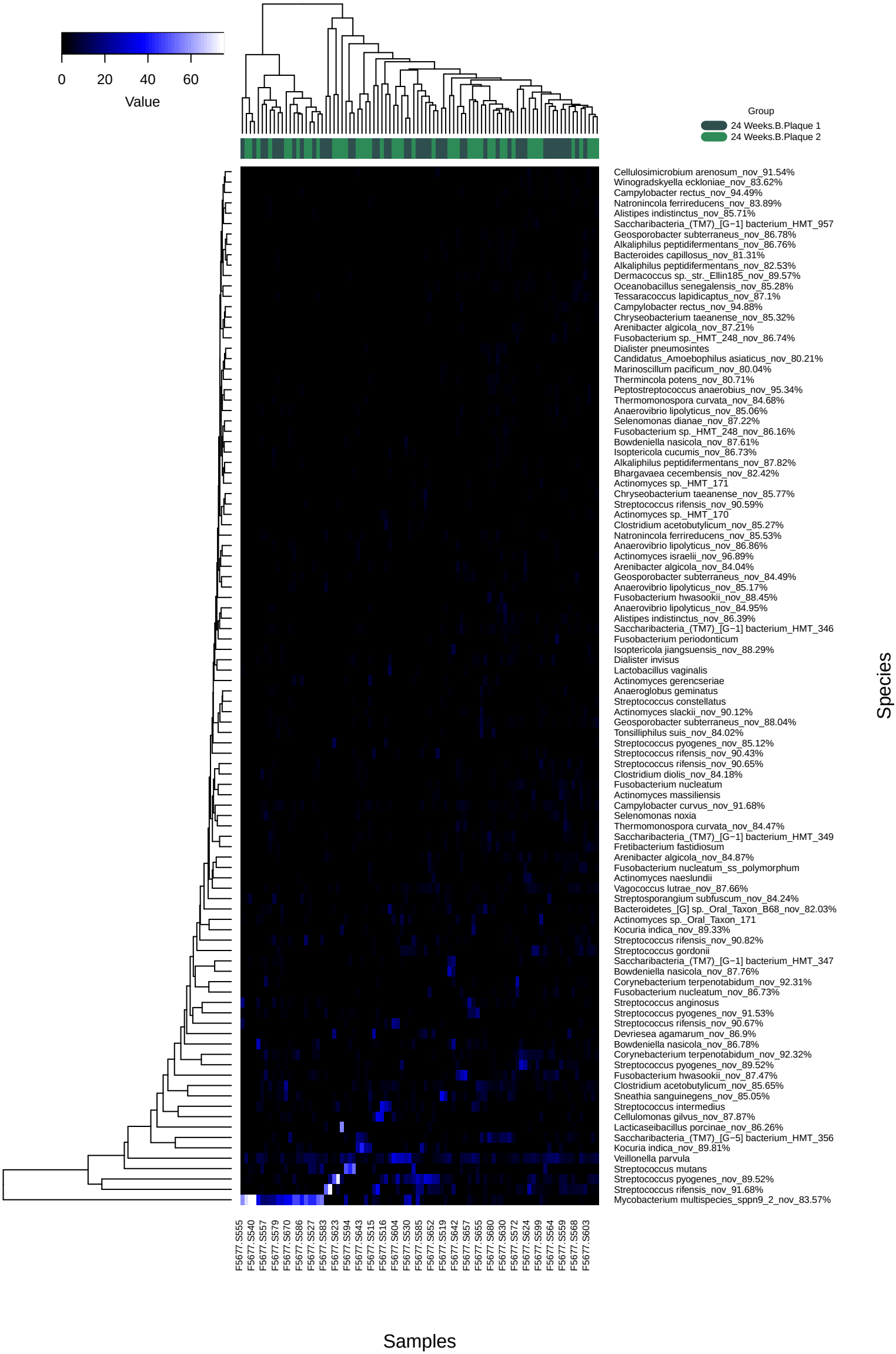
0
20
40
60
Value
Group
24 Weeks.B.Plaque 1
24 Weeks.B.Plaque 2
Cellulosimicrobium arenosum_nov_91.54%
Winogradskyella eckloniae_nov_83.62%
Campylobacter rectus_nov_94.49%
Natronincola ferrireducens_nov_83.89%
Alistipes indistinctus_nov_85.71%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_957
Geosporobacter subterraneus_nov_86.78%
Alkaliphilus peptidifermentans_nov_86.76%
Bacteroides capillosus_nov_81.31%
Alkaliphilus peptidifermentans_nov_82.53%
Dermacoccus sp._str._Ellin185_nov_89.57%
Oceanobacillus senegalensis_nov_85.28%
Tessaracoccus lapidicaptus_nov_87.1%
Campylobacter rectus_nov_94.88%
Chryseobacterium taeanense_nov_85.32%
Arenibacter algicola_nov_87.21%
Fusobacterium sp._HMT_248_nov_86.74%
Dialister pneumosintes
Candidatus_Amoebophilus asiaticus_nov_80.21%
Marinoscillum pacificum_nov_80.04%
Thermincola potens_nov_80.71%
Peptostreptococcus anaerobius_nov_95.34%
Thermomonospora curvata_nov_84.68%
Anaerovibrio lipolyticus_nov_85.06%
Selenomonas dianae_nov_87.22%
Fusobacterium sp._HMT_248_nov_86.16%
Bowdeniella nasicola_nov_87.61%
Isoptericola cucumis_nov_86.73%
Alkaliphilus peptidifermentans_nov_87.82%
Bhargavaea cecembensis_nov_82.42%
Actinomyces sp._HMT_171
Chryseobacterium taeanense_nov_85.77%
Streptococcus rifensis_nov_90.59%
Actinomyces sp._HMT_170
Clostridium acetobutylicum_nov_85.27%
Natronincola ferrireducens_nov_85.53%
Anaerovibrio lipolyticus_nov_86.86%
Actinomyces israelii_nov_96.89%
Arenibacter algicola_nov_84.04%
Geosporobacter subterraneus_nov_84.49%
Anaerovibrio lipolyticus_nov_85.17%
Fusobacterium hwasookii_nov_88.45%
Anaerovibrio lipolyticus_nov_84.95%
Alistipes indistinctus_nov_86.39%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Fusobacterium periodonticum
Isoptericola jiangsuensis_nov_88.29%
Dialister invisus
Lactobacillus vaginalis
Actinomyces gerencseriae
Anaeroglobus geminatus
Streptococcus constellatus
Actinomyces slackii_nov_90.12%
Geosporobacter subterraneus_nov_88.04%
Tonsilliphilus suis_nov_84.02%
Streptococcus pyogenes_nov_85.12%
Streptococcus rifensis_nov_90.43%
Streptococcus rifensis_nov_90.65%
Clostridium diolis_nov_84.18%
Fusobacterium nucleatum
Actinomyces massiliensis
Campylobacter curvus_nov_91.68%
Selenomonas noxia
Thermomonospora curvata_nov_84.47%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Fretibacterium fastidiosum
Arenibacter algicola_nov_84.87%
Fusobacterium nucleatum_ss_polymorphum
Actinomyces naeslundii
Vagococcus lutrae_nov_87.66%
Streptosporangium subfuscum_nov_84.24%
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Actinomyces sp._Oral_Taxon_171
Kocuria indica_nov_89.33%
Streptococcus rifensis_nov_90.82%
Streptococcus gordonii
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Bowdeniella nasicola_nov_87.76%
Corynebacterium terpenotabidum_nov_92.31%
Fusobacterium nucleatum_nov_86.73%
Streptococcus anginosus
Streptococcus pyogenes_nov_91.53%
Streptococcus rifensis_nov_90.67%
Devriesea agamarum_nov_86.9%
Bowdeniella nasicola_nov_86.78%
Corynebacterium terpenotabidum_nov_92.32%
Streptococcus pyogenes_nov_89.52%
Fusobacterium hwasookii_nov_87.47%
Clostridium acetobutylicum_nov_85.65%
Sneathia sanguinegens_nov_85.05%
Streptococcus intermedius
Cellulomonas gilvus_nov_87.87%
Lacticaseibacillus porcinae_nov_86.26%
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Kocuria indica_nov_89.81%
Veillonella parvula
Streptococcus mutans
Streptococcus pyogenes_nov_89.52%
Streptococcus rifensis_nov_91.68%
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S555
F5677.S540
F5677.S557
F5677.S579
F5677.S670
F5677.S586
F5677.S527
F5677.S583
F5677.S623
F5677.S594
F5677.S643
F5677.S515
F5677.S516
F5677.S604
F5677.S530
F5677.S585
F5677.S652
F5677.S519
F5677.S642
F5677.S657
F5677.S655
F5677.S680
F5677.S630
F5677.S572
F5677.S624
F5677.S599
F5677.S564
F5677.S559
F5677.S568
F5677.S603
Samples