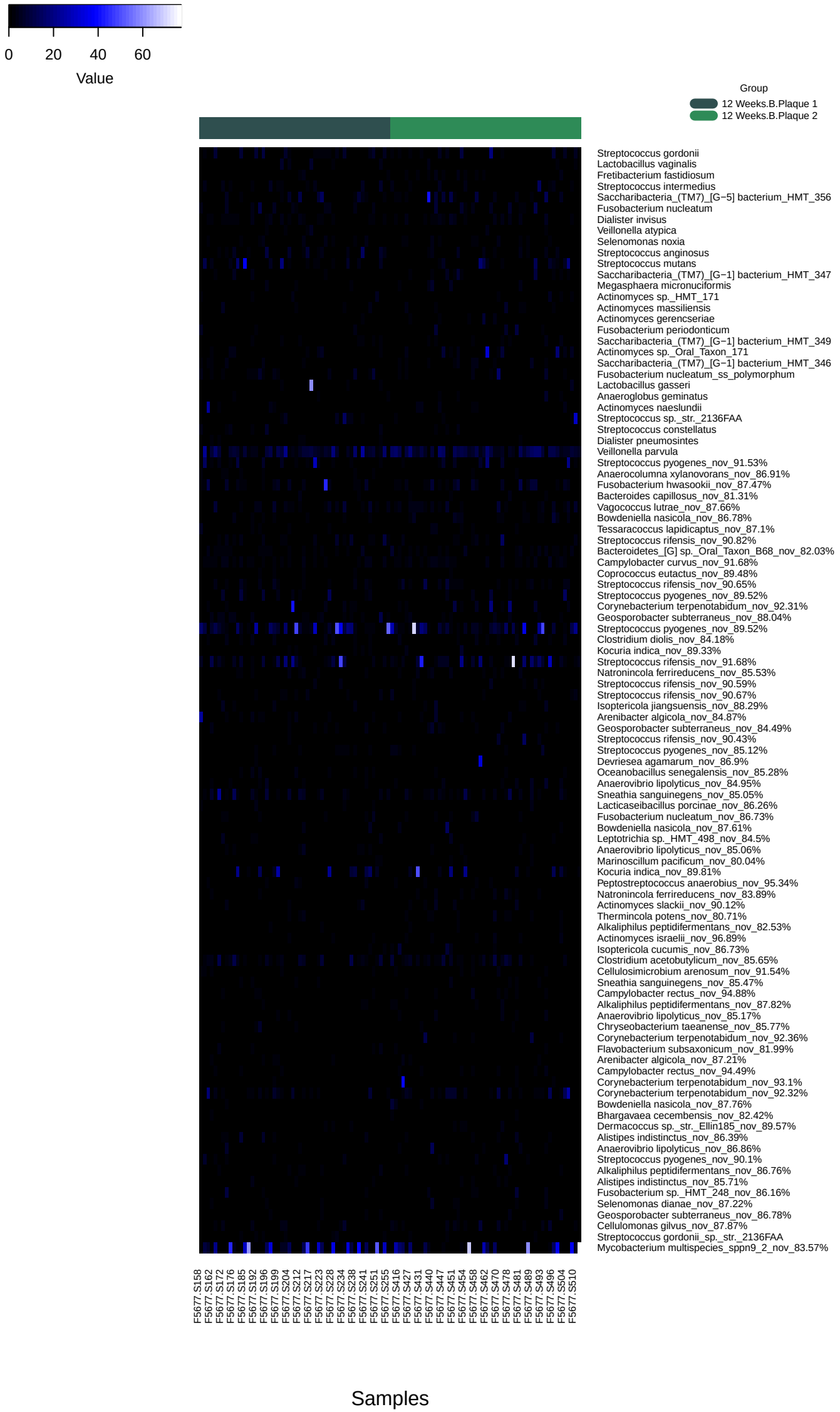
0
20
40
60
Value
Group
12 Weeks.B.Plaque 1
12 Weeks.B.Plaque 2
Streptococcus gordonii
Lactobacillus vaginalis
Fretibacterium fastidiosum
Streptococcus intermedius
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Fusobacterium nucleatum
Dialister invisus
Veillonella atypica
Selenomonas noxia
Streptococcus anginosus
Streptococcus mutans
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Megasphaera micronuciformis
Actinomyces sp._HMT_171
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%
Anaerocolumna xylanovorans_nov_86.91%
Fusobacterium hwasookii_nov_87.47%
Bacteroides capillosus_nov_81.31%
Vagococcus lutrae_nov_87.66%
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%
Isoptericola jiangsuensis_nov_88.29%
Arenibacter algicola_nov_84.87%
Geosporobacter subterraneus_nov_84.49%
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%
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%
Actinomyces slackii_nov_90.12%
Thermincola potens_nov_80.71%
Alkaliphilus peptidifermentans_nov_82.53%
Actinomyces israelii_nov_96.89%
Isoptericola cucumis_nov_86.73%
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%
Anaerovibrio lipolyticus_nov_85.17%
Chryseobacterium taeanense_nov_85.77%
Corynebacterium terpenotabidum_nov_92.36%
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%
Alkaliphilus peptidifermentans_nov_86.76%
Alistipes indistinctus_nov_85.71%
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.S158
F5677.S162
F5677.S172
F5677.S176
F5677.S185
F5677.S192
F5677.S196
F5677.S199
F5677.S204
F5677.S212
F5677.S217
F5677.S223
F5677.S228
F5677.S234
F5677.S238
F5677.S241
F5677.S251
F5677.S255
F5677.S416
F5677.S427
F5677.S431
F5677.S440
F5677.S447
F5677.S451
F5677.S454
F5677.S458
F5677.S462
F5677.S470
F5677.S478
F5677.S481
F5677.S489
F5677.S493
F5677.S496
F5677.S504
F5677.S510
Samples