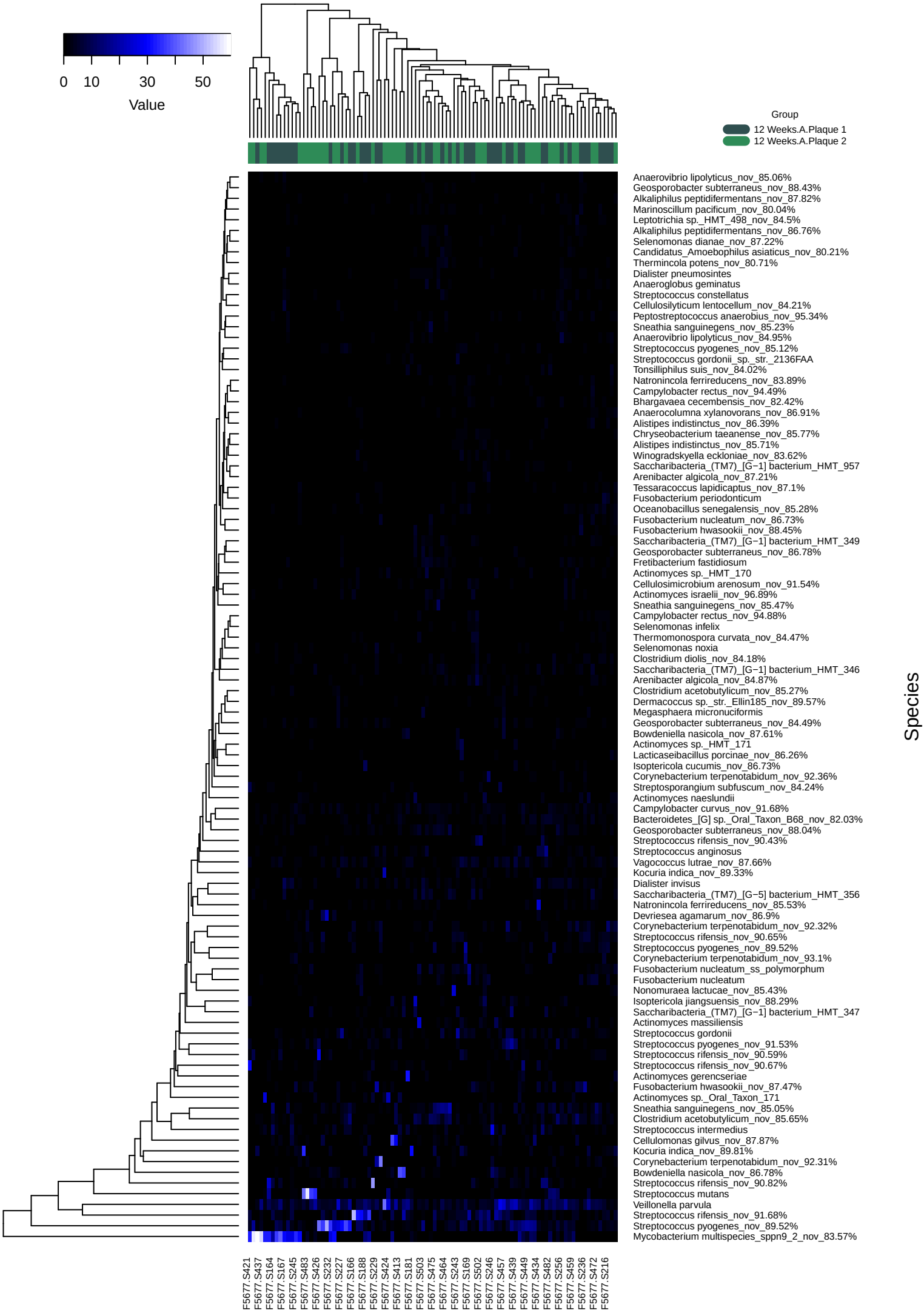
0
10
30
50
Value
Group
12 Weeks.A.Plaque 1
12 Weeks.A.Plaque 2
Anaerovibrio lipolyticus_nov_85.06%
Geosporobacter subterraneus_nov_88.43%
Alkaliphilus peptidifermentans_nov_87.82%
Marinoscillum pacificum_nov_80.04%
Leptotrichia sp._HMT_498_nov_84.5%
Alkaliphilus peptidifermentans_nov_86.76%
Selenomonas dianae_nov_87.22%
Candidatus_Amoebophilus asiaticus_nov_80.21%
Thermincola potens_nov_80.71%
Dialister pneumosintes
Anaeroglobus geminatus
Streptococcus constellatus
Cellulosilyticum lentocellum_nov_84.21%
Peptostreptococcus anaerobius_nov_95.34%
Sneathia sanguinegens_nov_85.23%
Anaerovibrio lipolyticus_nov_84.95%
Streptococcus pyogenes_nov_85.12%
Streptococcus gordonii_sp._str._2136FAA
Tonsilliphilus suis_nov_84.02%
Natronincola ferrireducens_nov_83.89%
Campylobacter rectus_nov_94.49%
Bhargavaea cecembensis_nov_82.42%
Anaerocolumna xylanovorans_nov_86.91%
Alistipes indistinctus_nov_86.39%
Chryseobacterium taeanense_nov_85.77%
Alistipes indistinctus_nov_85.71%
Winogradskyella eckloniae_nov_83.62%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_957
Arenibacter algicola_nov_87.21%
Tessaracoccus lapidicaptus_nov_87.1%
Fusobacterium periodonticum
Oceanobacillus senegalensis_nov_85.28%
Fusobacterium nucleatum_nov_86.73%
Fusobacterium hwasookii_nov_88.45%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Geosporobacter subterraneus_nov_86.78%
Fretibacterium fastidiosum
Actinomyces sp._HMT_170
Cellulosimicrobium arenosum_nov_91.54%
Actinomyces israelii_nov_96.89%
Sneathia sanguinegens_nov_85.47%
Campylobacter rectus_nov_94.88%
Selenomonas infelix
Thermomonospora curvata_nov_84.47%
Selenomonas noxia
Clostridium diolis_nov_84.18%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Arenibacter algicola_nov_84.87%
Clostridium acetobutylicum_nov_85.27%
Dermacoccus sp._str._Ellin185_nov_89.57%
Megasphaera micronuciformis
Geosporobacter subterraneus_nov_84.49%
Bowdeniella nasicola_nov_87.61%
Actinomyces sp._HMT_171
Lacticaseibacillus porcinae_nov_86.26%
Isoptericola cucumis_nov_86.73%
Corynebacterium terpenotabidum_nov_92.36%
Streptosporangium subfuscum_nov_84.24%
Actinomyces naeslundii
Campylobacter curvus_nov_91.68%
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Geosporobacter subterraneus_nov_88.04%
Streptococcus rifensis_nov_90.43%
Streptococcus anginosus
Vagococcus lutrae_nov_87.66%
Kocuria indica_nov_89.33%
Dialister invisus
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Natronincola ferrireducens_nov_85.53%
Devriesea agamarum_nov_86.9%
Corynebacterium terpenotabidum_nov_92.32%
Streptococcus rifensis_nov_90.65%
Streptococcus pyogenes_nov_89.52%
Corynebacterium terpenotabidum_nov_93.1%
Fusobacterium nucleatum_ss_polymorphum
Fusobacterium nucleatum
Nonomuraea lactucae_nov_85.43%
Isoptericola jiangsuensis_nov_88.29%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Actinomyces massiliensis
Streptococcus gordonii
Streptococcus pyogenes_nov_91.53%
Streptococcus rifensis_nov_90.59%
Streptococcus rifensis_nov_90.67%
Actinomyces gerencseriae
Fusobacterium hwasookii_nov_87.47%
Actinomyces sp._Oral_Taxon_171
Sneathia sanguinegens_nov_85.05%
Clostridium acetobutylicum_nov_85.65%
Streptococcus intermedius
Cellulomonas gilvus_nov_87.87%
Kocuria indica_nov_89.81%
Corynebacterium terpenotabidum_nov_92.31%
Bowdeniella nasicola_nov_86.78%
Streptococcus rifensis_nov_90.82%
Streptococcus mutans
Veillonella parvula
Streptococcus rifensis_nov_91.68%
Streptococcus pyogenes_nov_89.52%
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S421
F5677.S437
F5677.S164
F5677.S167
F5677.S245
F5677.S483
F5677.S426
F5677.S232
F5677.S227
F5677.S166
F5677.S188
F5677.S229
F5677.S424
F5677.S413
F5677.S181
F5677.S503
F5677.S475
F5677.S464
F5677.S243
F5677.S169
F5677.S502
F5677.S246
F5677.S457
F5677.S439
F5677.S449
F5677.S434
F5677.S482
F5677.S256
F5677.S459
F5677.S236
F5677.S472
F5677.S216
Samples