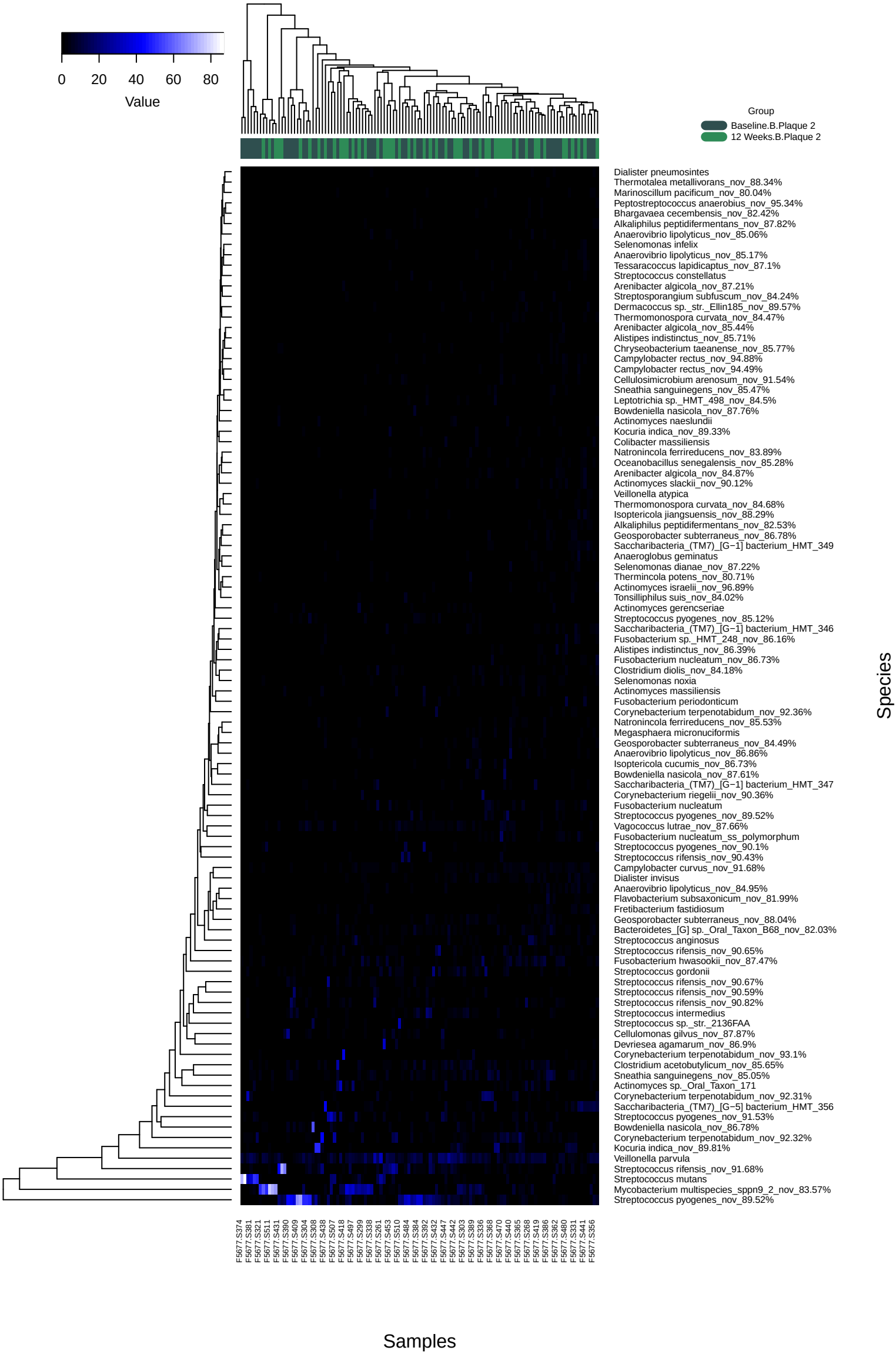
0
20
40
60
80
Value
Group
Baseline.B.Plaque 2
12 Weeks.B.Plaque 2
Dialister pneumosintes
Thermotalea metallivorans_nov_88.34%
Marinoscillum pacificum_nov_80.04%
Peptostreptococcus anaerobius_nov_95.34%
Bhargavaea cecembensis_nov_82.42%
Alkaliphilus peptidifermentans_nov_87.82%
Anaerovibrio lipolyticus_nov_85.06%
Selenomonas infelix
Anaerovibrio lipolyticus_nov_85.17%
Tessaracoccus lapidicaptus_nov_87.1%
Streptococcus constellatus
Arenibacter algicola_nov_87.21%
Streptosporangium subfuscum_nov_84.24%
Dermacoccus sp._str._Ellin185_nov_89.57%
Thermomonospora curvata_nov_84.47%
Arenibacter algicola_nov_85.44%
Alistipes indistinctus_nov_85.71%
Chryseobacterium taeanense_nov_85.77%
Campylobacter rectus_nov_94.88%
Campylobacter rectus_nov_94.49%
Cellulosimicrobium arenosum_nov_91.54%
Sneathia sanguinegens_nov_85.47%
Leptotrichia sp._HMT_498_nov_84.5%
Bowdeniella nasicola_nov_87.76%
Actinomyces naeslundii
Kocuria indica_nov_89.33%
Colibacter massiliensis
Natronincola ferrireducens_nov_83.89%
Oceanobacillus senegalensis_nov_85.28%
Arenibacter algicola_nov_84.87%
Actinomyces slackii_nov_90.12%
Veillonella atypica
Thermomonospora curvata_nov_84.68%
Isoptericola jiangsuensis_nov_88.29%
Alkaliphilus peptidifermentans_nov_82.53%
Geosporobacter subterraneus_nov_86.78%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Anaeroglobus geminatus
Selenomonas dianae_nov_87.22%
Thermincola potens_nov_80.71%
Actinomyces israelii_nov_96.89%
Tonsilliphilus suis_nov_84.02%
Actinomyces gerencseriae
Streptococcus pyogenes_nov_85.12%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Fusobacterium sp._HMT_248_nov_86.16%
Alistipes indistinctus_nov_86.39%
Fusobacterium nucleatum_nov_86.73%
Clostridium diolis_nov_84.18%
Selenomonas noxia
Actinomyces massiliensis
Fusobacterium periodonticum
Corynebacterium terpenotabidum_nov_92.36%
Natronincola ferrireducens_nov_85.53%
Megasphaera micronuciformis
Geosporobacter subterraneus_nov_84.49%
Anaerovibrio lipolyticus_nov_86.86%
Isoptericola cucumis_nov_86.73%
Bowdeniella nasicola_nov_87.61%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Corynebacterium riegelii_nov_90.36%
Fusobacterium nucleatum
Streptococcus pyogenes_nov_89.52%
Vagococcus lutrae_nov_87.66%
Fusobacterium nucleatum_ss_polymorphum
Streptococcus pyogenes_nov_90.1%
Streptococcus rifensis_nov_90.43%
Campylobacter curvus_nov_91.68%
Dialister invisus
Anaerovibrio lipolyticus_nov_84.95%
Flavobacterium subsaxonicum_nov_81.99%
Fretibacterium fastidiosum
Geosporobacter subterraneus_nov_88.04%
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Streptococcus anginosus
Streptococcus rifensis_nov_90.65%
Fusobacterium hwasookii_nov_87.47%
Streptococcus gordonii
Streptococcus rifensis_nov_90.67%
Streptococcus rifensis_nov_90.59%
Streptococcus rifensis_nov_90.82%
Streptococcus intermedius
Streptococcus sp._str._2136FAA
Cellulomonas gilvus_nov_87.87%
Devriesea agamarum_nov_86.9%
Corynebacterium terpenotabidum_nov_93.1%
Clostridium acetobutylicum_nov_85.65%
Sneathia sanguinegens_nov_85.05%
Actinomyces sp._Oral_Taxon_171
Corynebacterium terpenotabidum_nov_92.31%
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Streptococcus pyogenes_nov_91.53%
Bowdeniella nasicola_nov_86.78%
Corynebacterium terpenotabidum_nov_92.32%
Kocuria indica_nov_89.81%
Veillonella parvula
Streptococcus rifensis_nov_91.68%
Streptococcus mutans
Mycobacterium multispecies_sppn9_2_nov_83.57%
Streptococcus pyogenes_nov_89.52%
Species
F5677.S374
F5677.S381
F5677.S321
F5677.S511
F5677.S431
F5677.S390
F5677.S409
F5677.S304
F5677.S308
F5677.S438
F5677.S507
F5677.S418
F5677.S497
F5677.S299
F5677.S338
F5677.S261
F5677.S453
F5677.S510
F5677.S484
F5677.S384
F5677.S392
F5677.S432
F5677.S447
F5677.S442
F5677.S303
F5677.S389
F5677.S336
F5677.S368
F5677.S470
F5677.S440
F5677.S365
F5677.S268
F5677.S419
F5677.S386
F5677.S362
F5677.S480
F5677.S331
F5677.S441
F5677.S356
Samples