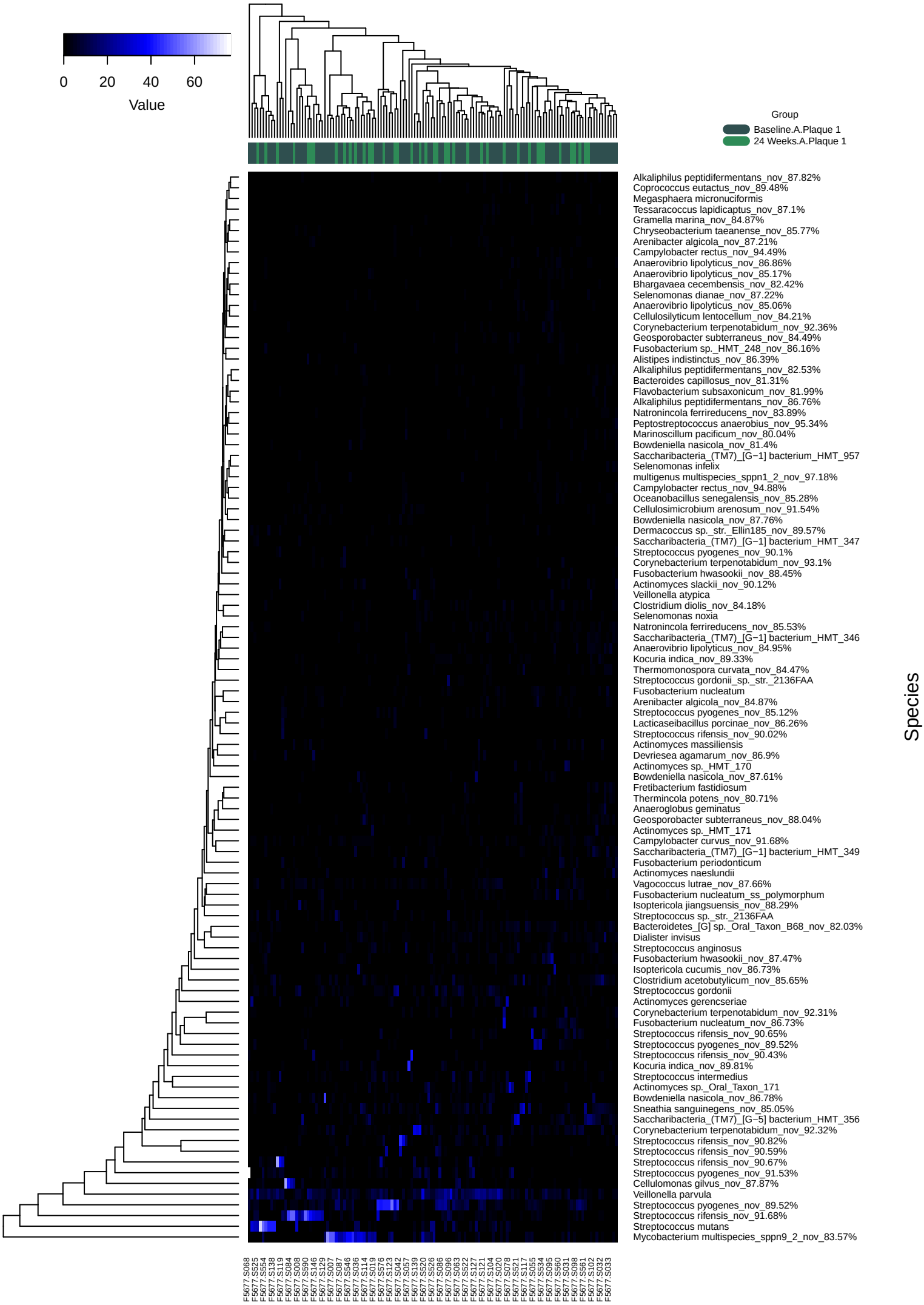
0
20
40
60
Value
Group
Baseline.A.Plaque 1
24 Weeks.A.Plaque 1
Alkaliphilus peptidifermentans_nov_87.82%
Coprococcus eutactus_nov_89.48%
Megasphaera micronuciformis
Tessaracoccus lapidicaptus_nov_87.1%
Gramella marina_nov_84.87%
Chryseobacterium taeanense_nov_85.77%
Arenibacter algicola_nov_87.21%
Campylobacter rectus_nov_94.49%
Anaerovibrio lipolyticus_nov_86.86%
Anaerovibrio lipolyticus_nov_85.17%
Bhargavaea cecembensis_nov_82.42%
Selenomonas dianae_nov_87.22%
Anaerovibrio lipolyticus_nov_85.06%
Cellulosilyticum lentocellum_nov_84.21%
Corynebacterium terpenotabidum_nov_92.36%
Geosporobacter subterraneus_nov_84.49%
Fusobacterium sp._HMT_248_nov_86.16%
Alistipes indistinctus_nov_86.39%
Alkaliphilus peptidifermentans_nov_82.53%
Bacteroides capillosus_nov_81.31%
Flavobacterium subsaxonicum_nov_81.99%
Alkaliphilus peptidifermentans_nov_86.76%
Natronincola ferrireducens_nov_83.89%
Peptostreptococcus anaerobius_nov_95.34%
Marinoscillum pacificum_nov_80.04%
Bowdeniella nasicola_nov_81.4%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_957
Selenomonas infelix
multigenus multispecies_sppn1_2_nov_97.18%
Campylobacter rectus_nov_94.88%
Oceanobacillus senegalensis_nov_85.28%
Cellulosimicrobium arenosum_nov_91.54%
Bowdeniella nasicola_nov_87.76%
Dermacoccus sp._str._Ellin185_nov_89.57%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Streptococcus pyogenes_nov_90.1%
Corynebacterium terpenotabidum_nov_93.1%
Fusobacterium hwasookii_nov_88.45%
Actinomyces slackii_nov_90.12%
Veillonella atypica
Clostridium diolis_nov_84.18%
Selenomonas noxia
Natronincola ferrireducens_nov_85.53%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Anaerovibrio lipolyticus_nov_84.95%
Kocuria indica_nov_89.33%
Thermomonospora curvata_nov_84.47%
Streptococcus gordonii_sp._str._2136FAA
Fusobacterium nucleatum
Arenibacter algicola_nov_84.87%
Streptococcus pyogenes_nov_85.12%
Lacticaseibacillus porcinae_nov_86.26%
Streptococcus rifensis_nov_90.02%
Actinomyces massiliensis
Devriesea agamarum_nov_86.9%
Actinomyces sp._HMT_170
Bowdeniella nasicola_nov_87.61%
Fretibacterium fastidiosum
Thermincola potens_nov_80.71%
Anaeroglobus geminatus
Geosporobacter subterraneus_nov_88.04%
Actinomyces sp._HMT_171
Campylobacter curvus_nov_91.68%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Fusobacterium periodonticum
Actinomyces naeslundii
Vagococcus lutrae_nov_87.66%
Fusobacterium nucleatum_ss_polymorphum
Isoptericola jiangsuensis_nov_88.29%
Streptococcus sp._str._2136FAA
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Dialister invisus
Streptococcus anginosus
Fusobacterium hwasookii_nov_87.47%
Isoptericola cucumis_nov_86.73%
Clostridium acetobutylicum_nov_85.65%
Streptococcus gordonii
Actinomyces gerencseriae
Corynebacterium terpenotabidum_nov_92.31%
Fusobacterium nucleatum_nov_86.73%
Streptococcus rifensis_nov_90.65%
Streptococcus pyogenes_nov_89.52%
Streptococcus rifensis_nov_90.43%
Kocuria indica_nov_89.81%
Streptococcus intermedius
Actinomyces sp._Oral_Taxon_171
Bowdeniella nasicola_nov_86.78%
Sneathia sanguinegens_nov_85.05%
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Corynebacterium terpenotabidum_nov_92.32%
Streptococcus rifensis_nov_90.82%
Streptococcus rifensis_nov_90.59%
Streptococcus rifensis_nov_90.67%
Streptococcus pyogenes_nov_91.53%
Cellulomonas gilvus_nov_87.87%
Veillonella parvula
Streptococcus pyogenes_nov_89.52%
Streptococcus rifensis_nov_91.68%
Streptococcus mutans
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S068
F5677.S525
F5677.S554
F5677.S138
F5677.S119
F5677.S084
F5677.S008
F5677.S590
F5677.S146
F5677.S129
F5677.S007
F5677.S087
F5677.S546
F5677.S036
F5677.S114
F5677.S019
F5677.S576
F5677.S123
F5677.S042
F5677.S057
F5677.S139
F5677.S520
F5677.S526
F5677.S086
F5677.S096
F5677.S063
F5677.S522
F5677.S127
F5677.S121
F5677.S104
F5677.S020
F5677.S078
F5677.S521
F5677.S117
F5677.S055
F5677.S534
F5677.S095
F5677.S560
F5677.S031
F5677.S098
F5677.S561
F5677.S102
F5677.S032
F5677.S033
Samples