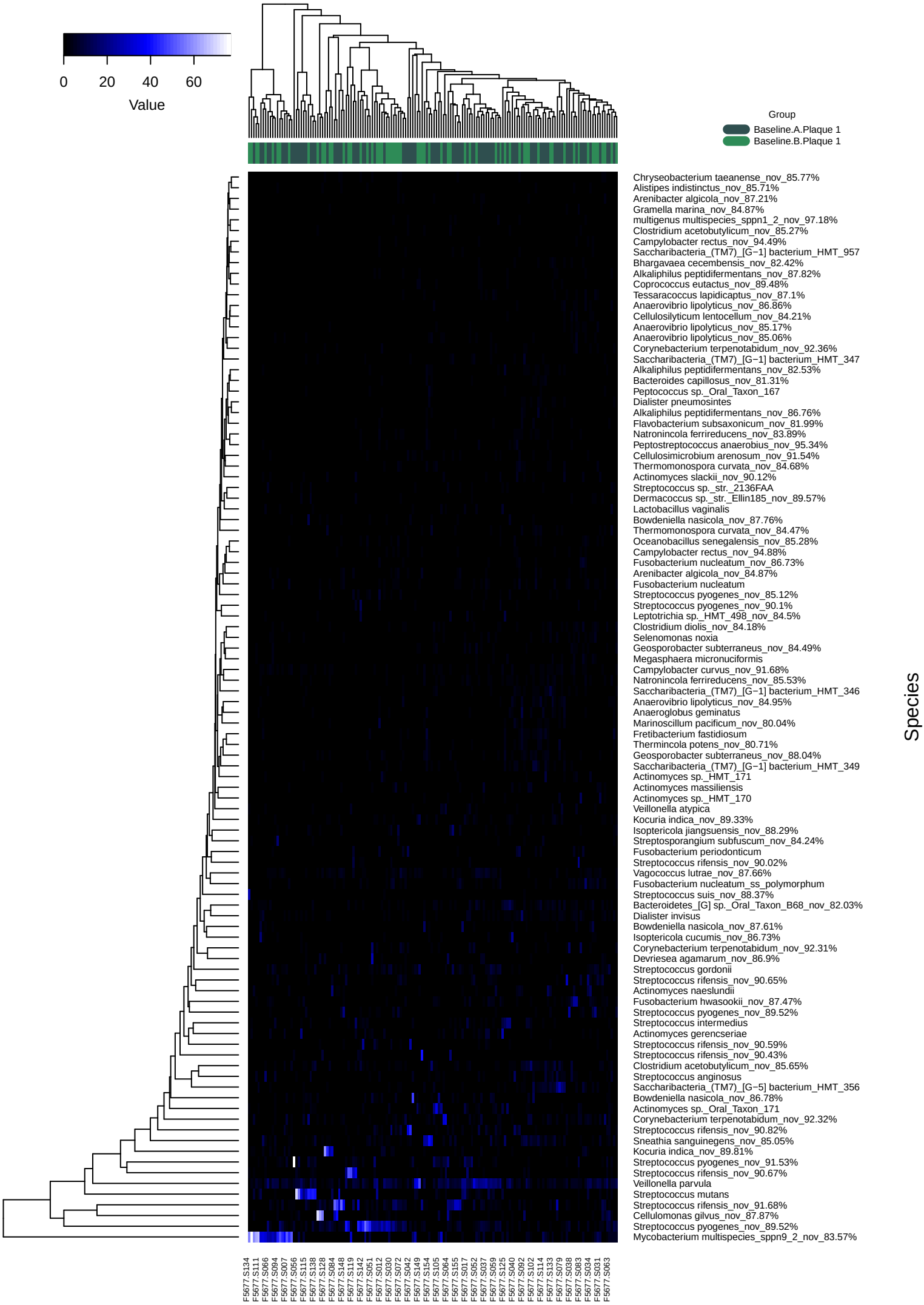
0
20
40
60
Value
Group
Baseline.A.Plaque 1
Baseline.B.Plaque 1
Chryseobacterium taeanense_nov_85.77%
Alistipes indistinctus_nov_85.71%
Arenibacter algicola_nov_87.21%
Gramella marina_nov_84.87%
multigenus multispecies_sppn1_2_nov_97.18%
Clostridium acetobutylicum_nov_85.27%
Campylobacter rectus_nov_94.49%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_957
Bhargavaea cecembensis_nov_82.42%
Alkaliphilus peptidifermentans_nov_87.82%
Coprococcus eutactus_nov_89.48%
Tessaracoccus lapidicaptus_nov_87.1%
Anaerovibrio lipolyticus_nov_86.86%
Cellulosilyticum lentocellum_nov_84.21%
Anaerovibrio lipolyticus_nov_85.17%
Anaerovibrio lipolyticus_nov_85.06%
Corynebacterium terpenotabidum_nov_92.36%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Alkaliphilus peptidifermentans_nov_82.53%
Bacteroides capillosus_nov_81.31%
Peptococcus sp._Oral_Taxon_167
Dialister pneumosintes
Alkaliphilus peptidifermentans_nov_86.76%
Flavobacterium subsaxonicum_nov_81.99%
Natronincola ferrireducens_nov_83.89%
Peptostreptococcus anaerobius_nov_95.34%
Cellulosimicrobium arenosum_nov_91.54%
Thermomonospora curvata_nov_84.68%
Actinomyces slackii_nov_90.12%
Streptococcus sp._str._2136FAA
Dermacoccus sp._str._Ellin185_nov_89.57%
Lactobacillus vaginalis
Bowdeniella nasicola_nov_87.76%
Thermomonospora curvata_nov_84.47%
Oceanobacillus senegalensis_nov_85.28%
Campylobacter rectus_nov_94.88%
Fusobacterium nucleatum_nov_86.73%
Arenibacter algicola_nov_84.87%
Fusobacterium nucleatum
Streptococcus pyogenes_nov_85.12%
Streptococcus pyogenes_nov_90.1%
Leptotrichia sp._HMT_498_nov_84.5%
Clostridium diolis_nov_84.18%
Selenomonas noxia
Geosporobacter subterraneus_nov_84.49%
Megasphaera micronuciformis
Campylobacter curvus_nov_91.68%
Natronincola ferrireducens_nov_85.53%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Anaerovibrio lipolyticus_nov_84.95%
Anaeroglobus geminatus
Marinoscillum pacificum_nov_80.04%
Fretibacterium fastidiosum
Thermincola potens_nov_80.71%
Geosporobacter subterraneus_nov_88.04%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Actinomyces sp._HMT_171
Actinomyces massiliensis
Actinomyces sp._HMT_170
Veillonella atypica
Kocuria indica_nov_89.33%
Isoptericola jiangsuensis_nov_88.29%
Streptosporangium subfuscum_nov_84.24%
Fusobacterium periodonticum
Streptococcus rifensis_nov_90.02%
Vagococcus lutrae_nov_87.66%
Fusobacterium nucleatum_ss_polymorphum
Streptococcus suis_nov_88.37%
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Dialister invisus
Bowdeniella nasicola_nov_87.61%
Isoptericola cucumis_nov_86.73%
Corynebacterium terpenotabidum_nov_92.31%
Devriesea agamarum_nov_86.9%
Streptococcus gordonii
Streptococcus rifensis_nov_90.65%
Actinomyces naeslundii
Fusobacterium hwasookii_nov_87.47%
Streptococcus pyogenes_nov_89.52%
Streptococcus intermedius
Actinomyces gerencseriae
Streptococcus rifensis_nov_90.59%
Streptococcus rifensis_nov_90.43%
Clostridium acetobutylicum_nov_85.65%
Streptococcus anginosus
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Bowdeniella nasicola_nov_86.78%
Actinomyces sp._Oral_Taxon_171
Corynebacterium terpenotabidum_nov_92.32%
Streptococcus rifensis_nov_90.82%
Sneathia sanguinegens_nov_85.05%
Kocuria indica_nov_89.81%
Streptococcus pyogenes_nov_91.53%
Streptococcus rifensis_nov_90.67%
Veillonella parvula
Streptococcus mutans
Streptococcus rifensis_nov_91.68%
Cellulomonas gilvus_nov_87.87%
Streptococcus pyogenes_nov_89.52%
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S134
F5677.S111
F5677.S066
F5677.S094
F5677.S007
F5677.S056
F5677.S115
F5677.S138
F5677.S128
F5677.S084
F5677.S148
F5677.S119
F5677.S142
F5677.S051
F5677.S012
F5677.S030
F5677.S072
F5677.S042
F5677.S149
F5677.S154
F5677.S105
F5677.S064
F5677.S155
F5677.S017
F5677.S052
F5677.S037
F5677.S059
F5677.S125
F5677.S040
F5677.S092
F5677.S102
F5677.S114
F5677.S133
F5677.S079
F5677.S038
F5677.S083
F5677.S034
F5677.S031
F5677.S063
Samples