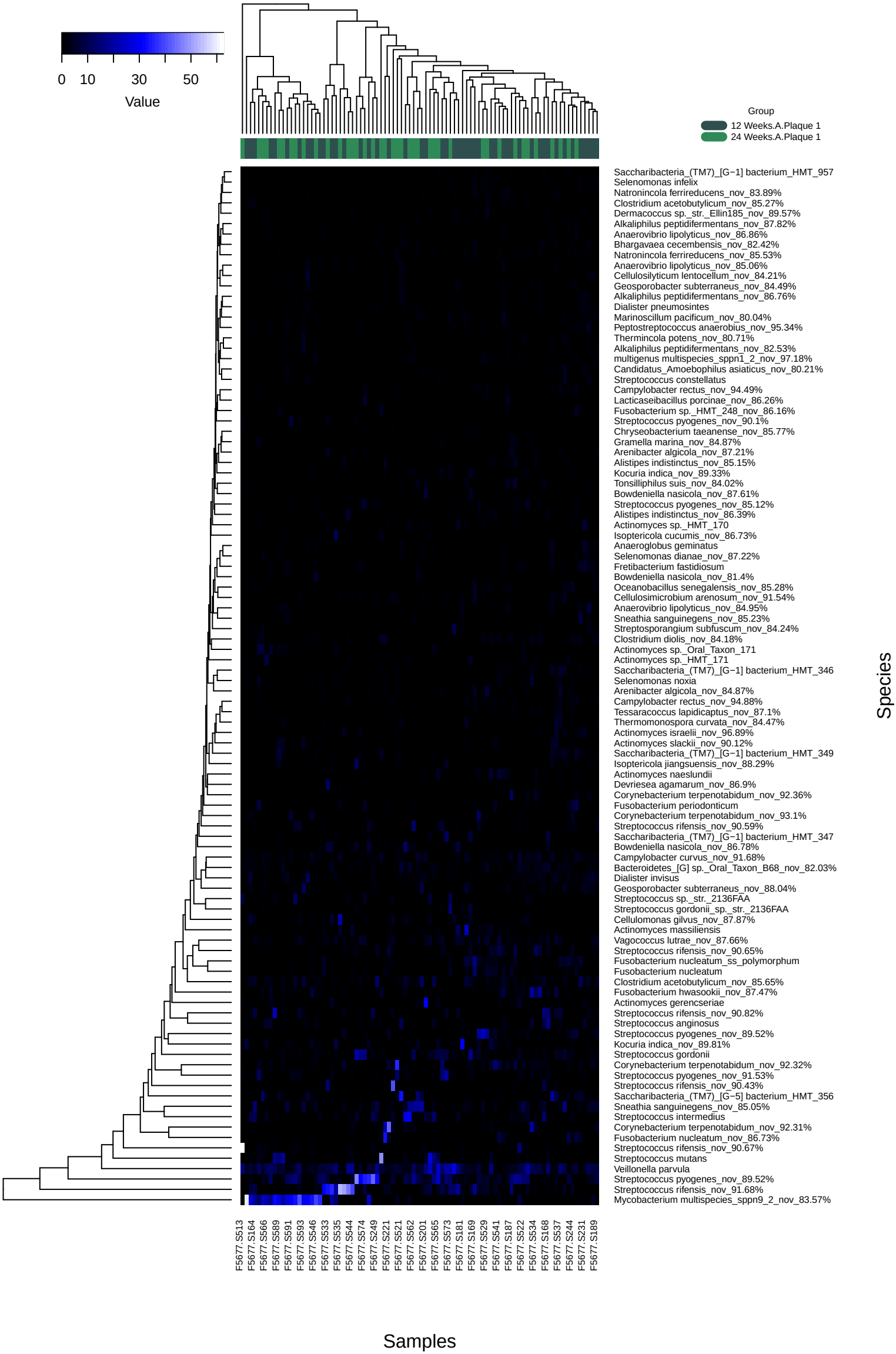
0
10
30
50
Value
Group
12 Weeks.A.Plaque 1
24 Weeks.A.Plaque 1
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_957
Selenomonas infelix
Natronincola ferrireducens_nov_83.89%
Clostridium acetobutylicum_nov_85.27%
Dermacoccus sp._str._Ellin185_nov_89.57%
Alkaliphilus peptidifermentans_nov_87.82%
Anaerovibrio lipolyticus_nov_86.86%
Bhargavaea cecembensis_nov_82.42%
Natronincola ferrireducens_nov_85.53%
Anaerovibrio lipolyticus_nov_85.06%
Cellulosilyticum lentocellum_nov_84.21%
Geosporobacter subterraneus_nov_84.49%
Alkaliphilus peptidifermentans_nov_86.76%
Dialister pneumosintes
Marinoscillum pacificum_nov_80.04%
Peptostreptococcus anaerobius_nov_95.34%
Thermincola potens_nov_80.71%
Alkaliphilus peptidifermentans_nov_82.53%
multigenus multispecies_sppn1_2_nov_97.18%
Candidatus_Amoebophilus asiaticus_nov_80.21%
Streptococcus constellatus
Campylobacter rectus_nov_94.49%
Lacticaseibacillus porcinae_nov_86.26%
Fusobacterium sp._HMT_248_nov_86.16%
Streptococcus pyogenes_nov_90.1%
Chryseobacterium taeanense_nov_85.77%
Gramella marina_nov_84.87%
Arenibacter algicola_nov_87.21%
Alistipes indistinctus_nov_85.15%
Kocuria indica_nov_89.33%
Tonsilliphilus suis_nov_84.02%
Bowdeniella nasicola_nov_87.61%
Streptococcus pyogenes_nov_85.12%
Alistipes indistinctus_nov_86.39%
Actinomyces sp._HMT_170
Isoptericola cucumis_nov_86.73%
Anaeroglobus geminatus
Selenomonas dianae_nov_87.22%
Fretibacterium fastidiosum
Bowdeniella nasicola_nov_81.4%
Oceanobacillus senegalensis_nov_85.28%
Cellulosimicrobium arenosum_nov_91.54%
Anaerovibrio lipolyticus_nov_84.95%
Sneathia sanguinegens_nov_85.23%
Streptosporangium subfuscum_nov_84.24%
Clostridium diolis_nov_84.18%
Actinomyces sp._Oral_Taxon_171
Actinomyces sp._HMT_171
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Selenomonas noxia
Arenibacter algicola_nov_84.87%
Campylobacter rectus_nov_94.88%
Tessaracoccus lapidicaptus_nov_87.1%
Thermomonospora curvata_nov_84.47%
Actinomyces israelii_nov_96.89%
Actinomyces slackii_nov_90.12%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Isoptericola jiangsuensis_nov_88.29%
Actinomyces naeslundii
Devriesea agamarum_nov_86.9%
Corynebacterium terpenotabidum_nov_92.36%
Fusobacterium periodonticum
Corynebacterium terpenotabidum_nov_93.1%
Streptococcus rifensis_nov_90.59%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Bowdeniella nasicola_nov_86.78%
Campylobacter curvus_nov_91.68%
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Dialister invisus
Geosporobacter subterraneus_nov_88.04%
Streptococcus sp._str._2136FAA
Streptococcus gordonii_sp._str._2136FAA
Cellulomonas gilvus_nov_87.87%
Actinomyces massiliensis
Vagococcus lutrae_nov_87.66%
Streptococcus rifensis_nov_90.65%
Fusobacterium nucleatum_ss_polymorphum
Fusobacterium nucleatum
Clostridium acetobutylicum_nov_85.65%
Fusobacterium hwasookii_nov_87.47%
Actinomyces gerencseriae
Streptococcus rifensis_nov_90.82%
Streptococcus anginosus
Streptococcus pyogenes_nov_89.52%
Kocuria indica_nov_89.81%
Streptococcus gordonii
Corynebacterium terpenotabidum_nov_92.32%
Streptococcus pyogenes_nov_91.53%
Streptococcus rifensis_nov_90.43%
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Sneathia sanguinegens_nov_85.05%
Streptococcus intermedius
Corynebacterium terpenotabidum_nov_92.31%
Fusobacterium nucleatum_nov_86.73%
Streptococcus rifensis_nov_90.67%
Streptococcus mutans
Veillonella parvula
Streptococcus pyogenes_nov_89.52%
Streptococcus rifensis_nov_91.68%
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S513
F5677.S164
F5677.S566
F5677.S589
F5677.S591
F5677.S593
F5677.S546
F5677.S533
F5677.S535
F5677.S544
F5677.S574
F5677.S249
F5677.S221
F5677.S521
F5677.S562
F5677.S201
F5677.S565
F5677.S573
F5677.S181
F5677.S169
F5677.S529
F5677.S541
F5677.S187
F5677.S522
F5677.S534
F5677.S168
F5677.S537
F5677.S244
F5677.S231
F5677.S189
Samples