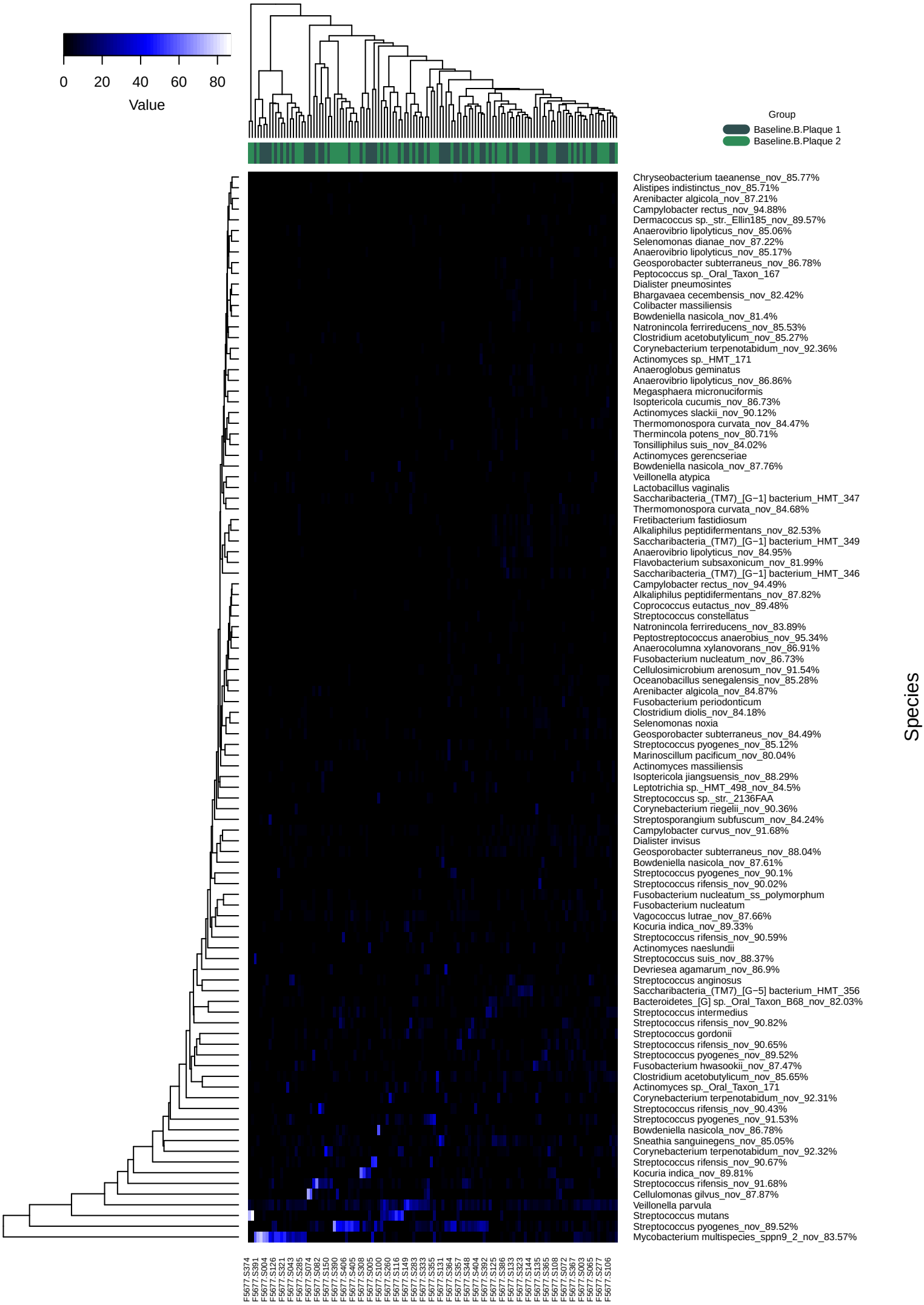
0
20
40
60
80
Value
Group
Baseline.B.Plaque 1
Baseline.B.Plaque 2
Chryseobacterium taeanense_nov_85.77%
Alistipes indistinctus_nov_85.71%
Arenibacter algicola_nov_87.21%
Campylobacter rectus_nov_94.88%
Dermacoccus sp._str._Ellin185_nov_89.57%
Anaerovibrio lipolyticus_nov_85.06%
Selenomonas dianae_nov_87.22%
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
Anaeroglobus geminatus
Anaerovibrio lipolyticus_nov_86.86%
Megasphaera micronuciformis
Isoptericola cucumis_nov_86.73%
Actinomyces slackii_nov_90.12%
Thermomonospora curvata_nov_84.47%
Thermincola potens_nov_80.71%
Tonsilliphilus suis_nov_84.02%
Actinomyces gerencseriae
Bowdeniella nasicola_nov_87.76%
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
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%
Oceanobacillus senegalensis_nov_85.28%
Arenibacter algicola_nov_84.87%
Fusobacterium periodonticum
Clostridium diolis_nov_84.18%
Selenomonas noxia
Geosporobacter subterraneus_nov_84.49%
Streptococcus pyogenes_nov_85.12%
Marinoscillum pacificum_nov_80.04%
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%
Fusobacterium nucleatum_ss_polymorphum
Fusobacterium nucleatum
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.S391
F5677.S004
F5677.S126
F5677.S321
F5677.S043
F5677.S285
F5677.S074
F5677.S082
F5677.S150
F5677.S390
F5677.S406
F5677.S405
F5677.S308
F5677.S005
F5677.S100
F5677.S260
F5677.S116
F5677.S149
F5677.S283
F5677.S333
F5677.S355
F5677.S131
F5677.S364
F5677.S357
F5677.S348
F5677.S404
F5677.S392
F5677.S125
F5677.S386
F5677.S133
F5677.S323
F5677.S144
F5677.S135
F5677.S365
F5677.S108
F5677.S072
F5677.S367
F5677.S003
F5677.S065
F5677.S277
F5677.S106
Samples