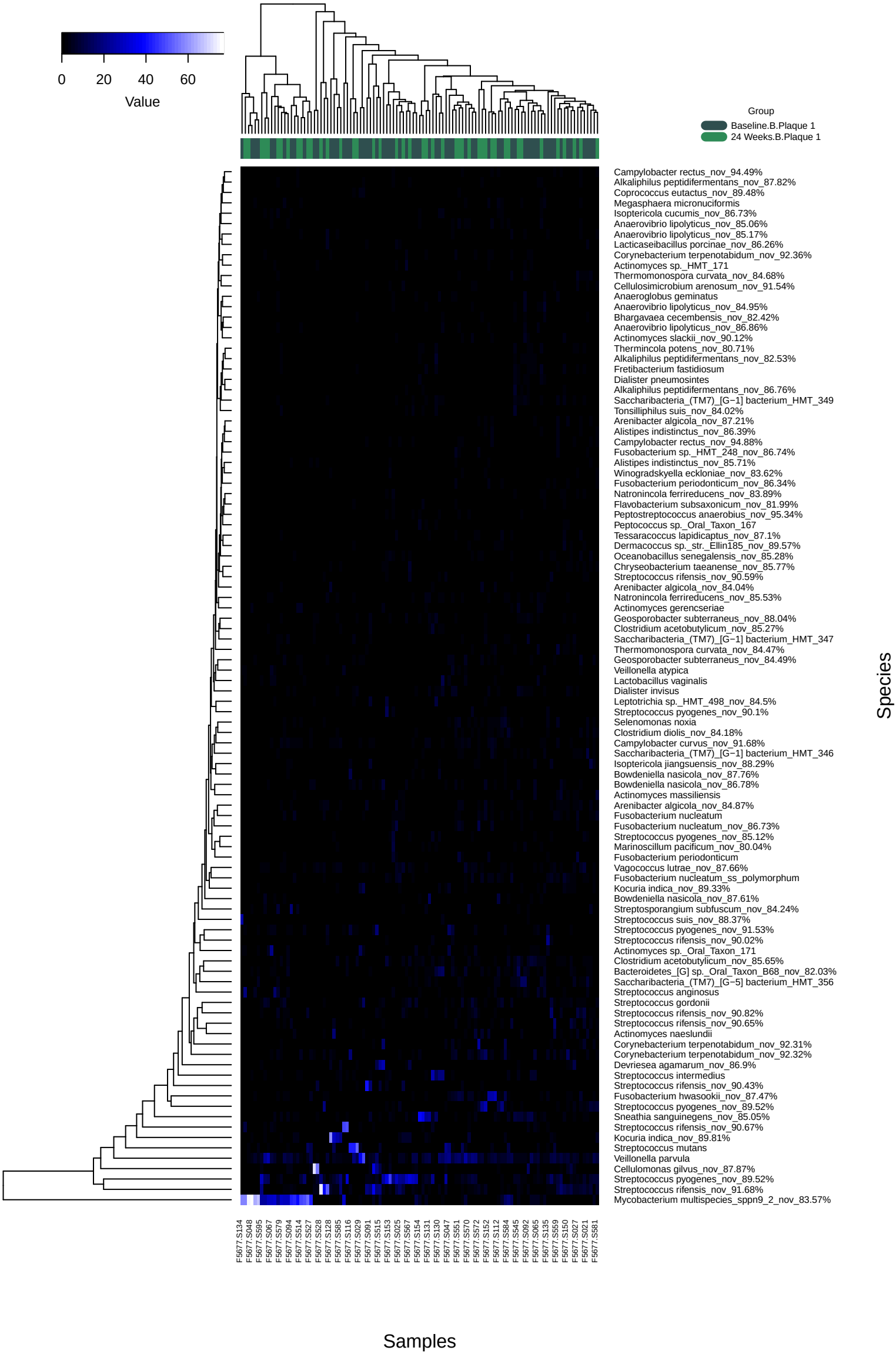
0
20
40
60
Value
Group
Baseline.B.Plaque 1
24 Weeks.B.Plaque 1
Campylobacter rectus_nov_94.49%
Alkaliphilus peptidifermentans_nov_87.82%
Coprococcus eutactus_nov_89.48%
Megasphaera micronuciformis
Isoptericola cucumis_nov_86.73%
Anaerovibrio lipolyticus_nov_85.06%
Anaerovibrio lipolyticus_nov_85.17%
Lacticaseibacillus porcinae_nov_86.26%
Corynebacterium terpenotabidum_nov_92.36%
Actinomyces sp._HMT_171
Thermomonospora curvata_nov_84.68%
Cellulosimicrobium arenosum_nov_91.54%
Anaeroglobus geminatus
Anaerovibrio lipolyticus_nov_84.95%
Bhargavaea cecembensis_nov_82.42%
Anaerovibrio lipolyticus_nov_86.86%
Actinomyces slackii_nov_90.12%
Thermincola potens_nov_80.71%
Alkaliphilus peptidifermentans_nov_82.53%
Fretibacterium fastidiosum
Dialister pneumosintes
Alkaliphilus peptidifermentans_nov_86.76%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Tonsilliphilus suis_nov_84.02%
Arenibacter algicola_nov_87.21%
Alistipes indistinctus_nov_86.39%
Campylobacter rectus_nov_94.88%
Fusobacterium sp._HMT_248_nov_86.74%
Alistipes indistinctus_nov_85.71%
Winogradskyella eckloniae_nov_83.62%
Fusobacterium periodonticum_nov_86.34%
Natronincola ferrireducens_nov_83.89%
Flavobacterium subsaxonicum_nov_81.99%
Peptostreptococcus anaerobius_nov_95.34%
Peptococcus sp._Oral_Taxon_167
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%
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%
Geosporobacter subterraneus_nov_84.49%
Veillonella atypica
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
Arenibacter algicola_nov_84.87%
Fusobacterium nucleatum
Fusobacterium nucleatum_nov_86.73%
Streptococcus pyogenes_nov_85.12%
Marinoscillum pacificum_nov_80.04%
Fusobacterium periodonticum
Vagococcus lutrae_nov_87.66%
Fusobacterium nucleatum_ss_polymorphum
Kocuria indica_nov_89.33%
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
Streptococcus gordonii
Streptococcus rifensis_nov_90.82%
Streptococcus rifensis_nov_90.65%
Actinomyces naeslundii
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.S595
F5677.S067
F5677.S579
F5677.S094
F5677.S514
F5677.S527
F5677.S528
F5677.S128
F5677.S585
F5677.S116
F5677.S029
F5677.S091
F5677.S515
F5677.S153
F5677.S025
F5677.S567
F5677.S154
F5677.S131
F5677.S130
F5677.S047
F5677.S551
F5677.S570
F5677.S572
F5677.S152
F5677.S112
F5677.S584
F5677.S545
F5677.S092
F5677.S065
F5677.S135
F5677.S559
F5677.S150
F5677.S027
F5677.S021
F5677.S581
Samples