhargavaea cecembensis_nov_82.42%
Alkaliphilus peptidifermentans_nov_87.82%
Dialister pneumosintes
Anaerovibrio lipolyticus_nov_85.06%
Marinoscillum pacificum_nov_80.04%
Alistipes indistinctus_nov_85.71%
Chryseobacterium taeanense_nov_85.77%
Arenibacter algicola_nov_87.21%
Fusobacterium sp._HMT_248_nov_86.74%
Clostridium acetobutylicum_nov_85.27%
Dermacoccus sp._str._Ellin185_nov_89.57%
Lactobacillus vaginalis
Actinomyces sp._HMT_170
Leptotrichia sp._HMT_498_nov_84.5%
Campylobacter rectus_nov_94.88%
Arenibacter algicola_nov_85.44%
Thermomonospora curvata_nov_84.47%
Selenomonas noxia
Cellulosimicrobium arenosum_nov_91.54%
Tessaracoccus lapidicaptus_nov_87.1%
Anaerocolumna xylanovorans_nov_86.91%
Oceanobacillus senegalensis_nov_85.28%
Natronincola ferrireducens_nov_83.89%
Streptococcus constellatus
Anaerovibrio lipolyticus_nov_85.17%
Actinomyces slackii_nov_90.12%
Streptosporangium subfuscum_nov_84.24%
Streptococcus pyogenes_nov_85.12%
Bowdeniella nasicola_nov_87.76%
Actinomyces massiliensis
Actinomyces sp._HMT_171
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Geosporobacter subterraneus_nov_86.78%
Fretibacterium fastidiosum
Alkaliphilus peptidifermentans_nov_86.76%
Alkaliphilus peptidifermentans_nov_82.53%
Thermincola potens_nov_80.71%
Actinomyces israelii_nov_96.89%
Anaeroglobus geminatus
Selenomonas dianae_nov_87.22%
Anaerovibrio lipolyticus_nov_84.95%
Campylobacter rectus_nov_94.49%
Alistipes indistinctus_nov_86.39%
Fusobacterium sp._HMT_248_nov_86.16%
Actinomyces gerencseriae
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Fusobacterium nucleatum_nov_86.73%
Clostridium diolis_nov_84.18%
Arenibacter algicola_nov_84.87%
Geosporobacter subterraneus_nov_88.04%
Sneathia sanguinegens_nov_85.47%
Actinomyces naeslundii
Geosporobacter subterraneus_nov_84.49%
Megasphaera micronuciformis
Anaerovibrio lipolyticus_nov_86.86%
Isoptericola cucumis_nov_86.73%
Bowdeniella nasicola_nov_87.61%
Fusobacterium periodonticum
Streptococcus pyogenes_nov_90.1%
Streptococcus rifensis_nov_90.65%
Corynebacterium terpenotabidum_nov_92.36%
Campylobacter curvus_nov_91.68%
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Dialister invisus
Streptococcus anginosus
Fusobacterium nucleatum
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Kocuria indica_nov_89.33%
Vagococcus lutrae_nov_87.66%
Fusobacterium nucleatum_ss_polymorphum
Streptococcus rifensis_nov_90.43%
Streptococcus intermedius
Nonomuraea lactucae_nov_85.43%
Isoptericola jiangsuensis_nov_88.29%
Fusobacterium hwasookii_nov_87.47%
Natronincola ferrireducens_nov_85.53%
Streptococcus gordonii
Streptococcus pyogenes_nov_89.52%
Corynebacterium terpenotabidum_nov_92.31%
Streptococcus rifensis_nov_90.67%
Streptococcus sp._str._2136FAA
Streptococcus pyogenes_nov_91.53%
Streptococcus rifensis_nov_90.59%
Corynebacterium terpenotabidum_nov_92.32%
Clostridium acetobutylicum_nov_85.65%
Sneathia sanguinegens_nov_85.05%
Corynebacterium terpenotabidum_nov_93.1%
Devriesea agamarum_nov_86.9%
Cellulomonas gilvus_nov_87.87%
Bowdeniella nasicola_nov_86.78%
Actinomyces sp._Oral_Taxon_171
Streptococcus rifensis_nov_90.82%
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Kocuria indica_nov_89.81%
Streptococcus mutans
Veillonella parvula
Streptococcus rifensis_nov_91.68%
Streptococcus pyogenes_nov_89.52%
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S455
F5677.S486
F5677.S495
F5677.S426
F5677.S510
F5677.S466
F5677.S491
F5677.S467
F5677.S483
F5677.S429
F5677.S418
F5677.S497
F5677.S465
F5677.S432
F5677.S482
F5677.S413
F5677.S509
F5677.S507
F5677.S468
F5677.S476
F5677.S453
F5677.S473
F5677.S442
F5677.S434
F5677.S474
F5677.S448
F5677.S498
F5677.S475
F5677.S485
F5677.S440
F5677.S452
F5677.S494
F5677.S503
F5677.S425
Samples