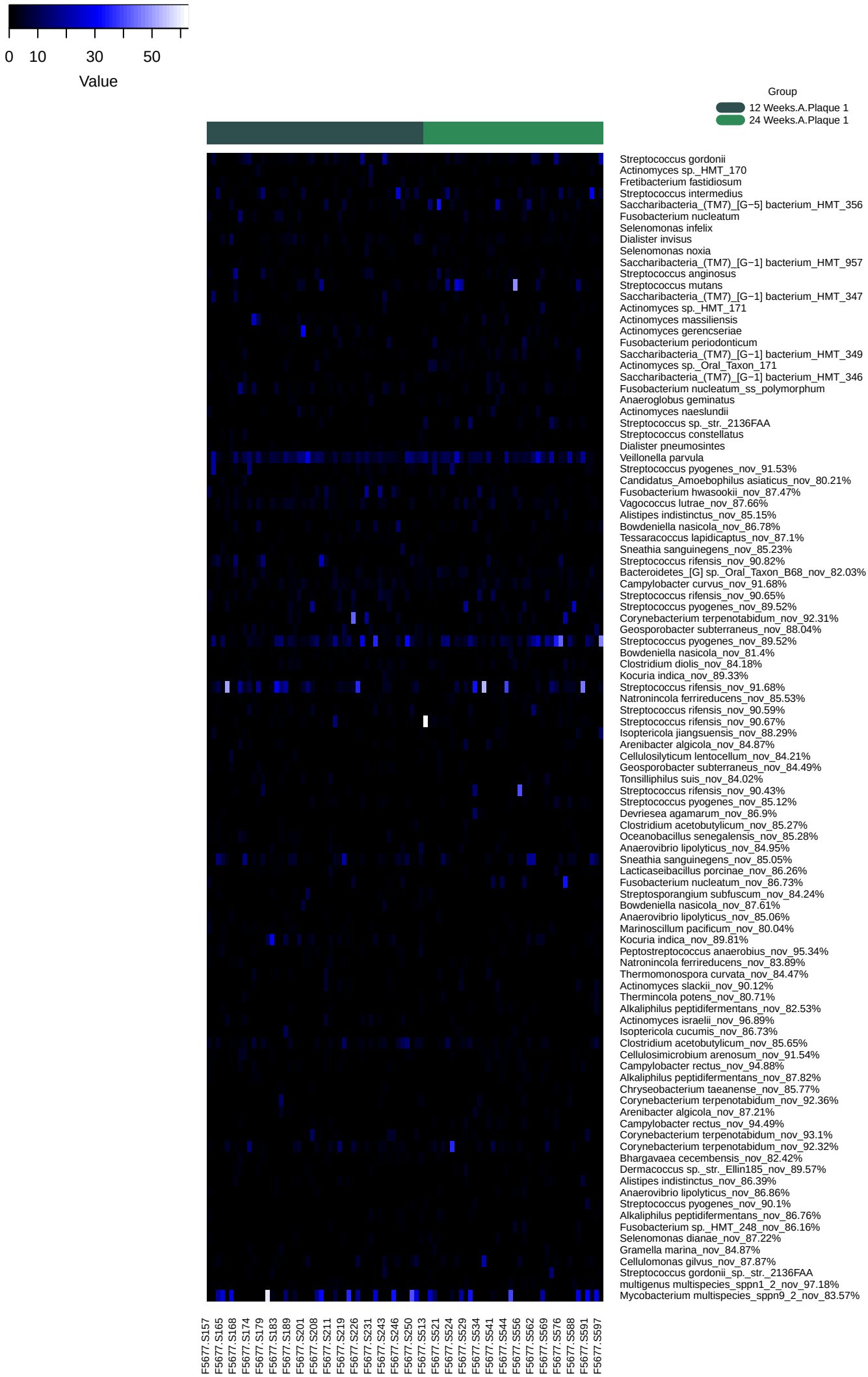
0
10
30
50
Value
Group
12 Weeks.A.Plaque 1
24 Weeks.A.Plaque 1
Streptococcus gordonii
Actinomyces sp._HMT_170
Fretibacterium fastidiosum
Streptococcus intermedius
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Fusobacterium nucleatum
Selenomonas infelix
Dialister invisus
Selenomonas noxia
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_957
Streptococcus anginosus
Streptococcus mutans
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Actinomyces sp._HMT_171
Actinomyces massiliensis
Actinomyces gerencseriae
Fusobacterium periodonticum
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Actinomyces sp._Oral_Taxon_171
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Fusobacterium nucleatum_ss_polymorphum
Anaeroglobus geminatus
Actinomyces naeslundii
Streptococcus sp._str._2136FAA
Streptococcus constellatus
Dialister pneumosintes
Veillonella parvula
Streptococcus pyogenes_nov_91.53%
Candidatus_Amoebophilus asiaticus_nov_80.21%
Fusobacterium hwasookii_nov_87.47%
Vagococcus lutrae_nov_87.66%
Alistipes indistinctus_nov_85.15%
Bowdeniella nasicola_nov_86.78%
Tessaracoccus lapidicaptus_nov_87.1%
Sneathia sanguinegens_nov_85.23%
Streptococcus rifensis_nov_90.82%
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Campylobacter curvus_nov_91.68%
Streptococcus rifensis_nov_90.65%
Streptococcus pyogenes_nov_89.52%
Corynebacterium terpenotabidum_nov_92.31%
Geosporobacter subterraneus_nov_88.04%
Streptococcus pyogenes_nov_89.52%
Bowdeniella nasicola_nov_81.4%
Clostridium diolis_nov_84.18%
Kocuria indica_nov_89.33%
Streptococcus rifensis_nov_91.68%
Natronincola ferrireducens_nov_85.53%
Streptococcus rifensis_nov_90.59%
Streptococcus rifensis_nov_90.67%
Isoptericola jiangsuensis_nov_88.29%
Arenibacter algicola_nov_84.87%
Cellulosilyticum lentocellum_nov_84.21%
Geosporobacter subterraneus_nov_84.49%
Tonsilliphilus suis_nov_84.02%
Streptococcus rifensis_nov_90.43%
Streptococcus pyogenes_nov_85.12%
Devriesea agamarum_nov_86.9%
Clostridium acetobutylicum_nov_85.27%
Oceanobacillus senegalensis_nov_85.28%
Anaerovibrio lipolyticus_nov_84.95%
Sneathia sanguinegens_nov_85.05%
Lacticaseibacillus porcinae_nov_86.26%
Fusobacterium nucleatum_nov_86.73%
Streptosporangium subfuscum_nov_84.24%
Bowdeniella nasicola_nov_87.61%
Anaerovibrio lipolyticus_nov_85.06%
Marinoscillum pacificum_nov_80.04%
Kocuria indica_nov_89.81%
Peptostreptococcus anaerobius_nov_95.34%
Natronincola ferrireducens_nov_83.89%
Thermomonospora curvata_nov_84.47%
Actinomyces slackii_nov_90.12%
Thermincola potens_nov_80.71%
Alkaliphilus peptidifermentans_nov_82.53%
Actinomyces israelii_nov_96.89%
Isoptericola cucumis_nov_86.73%
Clostridium acetobutylicum_nov_85.65%
Cellulosimicrobium arenosum_nov_91.54%
Campylobacter rectus_nov_94.88%
Alkaliphilus peptidifermentans_nov_87.82%
Chryseobacterium taeanense_nov_85.77%
Corynebacterium terpenotabidum_nov_92.36%
Arenibacter algicola_nov_87.21%
Campylobacter rectus_nov_94.49%
Corynebacterium terpenotabidum_nov_93.1%
Corynebacterium terpenotabidum_nov_92.32%
Bhargavaea cecembensis_nov_82.42%
Dermacoccus sp._str._Ellin185_nov_89.57%
Alistipes indistinctus_nov_86.39%
Anaerovibrio lipolyticus_nov_86.86%
Streptococcus pyogenes_nov_90.1%
Alkaliphilus peptidifermentans_nov_86.76%
Fusobacterium sp._HMT_248_nov_86.16%
Selenomonas dianae_nov_87.22%
Gramella marina_nov_84.87%
Cellulomonas gilvus_nov_87.87%
Streptococcus gordonii_sp._str._2136FAA
multigenus multispecies_sppn1_2_nov_97.18%
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S157
F5677.S165
F5677.S168
F5677.S174
F5677.S179
F5677.S183
F5677.S189
F5677.S201
F5677.S208
F5677.S211
F5677.S219
F5677.S226
F5677.S231
F5677.S243
F5677.S246
F5677.S250
F5677.S513
F5677.S521
F5677.S524
F5677.S529
F5677.S534
F5677.S541
F5677.S544
F5677.S556
F5677.S562
F5677.S569
F5677.S576
F5677.S588
F5677.S591
F5677.S597
Samples