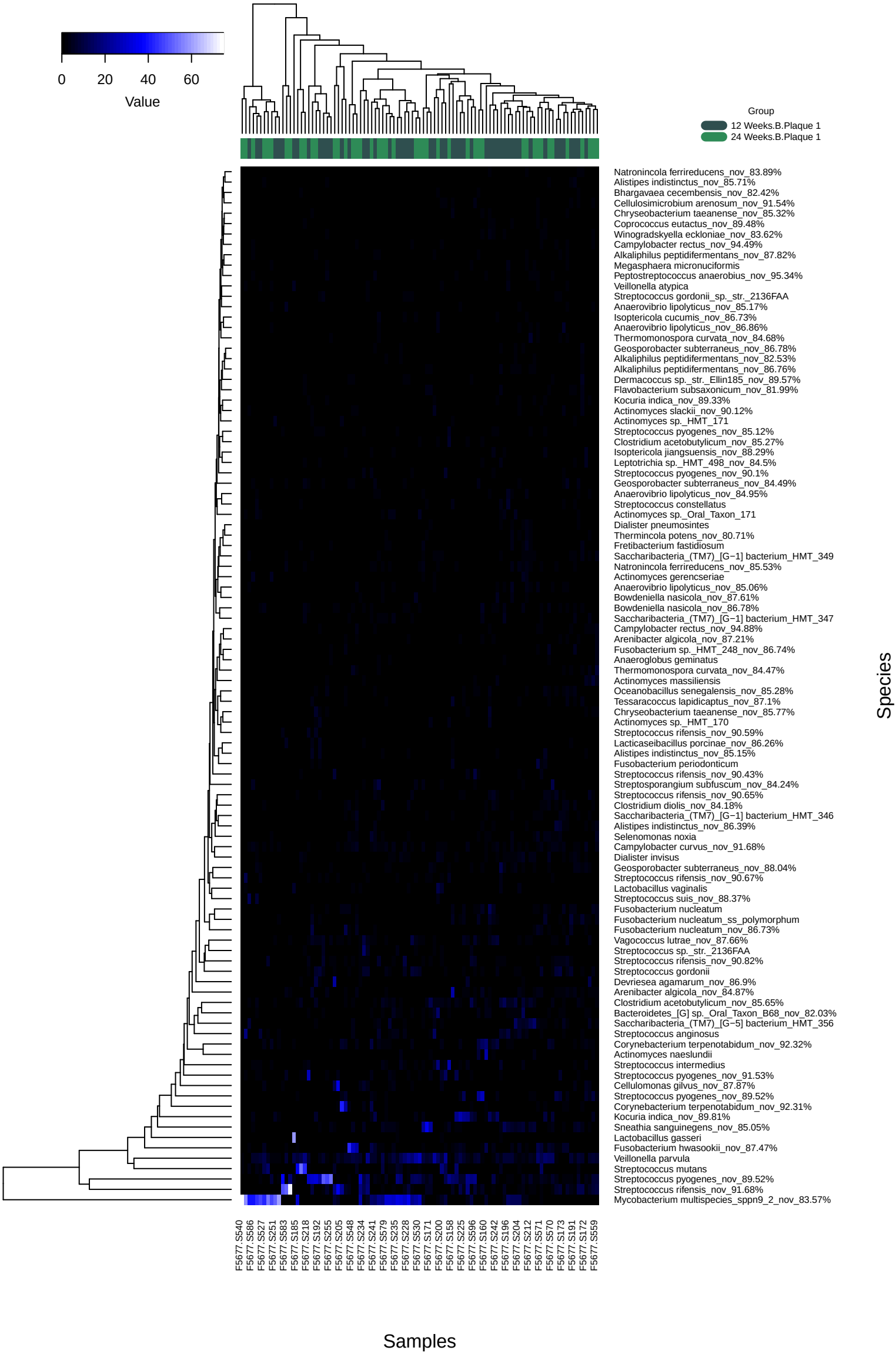
0
20
40
60
Value
Group
12 Weeks.B.Plaque 1
24 Weeks.B.Plaque 1
Natronincola ferrireducens_nov_83.89%
Alistipes indistinctus_nov_85.71%
Bhargavaea cecembensis_nov_82.42%
Cellulosimicrobium arenosum_nov_91.54%
Chryseobacterium taeanense_nov_85.32%
Coprococcus eutactus_nov_89.48%
Winogradskyella eckloniae_nov_83.62%
Campylobacter rectus_nov_94.49%
Alkaliphilus peptidifermentans_nov_87.82%
Megasphaera micronuciformis
Peptostreptococcus anaerobius_nov_95.34%
Veillonella atypica
Streptococcus gordonii_sp._str._2136FAA
Anaerovibrio lipolyticus_nov_85.17%
Isoptericola cucumis_nov_86.73%
Anaerovibrio lipolyticus_nov_86.86%
Thermomonospora curvata_nov_84.68%
Geosporobacter subterraneus_nov_86.78%
Alkaliphilus peptidifermentans_nov_82.53%
Alkaliphilus peptidifermentans_nov_86.76%
Dermacoccus sp._str._Ellin185_nov_89.57%
Flavobacterium subsaxonicum_nov_81.99%
Kocuria indica_nov_89.33%
Actinomyces slackii_nov_90.12%
Actinomyces sp._HMT_171
Streptococcus pyogenes_nov_85.12%
Clostridium acetobutylicum_nov_85.27%
Isoptericola jiangsuensis_nov_88.29%
Leptotrichia sp._HMT_498_nov_84.5%
Streptococcus pyogenes_nov_90.1%
Geosporobacter subterraneus_nov_84.49%
Anaerovibrio lipolyticus_nov_84.95%
Streptococcus constellatus
Actinomyces sp._Oral_Taxon_171
Dialister pneumosintes
Thermincola potens_nov_80.71%
Fretibacterium fastidiosum
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Natronincola ferrireducens_nov_85.53%
Actinomyces gerencseriae
Anaerovibrio lipolyticus_nov_85.06%
Bowdeniella nasicola_nov_87.61%
Bowdeniella nasicola_nov_86.78%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Campylobacter rectus_nov_94.88%
Arenibacter algicola_nov_87.21%
Fusobacterium sp._HMT_248_nov_86.74%
Anaeroglobus geminatus
Thermomonospora curvata_nov_84.47%
Actinomyces massiliensis
Oceanobacillus senegalensis_nov_85.28%
Tessaracoccus lapidicaptus_nov_87.1%
Chryseobacterium taeanense_nov_85.77%
Actinomyces sp._HMT_170
Streptococcus rifensis_nov_90.59%
Lacticaseibacillus porcinae_nov_86.26%
Alistipes indistinctus_nov_85.15%
Fusobacterium periodonticum
Streptococcus rifensis_nov_90.43%
Streptosporangium subfuscum_nov_84.24%
Streptococcus rifensis_nov_90.65%
Clostridium diolis_nov_84.18%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Alistipes indistinctus_nov_86.39%
Selenomonas noxia
Campylobacter curvus_nov_91.68%
Dialister invisus
Geosporobacter subterraneus_nov_88.04%
Streptococcus rifensis_nov_90.67%
Lactobacillus vaginalis
Streptococcus suis_nov_88.37%
Fusobacterium nucleatum
Fusobacterium nucleatum_ss_polymorphum
Fusobacterium nucleatum_nov_86.73%
Vagococcus lutrae_nov_87.66%
Streptococcus sp._str._2136FAA
Streptococcus rifensis_nov_90.82%
Streptococcus gordonii
Devriesea agamarum_nov_86.9%
Arenibacter algicola_nov_84.87%
Clostridium acetobutylicum_nov_85.65%
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Streptococcus anginosus
Corynebacterium terpenotabidum_nov_92.32%
Actinomyces naeslundii
Streptococcus intermedius
Streptococcus pyogenes_nov_91.53%
Cellulomonas gilvus_nov_87.87%
Streptococcus pyogenes_nov_89.52%
Corynebacterium terpenotabidum_nov_92.31%
Kocuria indica_nov_89.81%
Sneathia sanguinegens_nov_85.05%
Lactobacillus gasseri
Fusobacterium hwasookii_nov_87.47%
Veillonella parvula
Streptococcus mutans
Streptococcus pyogenes_nov_89.52%
Streptococcus rifensis_nov_91.68%
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S540
F5677.S586
F5677.S527
F5677.S251
F5677.S583
F5677.S185
F5677.S218
F5677.S192
F5677.S255
F5677.S205
F5677.S548
F5677.S234
F5677.S241
F5677.S579
F5677.S235
F5677.S228
F5677.S530
F5677.S171
F5677.S200
F5677.S158
F5677.S225
F5677.S596
F5677.S160
F5677.S242
F5677.S196
F5677.S204
F5677.S212
F5677.S571
F5677.S570
F5677.S173
F5677.S191
F5677.S172
F5677.S559
Samples