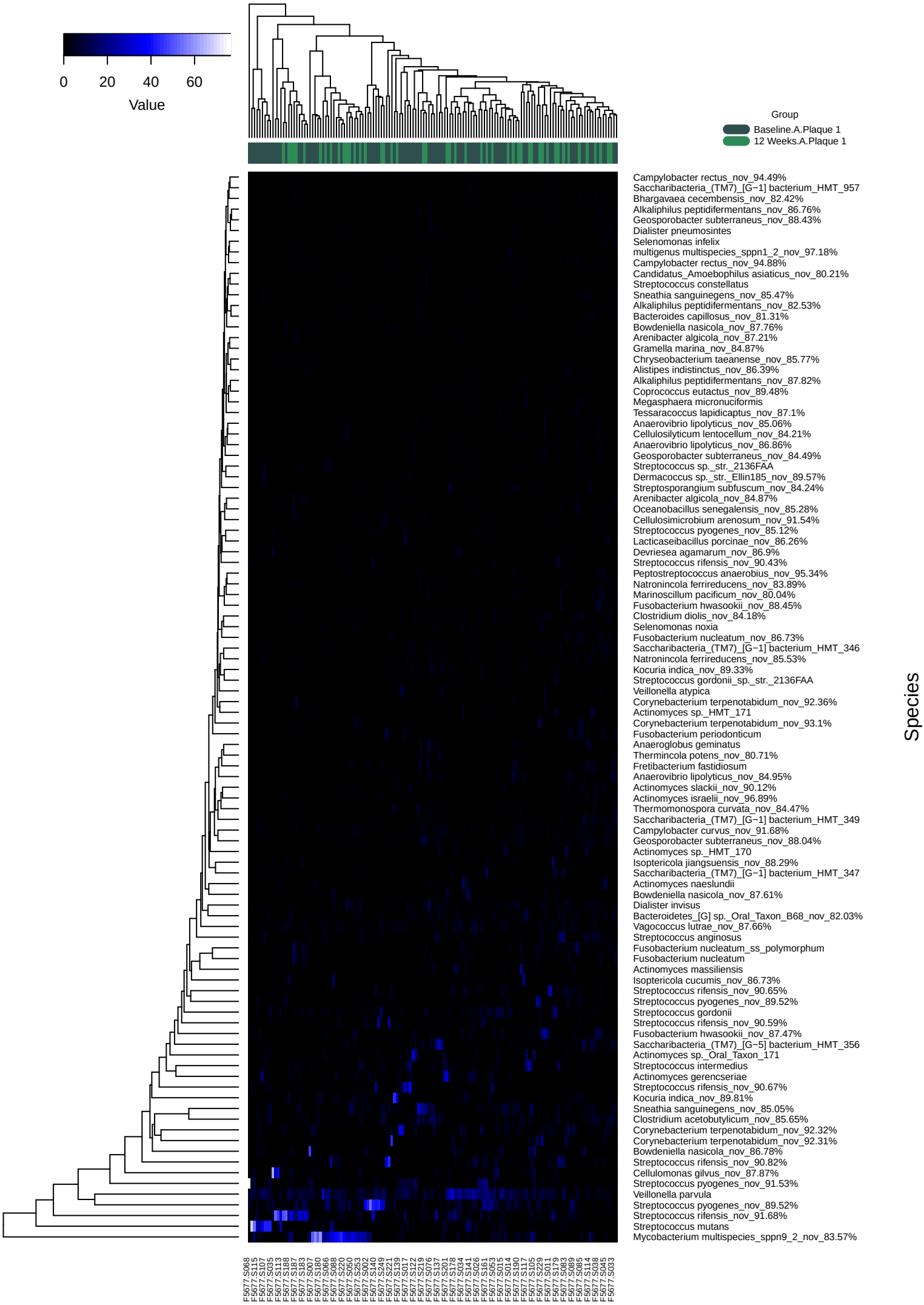
0
20
40
60
Value
Group
Baseline.A.Plaque 1
12 Weeks.A.Plaque 1
Campylobacter rectus_nov_94.49%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_957
Bhargavaea cecembensis_nov_82.42%
Alkaliphilus peptidifermentans_nov_86.76%
Geosporobacter subterraneus_nov_88.43%
Dialister pneumosintes
Selenomonas infelix
multigenus multispecies_sppn1_2_nov_97.18%
Campylobacter rectus_nov_94.88%
Candidatus_Amoebophilus asiaticus_nov_80.21%
Streptococcus constellatus
Sneathia sanguinegens_nov_85.47%
Alkaliphilus peptidifermentans_nov_82.53%
Bacteroides capillosus_nov_81.31%
Bowdeniella nasicola_nov_87.76%
Arenibacter algicola_nov_87.21%
Gramella marina_nov_84.87%
Chryseobacterium taeanense_nov_85.77%
Alistipes indistinctus_nov_86.39%
Alkaliphilus peptidifermentans_nov_87.82%
Coprococcus eutactus_nov_89.48%
Megasphaera micronuciformis
Tessaracoccus lapidicaptus_nov_87.1%
Anaerovibrio lipolyticus_nov_85.06%
Cellulosilyticum lentocellum_nov_84.21%
Anaerovibrio lipolyticus_nov_86.86%
Geosporobacter subterraneus_nov_84.49%
Streptococcus sp._str._2136FAA
Dermacoccus sp._str._Ellin185_nov_89.57%
Streptosporangium subfuscum_nov_84.24%
Arenibacter algicola_nov_84.87%
Oceanobacillus senegalensis_nov_85.28%
Cellulosimicrobium arenosum_nov_91.54%
Streptococcus pyogenes_nov_85.12%
Lacticaseibacillus porcinae_nov_86.26%
Devriesea agamarum_nov_86.9%
Streptococcus rifensis_nov_90.43%
Peptostreptococcus anaerobius_nov_95.34%
Natronincola ferrireducens_nov_83.89%
Marinoscillum pacificum_nov_80.04%
Fusobacterium hwasookii_nov_88.45%
Clostridium diolis_nov_84.18%
Selenomonas noxia
Fusobacterium nucleatum_nov_86.73%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Natronincola ferrireducens_nov_85.53%
Kocuria indica_nov_89.33%
Streptococcus gordonii_sp._str._2136FAA
Veillonella atypica
Corynebacterium terpenotabidum_nov_92.36%
Actinomyces sp._HMT_171
Corynebacterium terpenotabidum_nov_93.1%
Fusobacterium periodonticum
Anaeroglobus geminatus
Thermincola potens_nov_80.71%
Fretibacterium fastidiosum
Anaerovibrio lipolyticus_nov_84.95%
Actinomyces slackii_nov_90.12%
Actinomyces israelii_nov_96.89%
Thermomonospora curvata_nov_84.47%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Campylobacter curvus_nov_91.68%
Geosporobacter subterraneus_nov_88.04%
Actinomyces sp._HMT_170
Isoptericola jiangsuensis_nov_88.29%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Actinomyces naeslundii
Bowdeniella nasicola_nov_87.61%
Dialister invisus
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Vagococcus lutrae_nov_87.66%
Streptococcus anginosus
Fusobacterium nucleatum_ss_polymorphum
Fusobacterium nucleatum
Actinomyces massiliensis
Isoptericola cucumis_nov_86.73%
Streptococcus rifensis_nov_90.65%
Streptococcus pyogenes_nov_89.52%
Streptococcus gordonii
Streptococcus rifensis_nov_90.59%
Fusobacterium hwasookii_nov_87.47%
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Actinomyces sp._Oral_Taxon_171
Streptococcus intermedius
Actinomyces gerencseriae
Streptococcus rifensis_nov_90.67%
Kocuria indica_nov_89.81%
Sneathia sanguinegens_nov_85.05%
Clostridium acetobutylicum_nov_85.65%
Corynebacterium terpenotabidum_nov_92.32%
Corynebacterium terpenotabidum_nov_92.31%
Bowdeniella nasicola_nov_86.78%
Streptococcus rifensis_nov_90.82%
Cellulomonas gilvus_nov_87.87%
Streptococcus pyogenes_nov_91.53%
Veillonella parvula
Streptococcus pyogenes_nov_89.52%
Streptococcus rifensis_nov_91.68%
Streptococcus mutans
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S068
F5677.S115
F5677.S107
F5677.S035
F5677.S113
F5677.S188
F5677.S187
F5677.S183
F5677.S007
F5677.S180
F5677.S066
F5677.S088
F5677.S050
F5677.S253
F5677.S002
F5677.S140
F5677.S249
F5677.S221
F5677.S139
F5677.S017
F5677.S122
F5677.S219
F5677.S076
F5677.S137
F5677.S201
F5677.S178
F5677.S084
F5677.S141
F5677.S026
F5677.S161
F5677.S053
F5677.S015
F5677.S014
F5677.S190
F5677.S117
F5677.S105
F5677.S229
F5677.S011
F5677.S179
F5677.S083
F5677.S089
F5677.S085
F5677.S114
F5677.S038
F5677.S045
F5677.S033
Samples