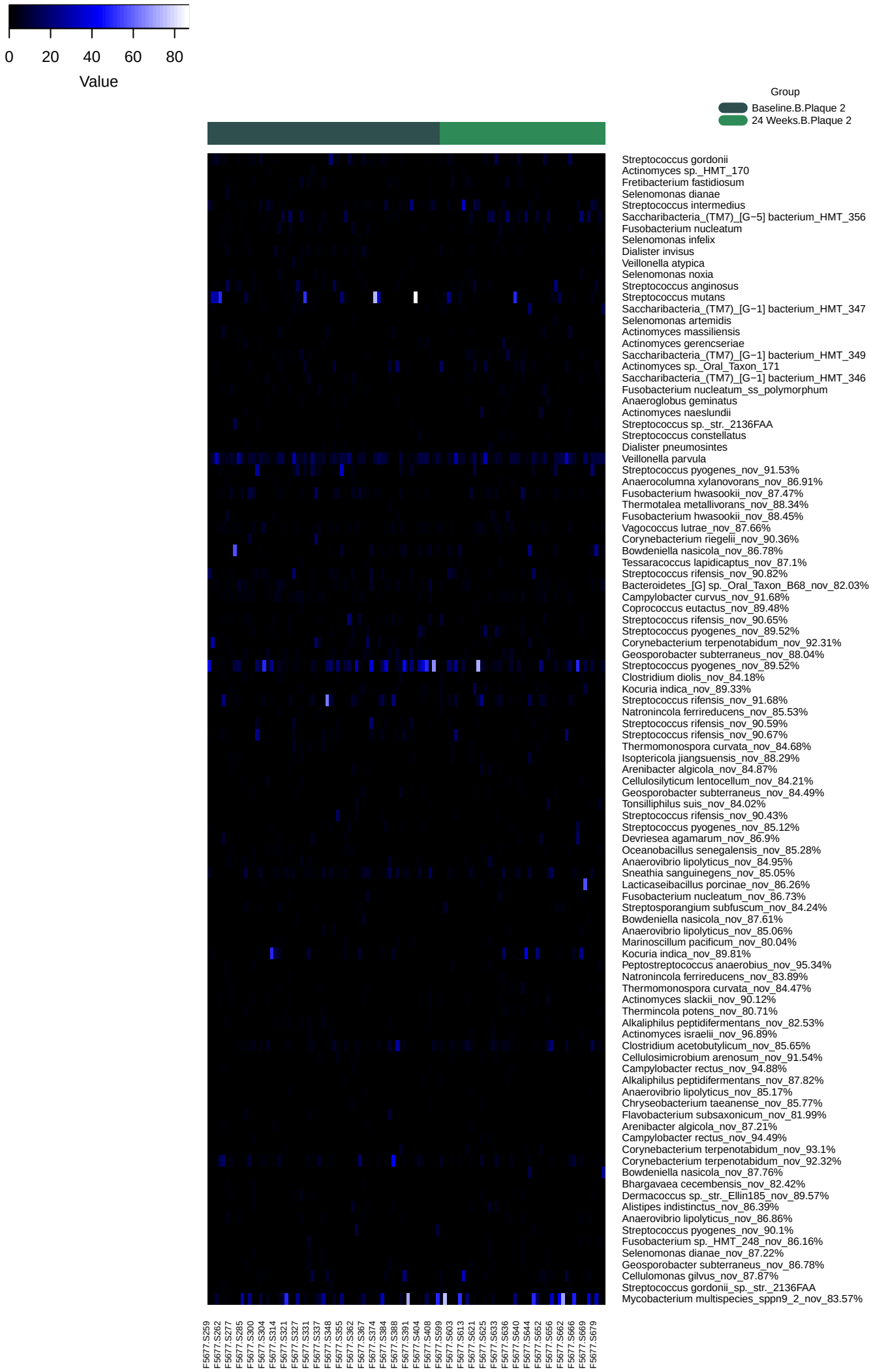
0
20
40
60
80
Value
Group
Baseline.B.Plaque 2
24 Weeks.B.Plaque 2
Streptococcus gordonii
Actinomyces sp._HMT_170
Fretibacterium fastidiosum
Selenomonas dianae
Streptococcus intermedius
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Fusobacterium nucleatum
Selenomonas infelix
Dialister invisus
Veillonella atypica
Selenomonas noxia
Streptococcus anginosus
Streptococcus mutans
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Selenomonas artemidis
Actinomyces massiliensis
Actinomyces gerencseriae
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Actinomyces sp._Oral_Taxon_171
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Fusobacterium nucleatum_ss_polymorphum
Anaeroglobus geminatus
Actinomyces naeslundii
Streptococcus sp._str._2136FAA
Streptococcus constellatus
Dialister pneumosintes
Veillonella parvula
Streptococcus pyogenes_nov_91.53%
Anaerocolumna xylanovorans_nov_86.91%
Fusobacterium hwasookii_nov_87.47%
Thermotalea metallivorans_nov_88.34%
Fusobacterium hwasookii_nov_88.45%
Vagococcus lutrae_nov_87.66%
Corynebacterium riegelii_nov_90.36%
Bowdeniella nasicola_nov_86.78%
Tessaracoccus lapidicaptus_nov_87.1%
Streptococcus rifensis_nov_90.82%
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Campylobacter curvus_nov_91.68%
Coprococcus eutactus_nov_89.48%
Streptococcus rifensis_nov_90.65%
Streptococcus pyogenes_nov_89.52%
Corynebacterium terpenotabidum_nov_92.31%
Geosporobacter subterraneus_nov_88.04%
Streptococcus pyogenes_nov_89.52%
Clostridium diolis_nov_84.18%
Kocuria indica_nov_89.33%
Streptococcus rifensis_nov_91.68%
Natronincola ferrireducens_nov_85.53%
Streptococcus rifensis_nov_90.59%
Streptococcus rifensis_nov_90.67%
Thermomonospora curvata_nov_84.68%
Isoptericola jiangsuensis_nov_88.29%
Arenibacter algicola_nov_84.87%
Cellulosilyticum lentocellum_nov_84.21%
Geosporobacter subterraneus_nov_84.49%
Tonsilliphilus suis_nov_84.02%
Streptococcus rifensis_nov_90.43%
Streptococcus pyogenes_nov_85.12%
Devriesea agamarum_nov_86.9%
Oceanobacillus senegalensis_nov_85.28%
Anaerovibrio lipolyticus_nov_84.95%
Sneathia sanguinegens_nov_85.05%
Lacticaseibacillus porcinae_nov_86.26%
Fusobacterium nucleatum_nov_86.73%
Streptosporangium subfuscum_nov_84.24%
Bowdeniella nasicola_nov_87.61%
Anaerovibrio lipolyticus_nov_85.06%
Marinoscillum pacificum_nov_80.04%
Kocuria indica_nov_89.81%
Peptostreptococcus anaerobius_nov_95.34%
Natronincola ferrireducens_nov_83.89%
Thermomonospora curvata_nov_84.47%
Actinomyces slackii_nov_90.12%
Thermincola potens_nov_80.71%
Alkaliphilus peptidifermentans_nov_82.53%
Actinomyces israelii_nov_96.89%
Clostridium acetobutylicum_nov_85.65%
Cellulosimicrobium arenosum_nov_91.54%
Campylobacter rectus_nov_94.88%
Alkaliphilus peptidifermentans_nov_87.82%
Anaerovibrio lipolyticus_nov_85.17%
Chryseobacterium taeanense_nov_85.77%
Flavobacterium subsaxonicum_nov_81.99%
Arenibacter algicola_nov_87.21%
Campylobacter rectus_nov_94.49%
Corynebacterium terpenotabidum_nov_93.1%
Corynebacterium terpenotabidum_nov_92.32%
Bowdeniella nasicola_nov_87.76%
Bhargavaea cecembensis_nov_82.42%
Dermacoccus sp._str._Ellin185_nov_89.57%
Alistipes indistinctus_nov_86.39%
Anaerovibrio lipolyticus_nov_86.86%
Streptococcus pyogenes_nov_90.1%
Fusobacterium sp._HMT_248_nov_86.16%
Selenomonas dianae_nov_87.22%
Geosporobacter subterraneus_nov_86.78%
Cellulomonas gilvus_nov_87.87%
Streptococcus gordonii_sp._str._2136FAA
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S259
F5677.S262
F5677.S277
F5677.S285
F5677.S300
F5677.S304
F5677.S314
F5677.S321
F5677.S327
F5677.S331
F5677.S337
F5677.S348
F5677.S355
F5677.S362
F5677.S367
F5677.S374
F5677.S384
F5677.S388
F5677.S391
F5677.S404
F5677.S408
F5677.S599
F5677.S603
F5677.S613
F5677.S621
F5677.S625
F5677.S633
F5677.S636
F5677.S640
F5677.S644
F5677.S652
F5677.S656
F5677.S662
F5677.S666
F5677.S669
F5677.S679
Samples