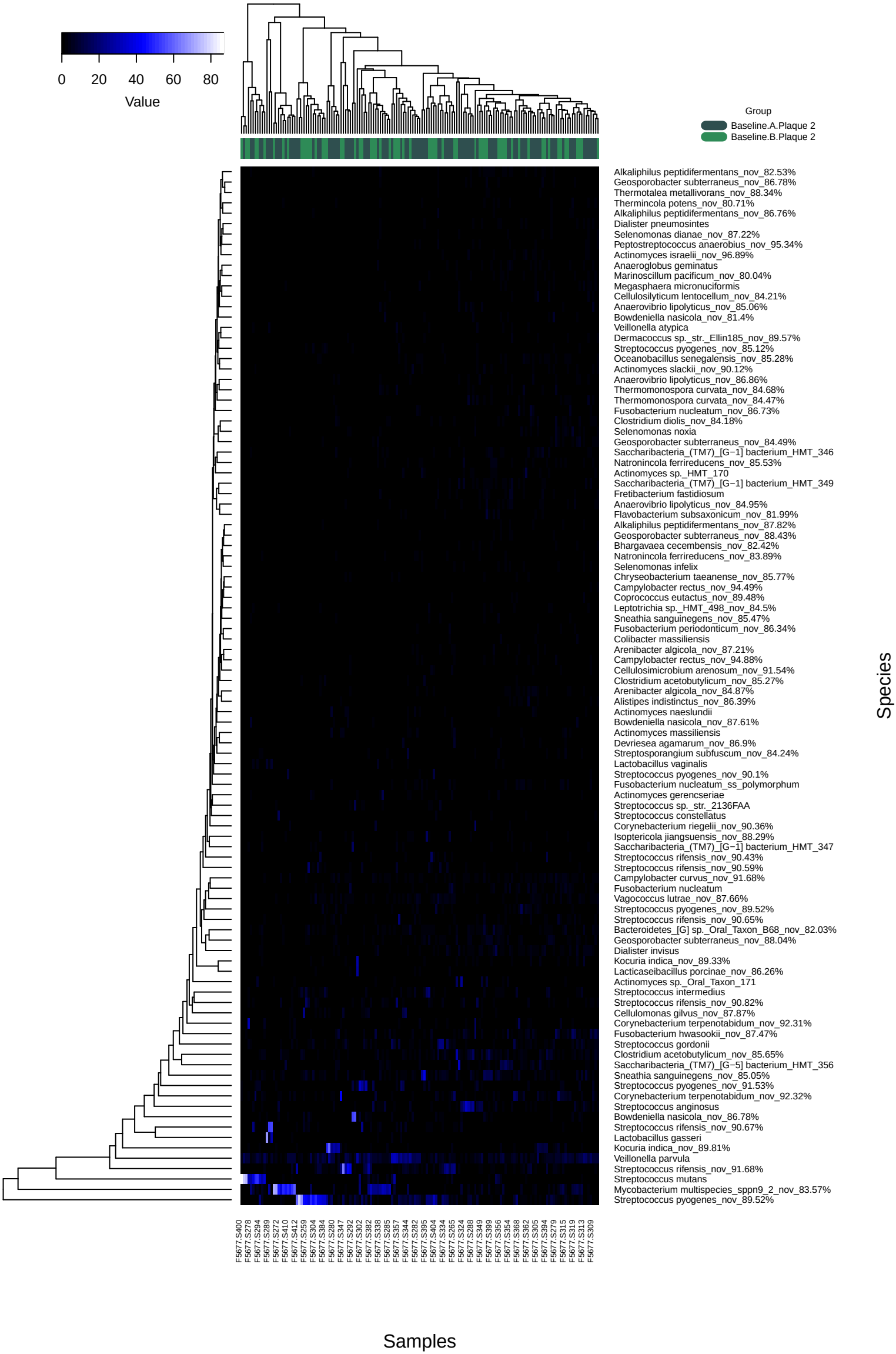
0
20
40
60
80
Value
Group
Baseline.A.Plaque 2
Baseline.B.Plaque 2
Alkaliphilus peptidifermentans_nov_82.53%
Geosporobacter subterraneus_nov_86.78%
Thermotalea metallivorans_nov_88.34%
Thermincola potens_nov_80.71%
Alkaliphilus peptidifermentans_nov_86.76%
Dialister pneumosintes
Selenomonas dianae_nov_87.22%
Peptostreptococcus anaerobius_nov_95.34%
Actinomyces israelii_nov_96.89%
Anaeroglobus geminatus
Marinoscillum pacificum_nov_80.04%
Megasphaera micronuciformis
Cellulosilyticum lentocellum_nov_84.21%
Anaerovibrio lipolyticus_nov_85.06%
Bowdeniella nasicola_nov_81.4%
Veillonella atypica
Dermacoccus sp._str._Ellin185_nov_89.57%
Streptococcus pyogenes_nov_85.12%
Oceanobacillus senegalensis_nov_85.28%
Actinomyces slackii_nov_90.12%
Anaerovibrio lipolyticus_nov_86.86%
Thermomonospora curvata_nov_84.68%
Thermomonospora curvata_nov_84.47%
Fusobacterium nucleatum_nov_86.73%
Clostridium diolis_nov_84.18%
Selenomonas noxia
Geosporobacter subterraneus_nov_84.49%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Natronincola ferrireducens_nov_85.53%
Actinomyces sp._HMT_170
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Fretibacterium fastidiosum
Anaerovibrio lipolyticus_nov_84.95%
Flavobacterium subsaxonicum_nov_81.99%
Alkaliphilus peptidifermentans_nov_87.82%
Geosporobacter subterraneus_nov_88.43%
Bhargavaea cecembensis_nov_82.42%
Natronincola ferrireducens_nov_83.89%
Selenomonas infelix
Chryseobacterium taeanense_nov_85.77%
Campylobacter rectus_nov_94.49%
Coprococcus eutactus_nov_89.48%
Leptotrichia sp._HMT_498_nov_84.5%
Sneathia sanguinegens_nov_85.47%
Fusobacterium periodonticum_nov_86.34%
Colibacter massiliensis
Arenibacter algicola_nov_87.21%
Campylobacter rectus_nov_94.88%
Cellulosimicrobium arenosum_nov_91.54%
Clostridium acetobutylicum_nov_85.27%
Arenibacter algicola_nov_84.87%
Alistipes indistinctus_nov_86.39%
Actinomyces naeslundii
Bowdeniella nasicola_nov_87.61%
Actinomyces massiliensis
Devriesea agamarum_nov_86.9%
Streptosporangium subfuscum_nov_84.24%
Lactobacillus vaginalis
Streptococcus pyogenes_nov_90.1%
Fusobacterium nucleatum_ss_polymorphum
Actinomyces gerencseriae
Streptococcus sp._str._2136FAA
Streptococcus constellatus
Corynebacterium riegelii_nov_90.36%
Isoptericola jiangsuensis_nov_88.29%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Streptococcus rifensis_nov_90.43%
Streptococcus rifensis_nov_90.59%
Campylobacter curvus_nov_91.68%
Fusobacterium nucleatum
Vagococcus lutrae_nov_87.66%
Streptococcus pyogenes_nov_89.52%
Streptococcus rifensis_nov_90.65%
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Geosporobacter subterraneus_nov_88.04%
Dialister invisus
Kocuria indica_nov_89.33%
Lacticaseibacillus porcinae_nov_86.26%
Actinomyces sp._Oral_Taxon_171
Streptococcus intermedius
Streptococcus rifensis_nov_90.82%
Cellulomonas gilvus_nov_87.87%
Corynebacterium terpenotabidum_nov_92.31%
Fusobacterium hwasookii_nov_87.47%
Streptococcus gordonii
Clostridium acetobutylicum_nov_85.65%
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Sneathia sanguinegens_nov_85.05%
Streptococcus pyogenes_nov_91.53%
Corynebacterium terpenotabidum_nov_92.32%
Streptococcus anginosus
Bowdeniella nasicola_nov_86.78%
Streptococcus rifensis_nov_90.67%
Lactobacillus gasseri
Kocuria indica_nov_89.81%
Veillonella parvula
Streptococcus rifensis_nov_91.68%
Streptococcus mutans
Mycobacterium multispecies_sppn9_2_nov_83.57%
Streptococcus pyogenes_nov_89.52%
Species
F5677.S400
F5677.S278
F5677.S294
F5677.S289
F5677.S272
F5677.S410
F5677.S412
F5677.S259
F5677.S304
F5677.S384
F5677.S280
F5677.S347
F5677.S292
F5677.S302
F5677.S382
F5677.S338
F5677.S285
F5677.S357
F5677.S344
F5677.S282
F5677.S395
F5677.S404
F5677.S334
F5677.S265
F5677.S324
F5677.S288
F5677.S349
F5677.S399
F5677.S356
F5677.S354
F5677.S368
F5677.S362
F5677.S305
F5677.S394
F5677.S279
F5677.S315
F5677.S319
F5677.S313
F5677.S309
Samples