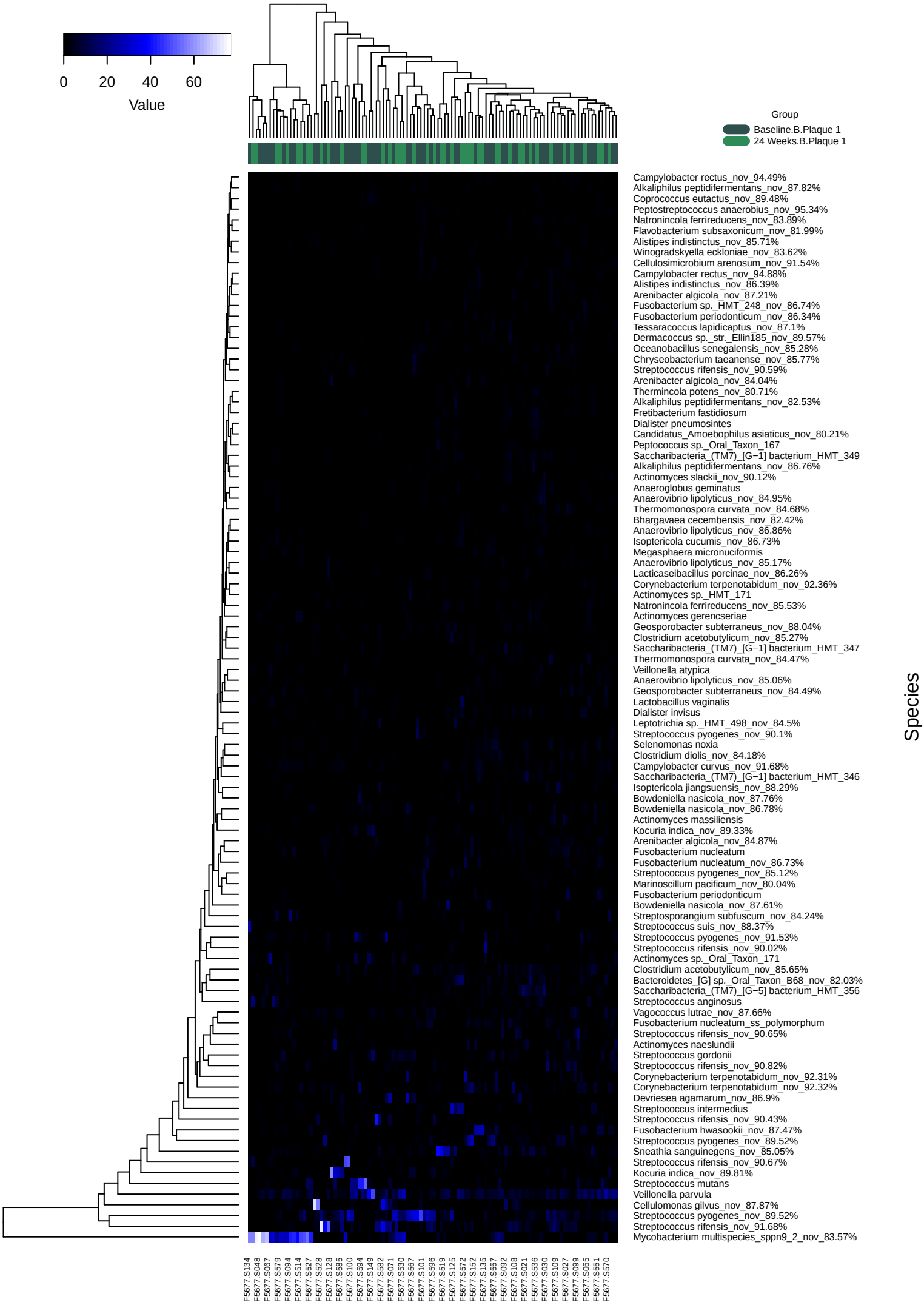
0
20
40
60
Value
Group
Baseline.B.Plaque 1
24 Weeks.B.Plaque 1
Campylobacter rectus_nov_94.49%
Alkaliphilus peptidifermentans_nov_87.82%
Coprococcus eutactus_nov_89.48%
Peptostreptococcus anaerobius_nov_95.34%
Natronincola ferrireducens_nov_83.89%
Flavobacterium subsaxonicum_nov_81.99%
Alistipes indistinctus_nov_85.71%
Winogradskyella ecklonia_nov_83.62%
Cellulosimicrobium arenosum_nov_91.54%
Campylobacter rectus_nov_94.88%
Alistipes indistinctus_nov_86.39%
Arenibacter algicola_nov_87.21%
Fusobacterium sp._HMT_248_nov_86.74%
Fusobacterium periodonticum_nov_86.34%
Tessaracoccus lapidicaptus_nov_87.1%
Dermacoccus sp._str._Ellin185_nov_89.57%
Oceanobacillus senegalensis_nov_85.28%
Chryseobacterium taeanense_nov_85.77%
Streptococcus rifensis_nov_90.59%
Arenibacter algicola_nov_84.04%
Thermincola potens_nov_80.71%
Alkaliphilus peptidifermentans_nov_82.53%
Fretibacterium fastidiosum
Dialister pneumosintes
Candidatus_Amoebophilus asiaticus_nov_80.21%
Peptococcus sp._Oral_Taxon_167
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Alkaliphilus peptidifermentans_nov_86.76%
Actinomyces slackii_nov_90.12%
Anaeroglobus geminatus
Anaerovibrio lipolyticus_nov_84.95%
Thermomonospora curvata_nov_84.68%
Bhargavaea cecembensis_nov_82.42%
Anaerovibrio lipolyticus_nov_86.86%
Isoptericola cucumis_nov_86.73%
Megasphaera micronuciformis
Anaerovibrio lipolyticus_nov_85.17%
Lacticaseibacillus porcinae_nov_86.26%
Corynebacterium terpenotabidum_nov_92.36%
Actinomyces sp._HMT_171
Natronincola ferrireducens_nov_85.53%
Actinomyces gerencseriae
Geosporobacter subterraneus_nov_88.04%
Clostridium acetobutylicum_nov_85.27%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Thermomonospora curvata_nov_84.47%
Veillonella atypica
Anaerovibrio lipolyticus_nov_85.06%
Geosporobacter subterraneus_nov_84.49%
Lactobacillus vaginalis
Dialister invisus
Leptotrichia sp._HMT_498_nov_84.5%
Streptococcus pyogenes_nov_90.1%
Selenomonas noxia
Clostridium diolis_nov_84.18%
Campylobacter curvus_nov_91.68%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Isoptericola jiangsuensis_nov_88.29%
Bowdeniella nasicola_nov_87.76%
Bowdeniella nasicola_nov_86.78%
Actinomyces massiliensis
Kocuria indica_nov_89.33%
Arenibacter algicola_nov_84.87%
Fusobacterium nucleatum
Fusobacterium nucleatum_nov_86.73%
Streptococcus pyogenes_nov_85.12%
Marinoscillum pacificum_nov_80.04%
Fusobacterium periodonticum
Bowdeniella nasicola_nov_87.61%
Streptosporangium subfuscum_nov_84.24%
Streptococcus suis_nov_88.37%
Streptococcus pyogenes_nov_91.53%
Streptococcus rifensis_nov_90.02%
Actinomyces sp._Oral_Taxon_171
Clostridium acetobutylicum_nov_85.65%
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Streptococcus anginosus
Vagococcus lutrae_nov_87.66%
Fusobacterium nucleatum_ss_polymorphum
Streptococcus rifensis_nov_90.65%
Actinomyces naeslundii
Streptococcus gordonii
Streptococcus rifensis_nov_90.82%
Corynebacterium terpenotabidum_nov_92.31%
Corynebacterium terpenotabidum_nov_92.32%
Devriesea agamarum_nov_86.9%
Streptococcus intermedius
Streptococcus rifensis_nov_90.43%
Fusobacterium hwasookii_nov_87.47%
Streptococcus pyogenes_nov_89.52%
Sneathia sanguinegens_nov_85.05%
Streptococcus rifensis_nov_90.67%
Kocuria indica_nov_89.81%
Streptococcus mutans
Veillonella parvula
Cellulomonas gilvus_nov_87.87%
Streptococcus pyogenes_nov_89.52%
Streptococcus rifensis_nov_91.68%
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S134
F5677.S048
F5677.S067
F5677.S579
F5677.S094
F5677.S514
F5677.S527
F5677.S528
F5677.S128
F5677.S585
F5677.S100
F5677.S594
F5677.S149
F5677.S582
F5677.S071
F5677.S530
F5677.S567
F5677.S101
F5677.S596
F5677.S519
F5677.S125
F5677.S572
F5677.S152
F5677.S135
F5677.S557
F5677.S092
F5677.S108
F5677.S021
F5677.S536
F5677.S030
F5677.S109
F5677.S027
F5677.S099
F5677.S065
F5677.S551
F5677.S570
Samples