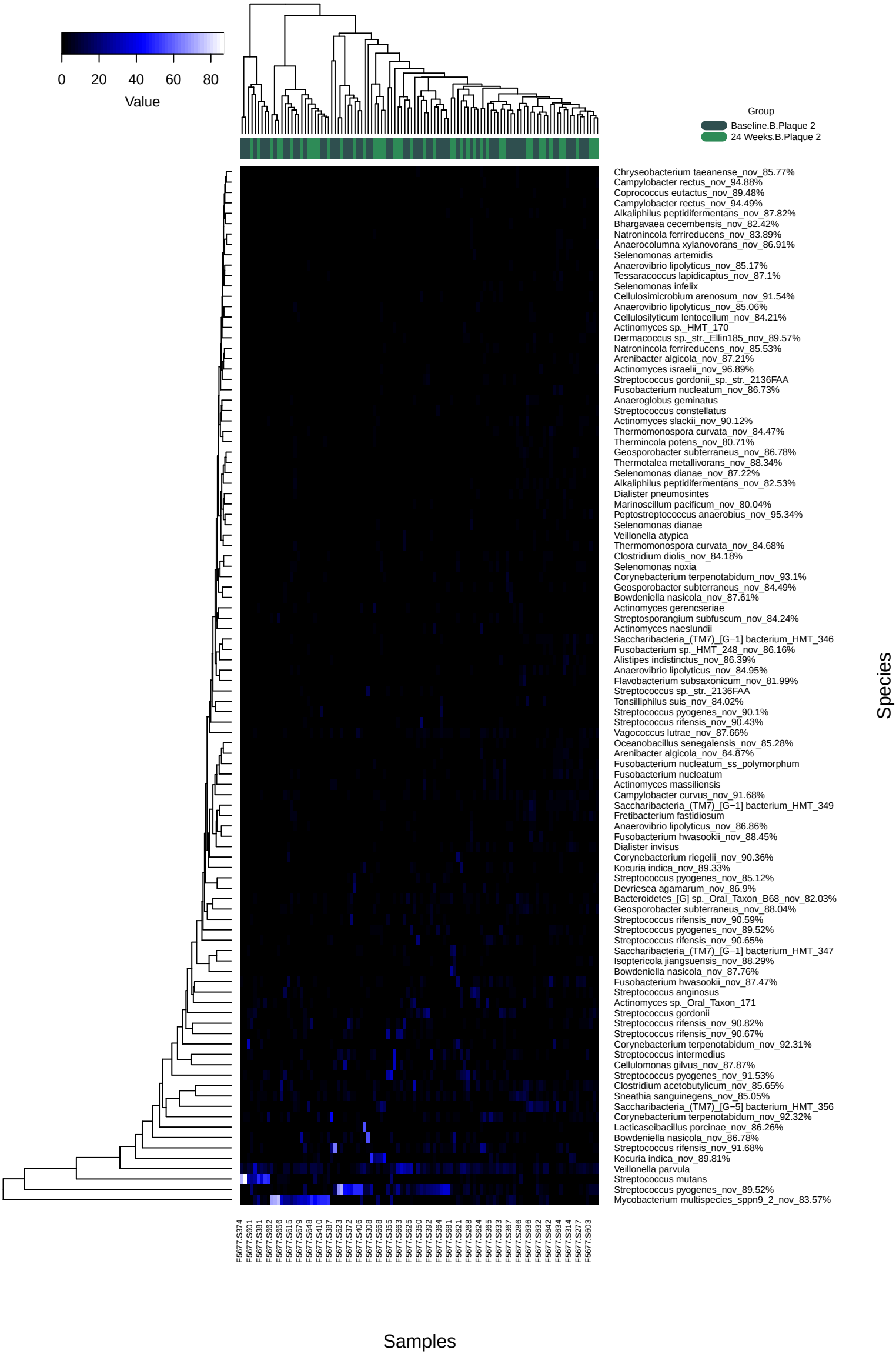
0
20
40
60
80
Value
Group
Baseline.B.Plaque 2
24 Weeks.B.Plaque 2
Chryseobacterium taeanense_nov_85.77%
Campylobacter rectus_nov_94.88%
Coprococcus eutactus_nov_89.48%
Campylobacter rectus_nov_94.49%
Alkaliphilus peptidifermentans_nov_87.82%
Bhargavaea cecembensis_nov_82.42%
Natronincola ferrireducens_nov_83.89%
Anaerocolumna xylanovorans_nov_86.91%
Selenomonas artemidis
Anaerovibrio lipolyticus_nov_85.17%
Tessaracoccus lapidicaptus_nov_87.1%
Selenomonas infelix
Cellulosimicrobium arenosum_nov_91.54%
Anaerovibrio lipolyticus_nov_85.06%
Cellulosilyticum lentocellum_nov_84.21%
Actinomyces sp._HMT_170
Dermacoccus sp._str._Ellin185_nov_89.57%
Natronincola ferrireducens_nov_85.53%
Arenibacter algicola_nov_87.21%
Actinomyces israelii_nov_96.89%
Streptococcus gordonii_sp._str._2136FAA
Fusobacterium nucleatum_nov_86.73%
Anaeroglobus geminatus
Streptococcus constellatus
Actinomyces slackii_nov_90.12%
Thermomonospora curvata_nov_84.47%
Thermincola potens_nov_80.71%
Geosporobacter subterraneus_nov_86.78%
Thermotalea metallivorans_nov_88.34%
Selenomonas dianae_nov_87.22%
Alkaliphilus peptidifermentans_nov_82.53%
Dialister pneumosintes
Marinoscillum pacificum_nov_80.04%
Peptostreptococcus anaerobius_nov_95.34%
Selenomonas dianae
Veillonella atypica
Thermomonospora curvata_nov_84.68%
Clostridium diolis_nov_84.18%
Selenomonas noxia
Corynebacterium terpenotabidum_nov_93.1%
Geosporobacter subterraneus_nov_84.49%
Bowdeniella nasicola_nov_87.61%
Actinomyces gerencseriae
Streptosporangium subfuscum_nov_84.24%
Actinomyces naeslundii
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Fusobacterium sp._HMT_248_nov_86.16%
Alistipes indistinctus_nov_86.39%
Anaerovibrio lipolyticus_nov_84.95%
Flavobacterium subsaxonicum_nov_81.99%
Streptococcus sp._str._2136FAA
Tonsilliphilus suis_nov_84.02%
Streptococcus pyogenes_nov_90.1%
Streptococcus rifensis_nov_90.43%
Vagococcus lutrae_nov_87.66%
Oceanobacillus senegalensis_nov_85.28%
Arenibacter algicola_nov_84.87%
Fusobacterium nucleatum_ss_polymorphum
Fusobacterium nucleatum
Actinomyces massiliensis
Campylobacter curvus_nov_91.68%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Fretibacterium fastidiosum
Anaerovibrio lipolyticus_nov_86.86%
Fusobacterium hwasookii_nov_88.45%
Dialister invisus
Corynebacterium riegelii_nov_90.36%
Kocuria indica_nov_89.33%
Streptococcus pyogenes_nov_85.12%
Devriesea agamarum_nov_86.9%
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Geosporobacter subterraneus_nov_88.04%
Streptococcus rifensis_nov_90.59%
Streptococcus pyogenes_nov_89.52%
Streptococcus rifensis_nov_90.65%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Isoptericola jiangsuensis_nov_88.29%
Bowdeniella nasicola_nov_87.76%
Fusobacterium hwasookii_nov_87.47%
Streptococcus anginosus
Actinomyces sp._Oral_Taxon_171
Streptococcus gordonii
Streptococcus rifensis_nov_90.82%
Streptococcus rifensis_nov_90.67%
Corynebacterium terpenotabidum_nov_92.31%
Streptococcus intermedius
Cellulomonas gilvus_nov_87.87%
Streptococcus pyogenes_nov_91.53%
Clostridium acetobutylicum_nov_85.65%
Sneathia sanguinegens_nov_85.05%
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Corynebacterium terpenotabidum_nov_92.32%
Lacticaseibacillus porcinae_nov_86.26%
Bowdeniella nasicola_nov_86.78%
Streptococcus rifensis_nov_91.68%
Kocuria indica_nov_89.81%
Veillonella parvula
Streptococcus mutans
Streptococcus pyogenes_nov_89.52%
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S374
F5677.S601
F5677.S381
F5677.S662
F5677.S656
F5677.S615
F5677.S679
F5677.S648
F5677.S410
F5677.S387
F5677.S623
F5677.S372
F5677.S406
F5677.S308
F5677.S668
F5677.S355
F5677.S663
F5677.S625
F5677.S350
F5677.S392
F5677.S364
F5677.S681
F5677.S621
F5677.S268
F5677.S624
F5677.S365
F5677.S633
F5677.S367
F5677.S286
F5677.S636
F5677.S632
F5677.S642
F5677.S634
F5677.S314
F5677.S277
F5677.S603
Samples