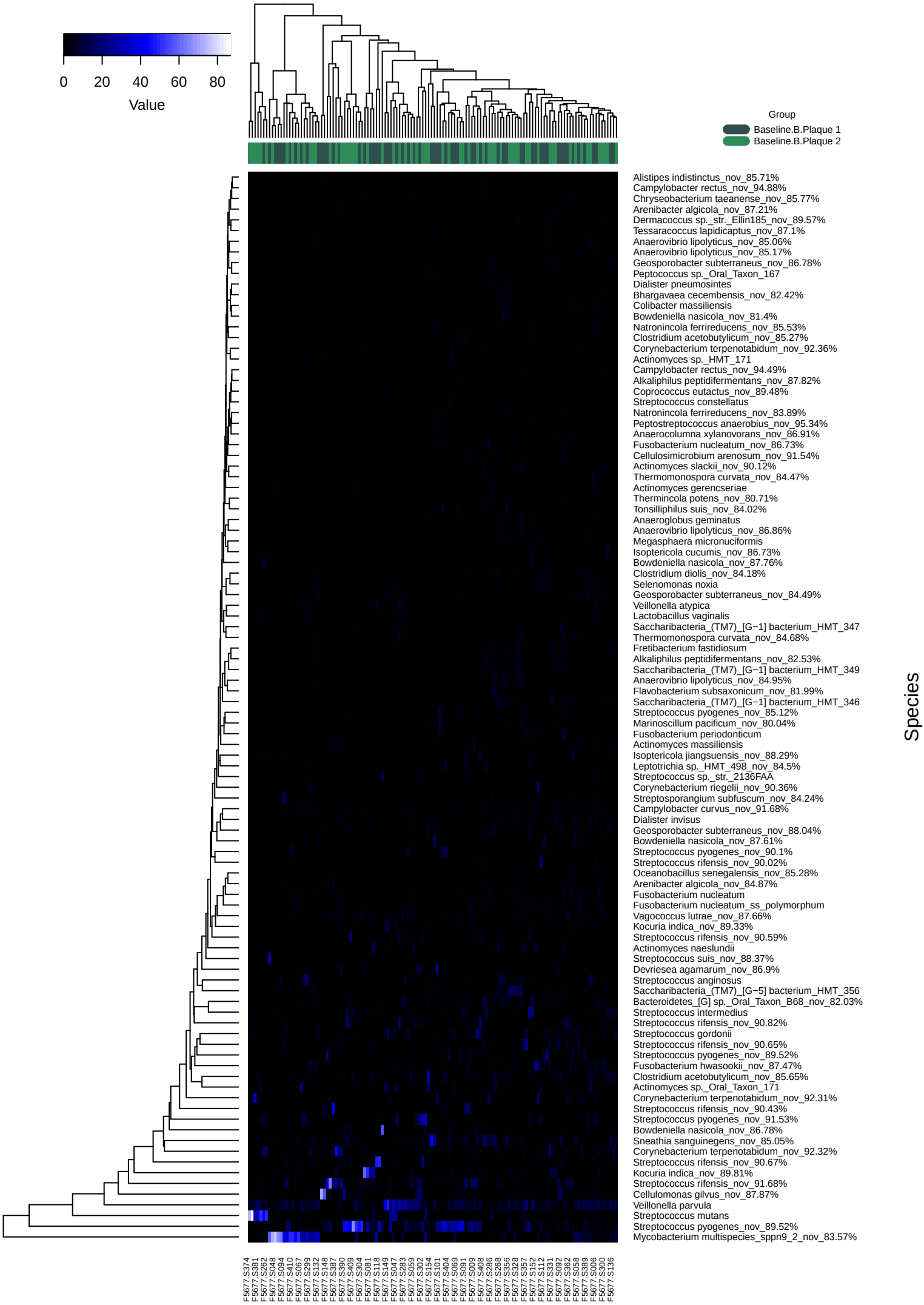
0
20
40
60
80
Value
Group
Baseline.B.Plaque 1
Baseline.B.Plaque 2
Alistipes indistinctus_nov_85.71%
Campylobacter rectus_nov_94.88%
Chryseobacterium taeanense_nov_85.77%
Arenibacter algicola_nov_87.21%
Dermacoccus sp._str._Ellin185_nov_89.57%
Tessaracoccus lapidicaptus_nov_87.1%
Anaerovibrio lipolyticus_nov_85.06%
Anaerovibrio lipolyticus_nov_85.17%
Geosporobacter subterraneus_nov_86.78%
Peptococcus sp._Oral_Taxon_167
Dialister pneumosintes
Bhargavaea cecembensis_nov_82.42%
Colibacter massiliensis
Bowdeniella nasicola_nov_81.4%
Natronincola ferrireducens_nov_85.53%
Clostridium acetobutylicum_nov_85.27%
Corynebacterium terpenotabidum_nov_92.36%
Actinomyces sp._HMT_171
Campylobacter rectus_nov_94.49%
Alkaliphilus peptidifermentans_nov_87.82%
Coprococcus eutactus_nov_89.48%
Streptococcus constellatus
Natronincola ferrireducens_nov_83.89%
Peptostreptococcus anaerobius_nov_95.34%
Anaerocolumna xylanovorans_nov_86.91%
Fusobacterium nucleatum_nov_86.73%
Cellulosimicrobium arenosum_nov_91.54%
Actinomyces slackii_nov_90.12%
Thermomonospora curvata_nov_84.47%
Actinomyces gerencseriae
Thermincola potens_nov_80.71%
Tonsilliphilus suis_nov_84.02%
Anaeroglobus geminatus
Anaerovibrio lipolyticus_nov_86.86%
Megasphaera micronuciformis
Isoptericola cucumis_nov_86.73%
Bowdeniella nasicola_nov_87.76%
Clostridium diolis_nov_84.18%
Selenomonas noxia
Geosporobacter subterraneus_nov_84.49%
Veillonella atypica
Lactobacillus vaginalis
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Thermomonospora curvata_nov_84.68%
Fretibacterium fastidiosum
Alkaliphilus peptidifermentans_nov_82.53%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Anaerovibrio lipolyticus_nov_84.95%
Flavobacterium subsaxonicum_nov_81.99%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Streptococcus pyogenes_nov_85.12%
Marinoscillum pacificum_nov_80.04%
Fusobacterium periodonticum
Actinomyces massiliensis
Isoptericola jiangsuensis_nov_88.29%
Leptotrichia sp._HMT_498_nov_84.5%
Streptococcus sp._str._2136FAA
Corynebacterium riegelii_nov_90.36%
Streptosporangium subfuscum_nov_84.24%
Campylobacter curvus_nov_91.68%
Dialister invisus
Geosporobacter subterraneus_nov_88.04%
Bowdeniella nasicola_nov_87.61%
Streptococcus pyogenes_nov_90.1%
Streptococcus rifensis_nov_90.02%
Oceanobacillus senegalensis_nov_85.28%
Arenibacter algicola_nov_84.87%
Fusobacterium nucleatum
Fusobacterium nucleatum_ss_polymorphum
Vagococcus lutrae_nov_87.66%
Kocuria indica_nov_89.33%
Streptococcus rifensis_nov_90.59%
Actinomyces naeslundii
Streptococcus suis_nov_88.37%
Devriesea agamarum_nov_86.9%
Streptococcus anginosus
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Streptococcus intermedius
Streptococcus rifensis_nov_90.82%
Streptococcus gordonii
Streptococcus rifensis_nov_90.65%
Streptococcus pyogenes_nov_89.52%
Fusobacterium hwasookii_nov_87.47%
Clostridium acetobutylicum_nov_85.65%
Actinomyces sp._Oral_Taxon_171
Corynebacterium terpenotabidum_nov_92.31%
Streptococcus rifensis_nov_90.43%
Streptococcus pyogenes_nov_91.53%
Bowdeniella nasicola_nov_86.78%
Sneathia sanguinegens_nov_85.05%
Corynebacterium terpenotabidum_nov_92.32%
Streptococcus rifensis_nov_90.67%
Kocuria indica_nov_89.81%
Streptococcus rifensis_nov_91.68%
Cellulomonas gilvus_nov_87.87%
Veillonella parvula
Streptococcus mutans
Streptococcus pyogenes_nov_89.52%
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S374
F5677.S381
F5677.S262
F5677.S048
F5677.S094
F5677.S410
F5677.S067
F5677.S299
F5677.S132
F5677.S148
F5677.S387
F5677.S390
F5677.S409
F5677.S304
F5677.S061
F5677.S118
F5677.S149
F5677.S047
F5677.S283
F5677.S059
F5677.S302
F5677.S154
F5677.S101
F5677.S404
F5677.S069
F5677.S091
F5677.S009
F5677.S408
F5677.S286
F5677.S268
F5677.S356
F5677.S328
F5677.S357
F5677.S152
F5677.S112
F5677.S331
F5677.S092
F5677.S362
F5677.S058
F5677.S389
F5677.S006
F5677.S300
F5677.S136
Samples