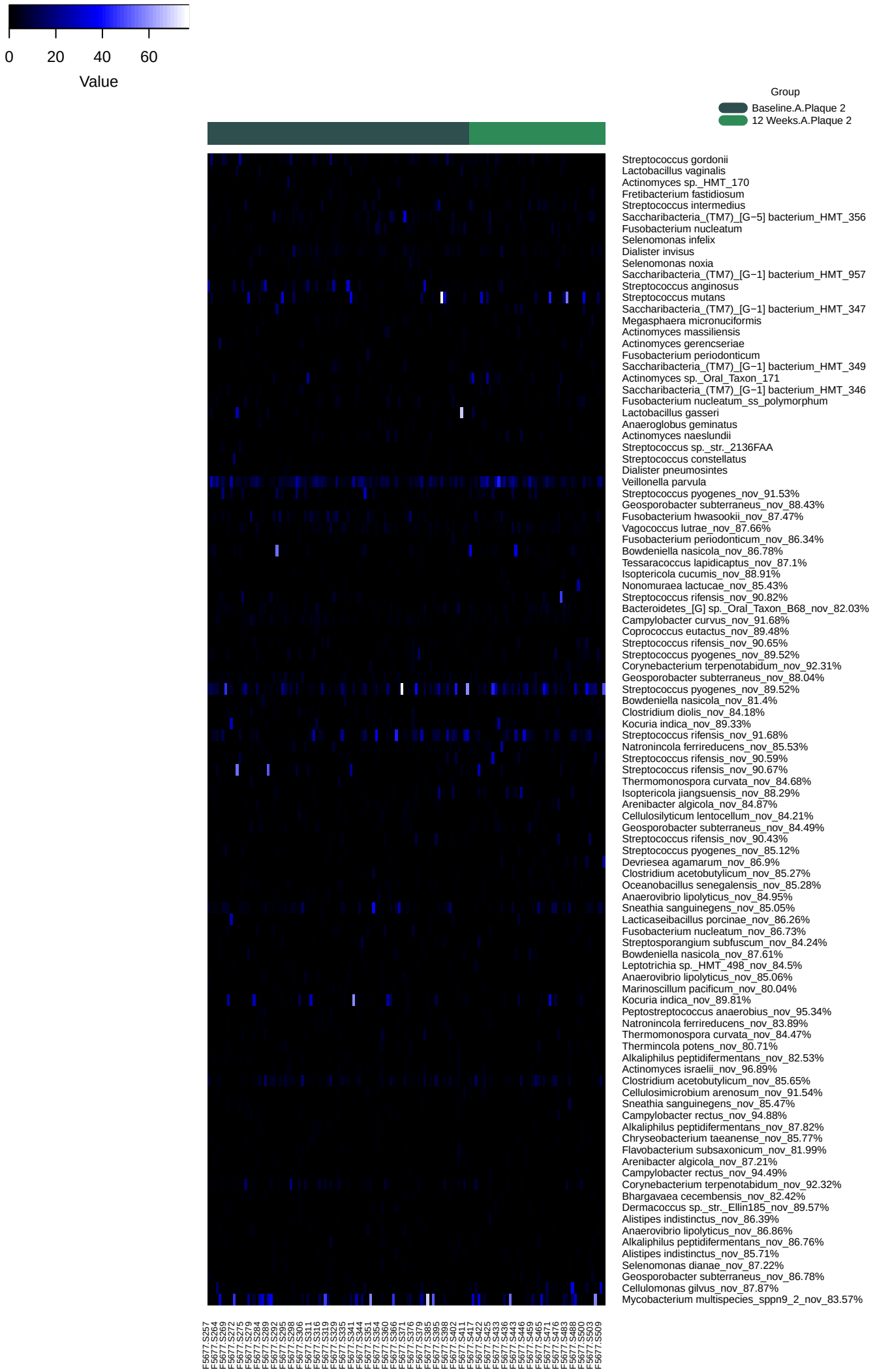
0
20
40
60
Value
Group
Baseline.A.Plaque 2
12 Weeks.A.Plaque 2
Streptococcus gordonii
Lactobacillus vaginalis
Actinomyces sp._HMT_170
Fretibacterium fastidiosum
Streptococcus intermedius
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Fusobacterium nucleatum
Selenomonas infelix
Dialister invisus
Selenomonas noxia
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_957
Streptococcus anginosus
Streptococcus mutans
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Megasphaera micronuciformis
Actinomyces massiliensis
Actinomyces gerencseriae
Fusobacterium periodonticum
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Actinomyces sp._Oral_Taxon_171
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Fusobacterium nucleatum_ss_polymorphum
Lactobacillus gasseri
Anaeroglobus geminatus
Actinomyces naeslundii
Streptococcus sp._str._2136FAA
Streptococcus constellatus
Dialister pneumosintes
Veillonella parvula
Streptococcus pyogenes_nov_91.53%
Geosporobacter subterraneus_nov_88.43%
Fusobacterium hwasookii_nov_87.47%
Vagococcus lutrae_nov_87.66%
Fusobacterium periodonticum_nov_86.34%
Bowdeniella nasicola_nov_86.78%
Tessaracoccus lapidicaptus_nov_87.1%
Isoptericola cucumis_nov_88.91%
Nonomuraea lactucae_nov_85.43%
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%
Bowdeniella nasicola_nov_81.4%
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%
Streptococcus rifensis_nov_90.43%
Streptococcus pyogenes_nov_85.12%
Devriesea agamarum_nov_86.9%
Clostridium acetobutylicum_nov_85.27%
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%
Leptotrichia sp._HMT_498_nov_84.5%
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%
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%
Sneathia sanguinegens_nov_85.47%
Campylobacter rectus_nov_94.88%
Alkaliphilus peptidifermentans_nov_87.82%
Chryseobacterium taeanense_nov_85.77%
Flavobacterium subsaxonicum_nov_81.99%
Arenibacter algicola_nov_87.21%
Campylobacter rectus_nov_94.49%
Corynebacterium terpenotabidum_nov_92.32%
Bhargavaea cecembensis_nov_82.42%
Dermacoccus sp._str._Ellin185_nov_89.57%
Alistipes indistinctus_nov_86.39%
Anaerovibrio lipolyticus_nov_86.86%
Alkaliphilus peptidifermentans_nov_86.76%
Alistipes indistinctus_nov_85.71%
Selenomonas dianae_nov_87.22%
Geosporobacter subterraneus_nov_86.78%
Cellulomonas gilvus_nov_87.87%
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S257
F5677.S264
F5677.S269
F5677.S272
F5677.S275
F5677.S279
F5677.S284
F5677.S289
F5677.S292
F5677.S295
F5677.S298
F5677.S306
F5677.S311
F5677.S316
F5677.S319
F5677.S329
F5677.S335
F5677.S341
F5677.S344
F5677.S351
F5677.S354
F5677.S360
F5677.S366
F5677.S371
F5677.S376
F5677.S379
F5677.S385
F5677.S395
F5677.S398
F5677.S402
F5677.S411
F5677.S417
F5677.S422
F5677.S425
F5677.S433
F5677.S436
F5677.S443
F5677.S446
F5677.S459
F5677.S465
F5677.S471
F5677.S476
F5677.S483
F5677.S488
F5677.S500
F5677.S503
F5677.S509
Samples