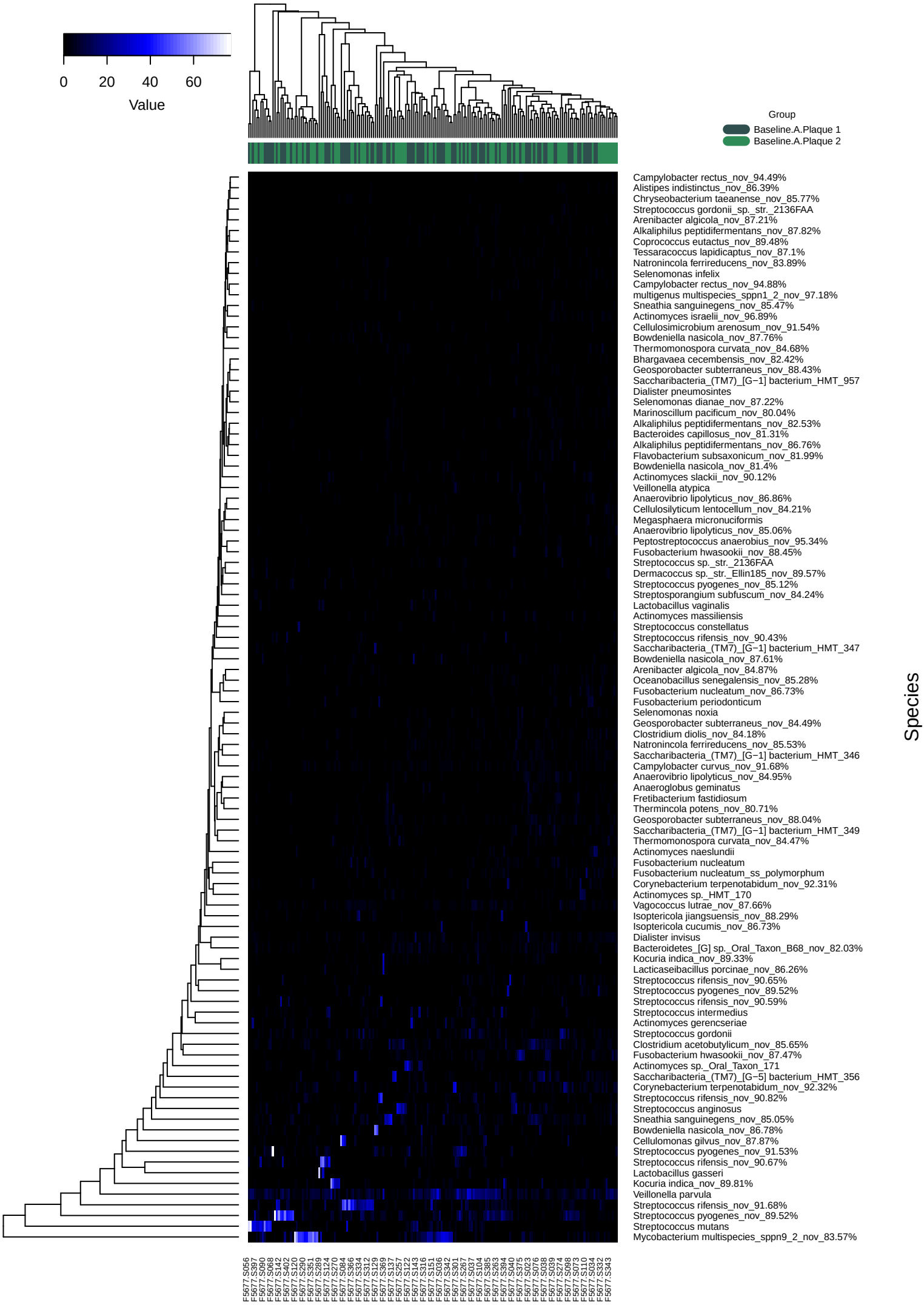
0
20
40
60
Value
Group
Baseline.A.Plaque 1
Baseline.A.Plaque 2
Campylobacter rectus_nov_94.49%
Alistipes indistinctus_nov_86.39%
Chryseobacterium taeanense_nov_85.77%
Streptococcus gordonii_sp._str._2136FAA
Arenibacter algicola_nov_87.21%
Alkaliphilus peptidifermentans_nov_87.82%
Coprococcus eutactus_nov_89.48%
Tessaracoccus lapidicaptus_nov_87.1%
Natronincola ferrireducens_nov_83.89%
Selenomonas infelix
Campylobacter rectus_nov_94.88%
multigenus multispecies_sppn1_2_nov_97.18%
Sneathia sanguinegens_nov_85.47%
Actinomyces israelii_nov_96.89%
Cellulosimicrobium arenosum_nov_91.54%
Bowdeniella nasicola_nov_87.76%
Thermomonospora curvata_nov_84.68%
Bhargavaea cecembensis_nov_82.42%
Geosporobacter subterraneus_nov_88.43%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_957
Dialister pneumosintes
Selenomonas dianae_nov_87.22%
Marinoscillum pacificum_nov_80.04%
Alkaliphilus peptidifermentans_nov_82.53%
Bacteroides capillosus_nov_81.31%
Alkaliphilus peptidifermentans_nov_86.76%
Flavobacterium subsaxonicum_nov_81.99%
Bowdeniella nasicola_nov_81.4%
Actinomyces slackii_nov_90.12%
Veillonella atypica
Anaerovibrio lipolyticus_nov_86.86%
Cellulosilyticum lentocellum_nov_84.21%
Megasphaera micronuciformis
Anaerovibrio lipolyticus_nov_85.06%
Peptostreptococcus anaerobius_nov_95.34%
Fusobacterium hwasookii_nov_88.45%
Streptococcus sp._str._2136FAA
Dermacoccus sp._str._Ellin185_nov_89.57%
Streptococcus pyogenes_nov_85.12%
Streptosporangium subfuscum_nov_84.24%
Lactobacillus vaginalis
Actinomyces massiliensis
Streptococcus constellatus
Streptococcus rifensis_nov_90.43%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Bowdeniella nasicola_nov_87.61%
Arenibacter algicola_nov_84.87%
Oceanobacillus senegalensis_nov_85.28%
Fusobacterium nucleatum_nov_86.73%
Fusobacterium periodonticum
Selenomonas noxia
Geosporobacter subterraneus_nov_84.49%
Clostridium diolis_nov_84.18%
Natronincola ferrireducens_nov_85.53%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Campylobacter curvus_nov_91.68%
Anaerovibrio lipolyticus_nov_84.95%
Anaeroglobus geminatus
Fretibacterium fastidiosum
Thermincola potens_nov_80.71%
Geosporobacter subterraneus_nov_88.04%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Thermomonospora curvata_nov_84.47%
Actinomyces naeslundii
Fusobacterium nucleatum
Fusobacterium nucleatum_ss_polymorphum
Corynebacterium terpenotabidum_nov_92.31%
Actinomyces sp._HMT_170
Vagococcus lutrae_nov_87.66%
Isoptericola jiangsuensis_nov_88.29%
Isoptericola cucumis_nov_86.73%
Dialister invisus
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Kocuria indica_nov_89.33%
Lacticaseibacillus porcinae_nov_86.26%
Streptococcus rifensis_nov_90.65%
Streptococcus pyogenes_nov_89.52%
Streptococcus rifensis_nov_90.59%
Streptococcus intermedius
Actinomyces gerencseriae
Streptococcus gordonii
Clostridium acetobutylicum_nov_85.65%
Fusobacterium hwasookii_nov_87.47%
Actinomyces sp._Oral_Taxon_171
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Corynebacterium terpenotabidum_nov_92.32%
Streptococcus rifensis_nov_90.82%
Streptococcus anginosus
Sneathia sanguinegens_nov_85.05%
Bowdeniella nasicola_nov_86.78%
Cellulomonas gilvus_nov_87.87%
Streptococcus pyogenes_nov_91.53%
Streptococcus rifensis_nov_90.67%
Lactobacillus gasseri
Kocuria indica_nov_89.81%
Veillonella parvula
Streptococcus rifensis_nov_91.68%
Streptococcus pyogenes_nov_89.52%
Streptococcus mutans
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S056
F5677.S397
F5677.S090
F5677.S068
F5677.S142
F5677.S402
F5677.S120
F5677.S290
F5677.S351
F5677.S289
F5677.S124
F5677.S270
F5677.S084
F5677.S366
F5677.S364
F5677.S312
F5677.S129
F5677.S369
F5677.S137
F5677.S257
F5677.S122
F5677.S143
F5677.S316
F5677.S151
F5677.S036
F5677.S342
F5677.S301
F5677.S267
F5677.S037
F5677.S104
F5677.S385
F5677.S263
F5677.S394
F5677.S040
F5677.S375
F5677.S023
F5677.S076
F5677.S038
F5677.S039
F5677.S274
F5677.S098
F5677.S073
F5677.S110
F5677.S034
F5677.S332
F5677.S343
Samples