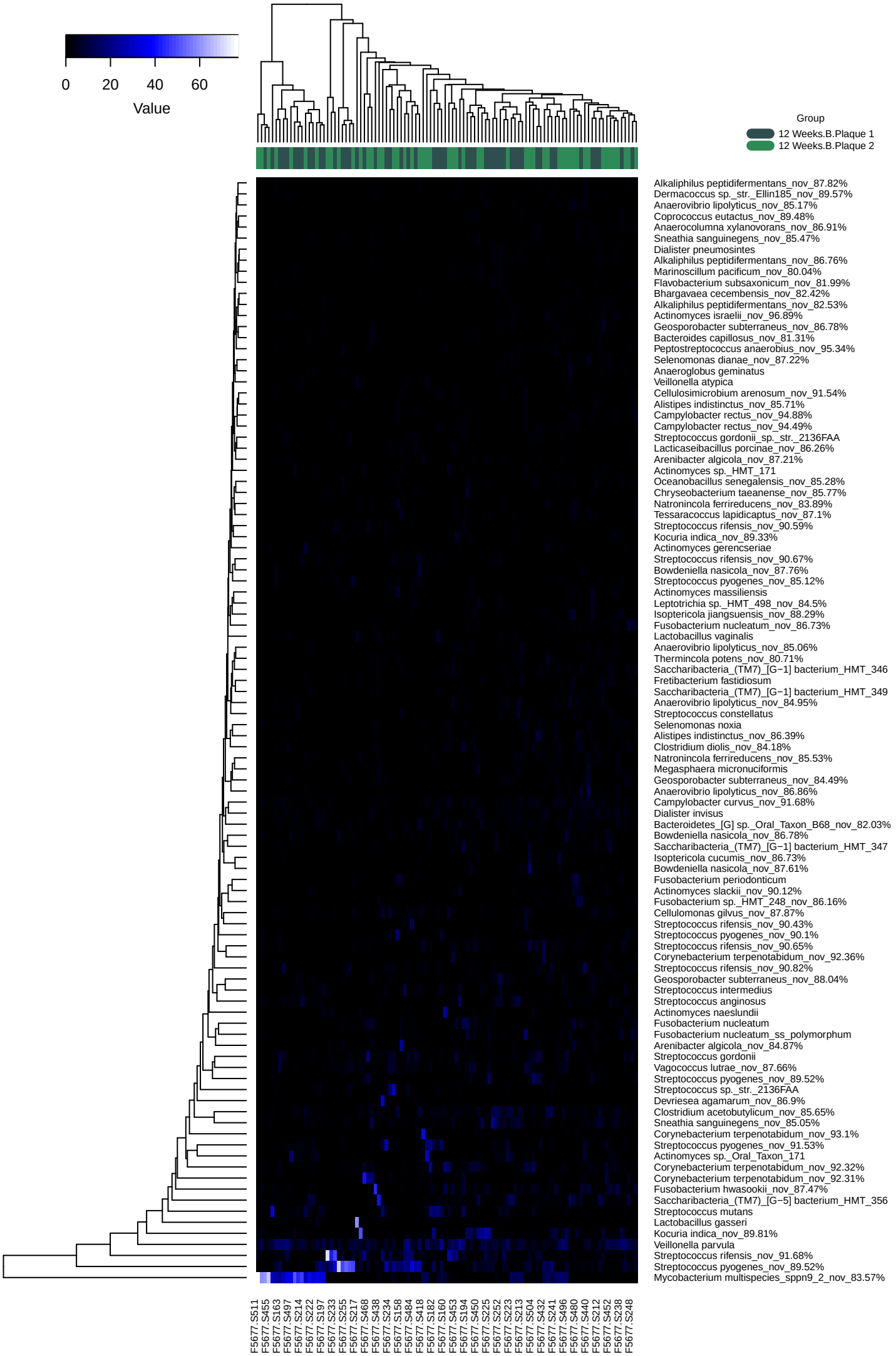
0
20
40
60
Value
Group
12 Weeks.B.Plaque 1
12 Weeks.B.Plaque 2
Alkaliphilus peptidifermentans_nov_87.82%
Dermacoccus sp._str._Ellin185_nov_89.57%
Anaerovibrio lipolyticus_nov_85.17%
Coprococcus eutactus_nov_89.48%
Anaerocolumna xylanovorans_nov_86.91%
Sneathia sanguinegens_nov_85.47%
Dialister pneumosintes
Alkaliphilus peptidifermentans_nov_86.76%
Marinoscillum pacificum_nov_80.04%
Flavobacterium subsaxonicum_nov_81.99%
Bhargavaea cecembensis_nov_82.42%
Alkaliphilus peptidifermentans_nov_82.53%
Actinomyces israelii_nov_96.89%
Geosporobacter subterraneus_nov_86.78%
Bacteroides capillosus_nov_81.31%
Peptostreptococcus anaerobius_nov_95.34%
Selenomonas dianae_nov_87.22%
Anaeroglobus geminatus
Veillonella atypica
Cellulosimicrobium arenosum_nov_91.54%
Alistipes indistinctus_nov_85.71%
Campylobacter rectus_nov_94.88%
Campylobacter rectus_nov_94.49%
Streptococcus gordonii_sp._str._2136FAA
Lacticaseibacillus porcinae_nov_86.26%
Arenibacter algicola_nov_87.21%
Actinomyces sp._HMT_171
Oceanobacillus senegalensis_nov_85.28%
Chryseobacterium taeanense_nov_85.77%
Natronincola ferrireducens_nov_83.89%
Tessaracoccus lapidicaptus_nov_87.1%
Streptococcus rifensis_nov_90.59%
Kocuria indica_nov_89.33%
Actinomyces gerencseriae
Streptococcus rifensis_nov_90.67%
Bowdeniella nasicola_nov_87.76%
Streptococcus pyogenes_nov_85.12%
Actinomyces massiliensis
Leptotrichia sp._HMT_498_nov_84.5%
Isoptericola jiangsuensis_nov_88.29%
Fusobacterium nucleatum_nov_86.73%
Lactobacillus vaginalis
Anaerovibrio lipolyticus_nov_85.06%
Thermincola potens_nov_80.71%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Fretibacterium fastidiosum
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Anaerovibrio lipolyticus_nov_84.95%
Streptococcus constellatus
Selenomonas noxia
Alistipes indistinctus_nov_86.39%
Clostridium diolis_nov_84.18%
Natronincola ferrireducens_nov_85.53%
Megasphaera micronuciformis
Geosporobacter subterraneus_nov_84.49%
Anaerovibrio lipolyticus_nov_86.86%
Campylobacter curvus_nov_91.68%
Dialister invisus
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Bowdeniella nasicola_nov_86.78%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Isoptericola cucumis_nov_86.73%
Bowdeniella nasicola_nov_87.61%
Fusobacterium periodonticum
Actinomyces slackii_nov_90.12%
Fusobacterium sp._HMT_248_nov_86.16%
Cellulomonas gilvus_nov_87.87%
Streptococcus rifensis_nov_90.43%
Streptococcus pyogenes_nov_90.1%
Streptococcus rifensis_nov_90.65%
Corynebacterium terpenotabidum_nov_92.36%
Streptococcus rifensis_nov_90.82%
Geosporobacter subterraneus_nov_88.04%
Streptococcus intermedius
Streptococcus anginosus
Actinomyces naeslundii
Fusobacterium nucleatum
Fusobacterium nucleatum_ss_polymorphum
Arenibacter algicola_nov_84.87%
Streptococcus gordonii
Vagococcus lutrae_nov_87.66%
Streptococcus pyogenes_nov_89.52%
Streptococcus sp._str._2136FAA
Devriesea agamarum_nov_86.9%
Clostridium acetobutylicum_nov_85.65%
Sneathia sanguinegens_nov_85.05%
Corynebacterium terpenotabidum_nov_93.1%
Streptococcus pyogenes_nov_91.53%
Actinomyces sp._Oral_Taxon_171
Corynebacterium terpenotabidum_nov_92.32%
Corynebacterium terpenotabidum_nov_92.31%
Fusobacterium hwasookii_nov_87.47%
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Streptococcus mutans
Lactobacillus gasseri
Kocuria indica_nov_89.81%
Veillonella parvula
Streptococcus rifensis_nov_91.68%
Streptococcus pyogenes_nov_89.52%
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S511
F5677.S455
F5677.S163
F5677.S497
F5677.S214
F5677.S222
F5677.S197
F5677.S233
F5677.S255
F5677.S217
F5677.S468
F5677.S438
F5677.S234
F5677.S158
F5677.S484
F5677.S418
F5677.S182
F5677.S160
F5677.S453
F5677.S194
F5677.S450
F5677.S225
F5677.S252
F5677.S223
F5677.S213
F5677.S504
F5677.S432
F5677.S241
F5677.S496
F5677.S480
F5677.S440
F5677.S212
F5677.S452
F5677.S238
F5677.S248
Samples