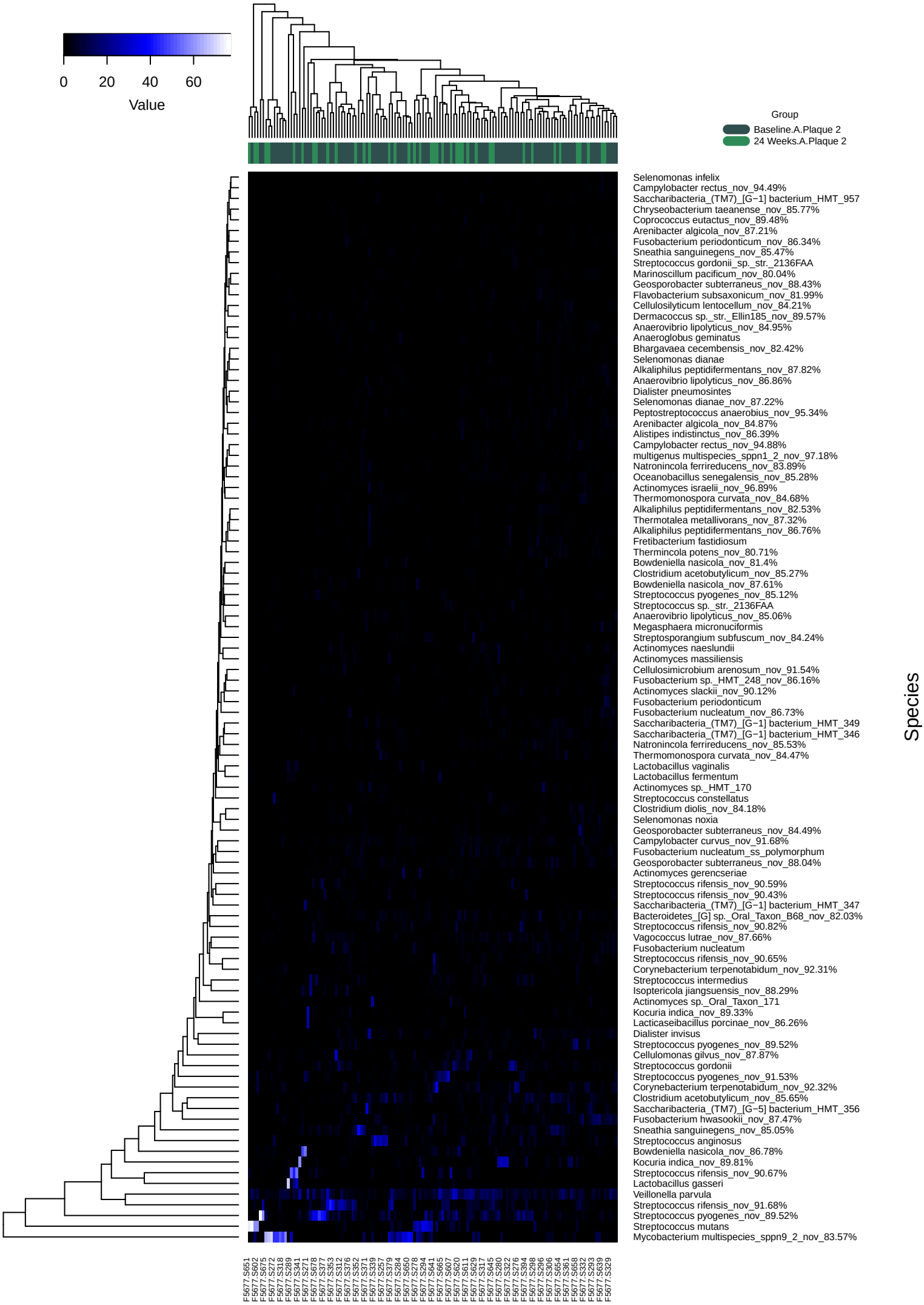
0
20
40
60
Value
Group
Baseline.A.Plaque 2
24 Weeks.A.Plaque 2
Selenomonas infelix
Campylobacter rectus_nov_94.49%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_957
Chryseobacterium taeanense_nov_85.77%
Coprococcus eutactus_nov_89.48%
Arenibacter algicola_nov_87.21%
Fusobacterium periodonticum_nov_86.34%
Sneathia sanguinegens_nov_85.47%
Streptococcus gordonii_sp._str._2136FAA
Marinoscillum pacificum_nov_80.04%
Geosporobacter subterraneus_nov_88.43%
Flavobacterium subsaxonicum_nov_81.99%
Cellulosilyticum lentocellum_nov_84.21%
Dermacoccus sp._str._Ellin185_nov_89.57%
Anaerovibrio lipolyticus_nov_84.95%
Anaeroglobus geminatus
Bhargavaea cecembensis_nov_82.42%
Selenomonas dianae
Alkaliphilus peptidifermentans_nov_87.82%
Anaerovibrio lipolyticus_nov_86.86%
Dialister pneumosintes
Selenomonas dianae_nov_87.22%
Peptostreptococcus anaerobius_nov_95.34%
Arenibacter algicola_nov_84.87%
Alistipes indistinctus_nov_86.39%
Campylobacter rectus_nov_94.88%
multigenus multispecies_sppn1_2_nov_97.18%
Natronincola ferrireducens_nov_83.89%
Oceanobacillus senegalensis_nov_85.28%
Actinomyces israelii_nov_96.89%
Thermomonospora curvata_nov_84.68%
Alkaliphilus peptidifermentans_nov_82.53%
Thermotalea metallivorans_nov_87.32%
Alkaliphilus peptidifermentans_nov_86.76%
Fretibacterium fastidiosum
Thermincola potens_nov_80.71%
Bowdeniella nasicola_nov_81.4%
Clostridium acetobutylicum_nov_85.27%
Bowdeniella nasicola_nov_87.61%
Streptococcus pyogenes_nov_85.12%
Streptococcus sp._str._2136FAA
Anaerovibrio lipolyticus_nov_85.06%
Megasphaera micronuciformis
Streptosporangium subfuscum_nov_84.24%
Actinomyces naeslundii
Actinomyces massiliensis
Cellulosimicrobium arenosum_nov_91.54%
Fusobacterium sp._HMT_248_nov_86.16%
Actinomyces slackii_nov_90.12%
Fusobacterium peridonticum
Fusobacterium nucleatum_nov_86.73%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Natronincola ferrireducens_nov_85.53%
Thermomonospora curvata_nov_84.47%
Lactobacillus vaginalis
Lactobacillus fermentum
Actinomyces sp._HMT_170
Streptococcus constellatus
Clostridium diolis_nov_84.18%
Selenomonas noxia
Geosporobacter subterraneus_nov_84.49%
Campylobacter curvus_nov_91.68%
Fusobacterium nucleatum_ss_polymorphum
Geosporobacter subterraneus_nov_88.04%
Actinomyces gerencseriae
Streptococcus rifensis_nov_90.59%
Streptococcus rifensis_nov_90.43%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Streptococcus rifensis_nov_90.82%
Vagococcus lutrae_nov_87.66%
Fusobacterium nucleatum
Streptococcus rifensis_nov_90.65%
Corynebacterium terpenotabidum_nov_92.31%
Streptococcus intermedius
Isoptericola jiangsuensis_nov_88.29%
Actinomyces sp._Oral_Taxon_171
Kocuria indica_nov_89.33%
Lacticaseibacillus porcinae_nov_86.26%
Dialister invisus
Streptococcus pyogenes_nov_89.52%
Cellulomonas gilvus_nov_87.87%
Streptococcus gordonii
Streptococcus pyogenes_nov_91.53%
Corynebacterium terpenotabidum_nov_92.32%
Clostridium acetobutylicum_nov_85.65%
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Fusobacterium hwasookii_nov_87.47%
Sneathia sanguinegens_nov_85.05%
Streptococcus anginosus
Bowdeniella nasicola_nov_86.78%
Kocuria indica_nov_89.81%
Streptococcus rifensis_nov_90.67%
Lactobacillus gasseri
Veillonella parvula
Streptococcus rifensis_nov_91.68%
Streptococcus pyogenes_nov_89.52%
Streptococcus mutans
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S651
F5677.S602
F5677.S675
F5677.S272
F5677.S318
F5677.S289
F5677.S341
F5677.S271
F5677.S678
F5677.S377
F5677.S353
F5677.S312
F5677.S376
F5677.S352
F5677.S371
F5677.S339
F5677.S257
F5677.S379
F5677.S284
F5677.S650
F5677.S278
F5677.S294
F5677.S641
F5677.S665
F5677.S607
F5677.S620
F5677.S611
F5677.S629
F5677.S317
F5677.S645
F5677.S280
F5677.S322
F5677.S276
F5677.S394
F5677.S298
F5677.S296
F5677.S306
F5677.S654
F5677.S361
F5677.S658
F5677.S332
F5677.S293
F5677.S639
F5677.S329
Samples