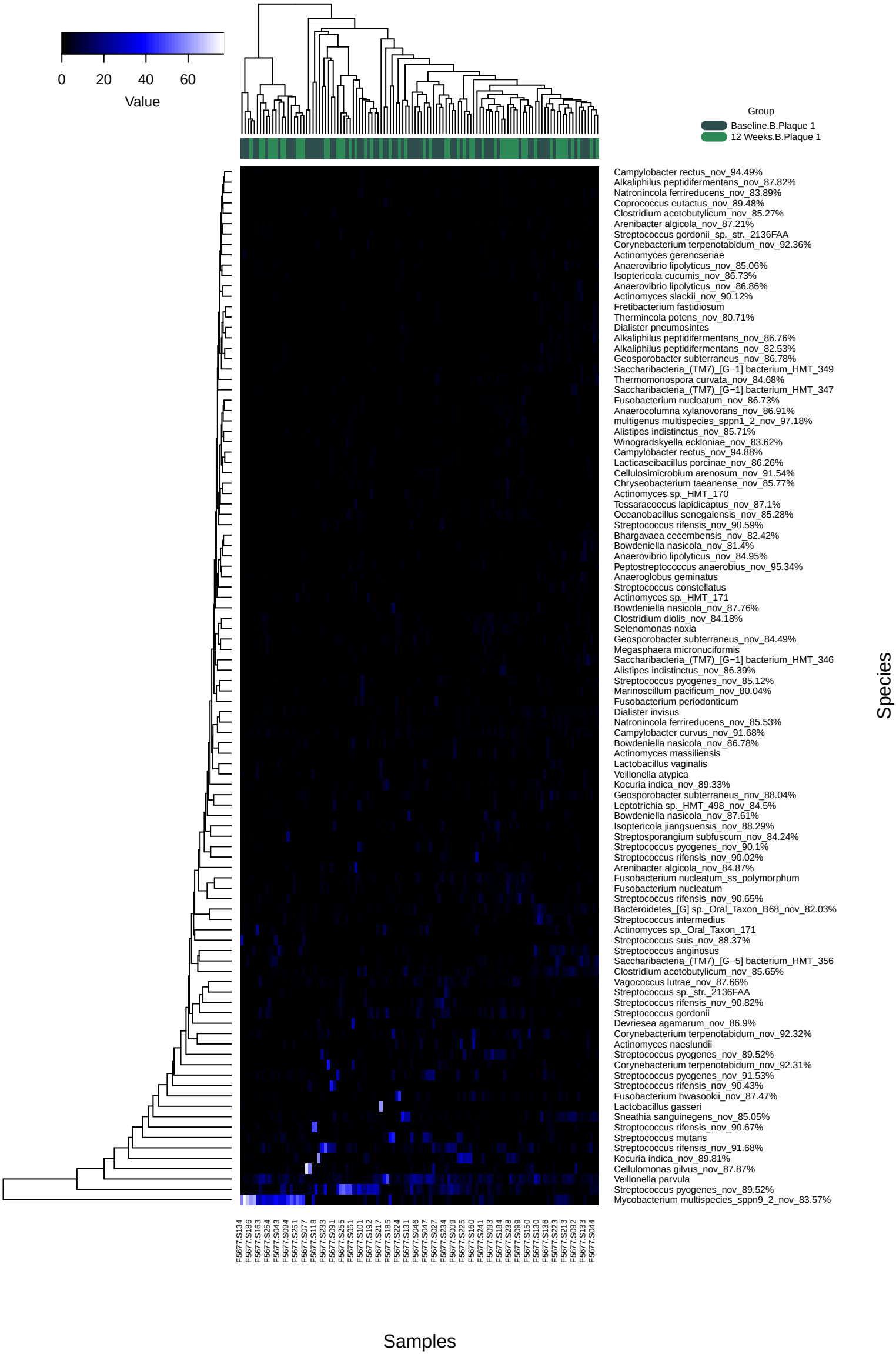
0
20
40
60
Value
Group
Baseline.B.Plaque 1
12 Weeks.B.Plaque 1
Campylobacter rectus_nov_94.49%
Alkaliphilus peptidifermentans_nov_87.82%
Natronincola ferrireducens_nov_83.89%
Coprococcus eutactus_nov_89.48%
Clostridium acetobutylicum_nov_85.27%
Arenibacter algicola_nov_87.21%
Streptococcus gordonii_sp._str._2136FAA
Corynebacterium terpenotabidum_nov_92.36%
Actinomyces gerencseriae
Anaerovibrio lipolyticus_nov_85.06%
Isoptericola cucumis_nov_86.73%
Anaerovibrio lipolyticus_nov_86.86%
Actinomyces slackii_nov_90.12%
Fretibacterium fastidiosum
Thermincola potens_nov_80.71%
Dialister pneumosintes
Alkaliphilus peptidifermentans_nov_86.76%
Alkaliphilus peptidifermentans_nov_82.53%
Geosporobacter subterraneus_nov_86.78%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Thermomonospora curvata_nov_84.68%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Fusobacterium nucleatum_nov_86.73%
Anaerocolumna xylanovorans_nov_86.91%
multigenus multispecies_sppn1_2_nov_97.18%
Alistipes indistinctus_nov_85.71%
Winogradskyella eckloniae_nov_83.62%
Campylobacter rectus_nov_94.88%
Lacticaseibacillus porcinae_nov_86.26%
Cellulosimicrobium arenosum_nov_91.54%
Chryseobacterium taeanense_nov_85.77%
Actinomyces sp._HMT_170
Tessaracoccus lapidicaptus_nov_87.1%
Oceanobacillus senegalensis_nov_85.28%
Streptococcus rifensis_nov_90.59%
Bhargavaea cecembensis_nov_82.42%
Bowdeniella nasicola_nov_81.4%
Anaerovibrio lipolyticus_nov_84.95%
Peptostreptococcus anaerobius_nov_95.34%
Anaeroglobus geminatus
Streptococcus constellatus
Actinomyces sp._HMT_171
Bowdeniella nasicola_nov_87.76%
Clostridium diolis_nov_84.18%
Selenomonas noxia
Geosporobacter subterraneus_nov_84.49%
Megasphaera micronuciformis
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Alistipes indistinctus_nov_86.39%
Streptococcus pyogenes_nov_85.12%
Marinoscillum pacificum_nov_80.04%
Fusobacterium periodonticum
Dialister invisus
Natronincola ferrireducens_nov_85.53%
Campylobacter curvus_nov_91.68%
Bowdeniella nasicola_nov_86.78%
Actinomyces massiliensis
Lactobacillus vaginalis
Veillonella atypica
Kocuria indica_nov_89.33%
Geosporobacter subterraneus_nov_88.04%
Leptotrichia sp._HMT_498_nov_84.5%
Bowdeniella nasicola_nov_87.61%
Isoptericola jiangsuensis_nov_88.29%
Streptosporangium subfuscum_nov_84.24%
Streptococcus pyogenes_nov_90.1%
Streptococcus rifensis_nov_90.02%
Arenibacter algicola_nov_84.87%
Fusobacterium nucleatum_ss_polymorphum
Fusobacterium nucleatum
Streptococcus rifensis_nov_90.65%
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Streptococcus intermedius
Actinomyces sp._Oral_Taxon_171
Streptococcus suis_nov_88.37%
Streptococcus anginosus
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Clostridium acetobutylicum_nov_85.65%
Vagococcus lutrae_nov_87.66%
Streptococcus sp._str._2136FAA
Streptococcus rifensis_nov_90.82%
Streptococcus gordonii
Devriesea agamarum_nov_86.9%
Corynebacterium terpenotabidum_nov_92.32%
Actinomyces naeslundii
Streptococcus pyogenes_nov_89.52%
Corynebacterium terpenotabidum_nov_92.31%
Streptococcus pyogenes_nov_91.53%
Streptococcus rifensis_nov_90.43%
Fusobacterium hwasookii_nov_87.47%
Lactobacillus gasseri
Sneathia sanguinegens_nov_85.05%
Streptococcus rifensis_nov_90.67%
Streptococcus mutans
Streptococcus rifensis_nov_91.68%
Kocuria indica_nov_89.81%
Cellulomonas gilvus_nov_87.87%
Veillonella parvula
Streptococcus pyogenes_nov_89.52%
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S134
F5677.S186
F5677.S163
F5677.S254
F5677.S043
F5677.S094
F5677.S251
F5677.S077
F5677.S118
F5677.S233
F5677.S091
F5677.S255
F5677.S051
F5677.S101
F5677.S192
F5677.S217
F5677.S185
F5677.S224
F5677.S131
F5677.S046
F5677.S047
F5677.S027
F5677.S234
F5677.S009
F5677.S225
F5677.S160
F5677.S241
F5677.S093
F5677.S184
F5677.S238
F5677.S099
F5677.S150
F5677.S130
F5677.S136
F5677.S223
F5677.S213
F5677.S092
F5677.S133
F5677.S044
Samples