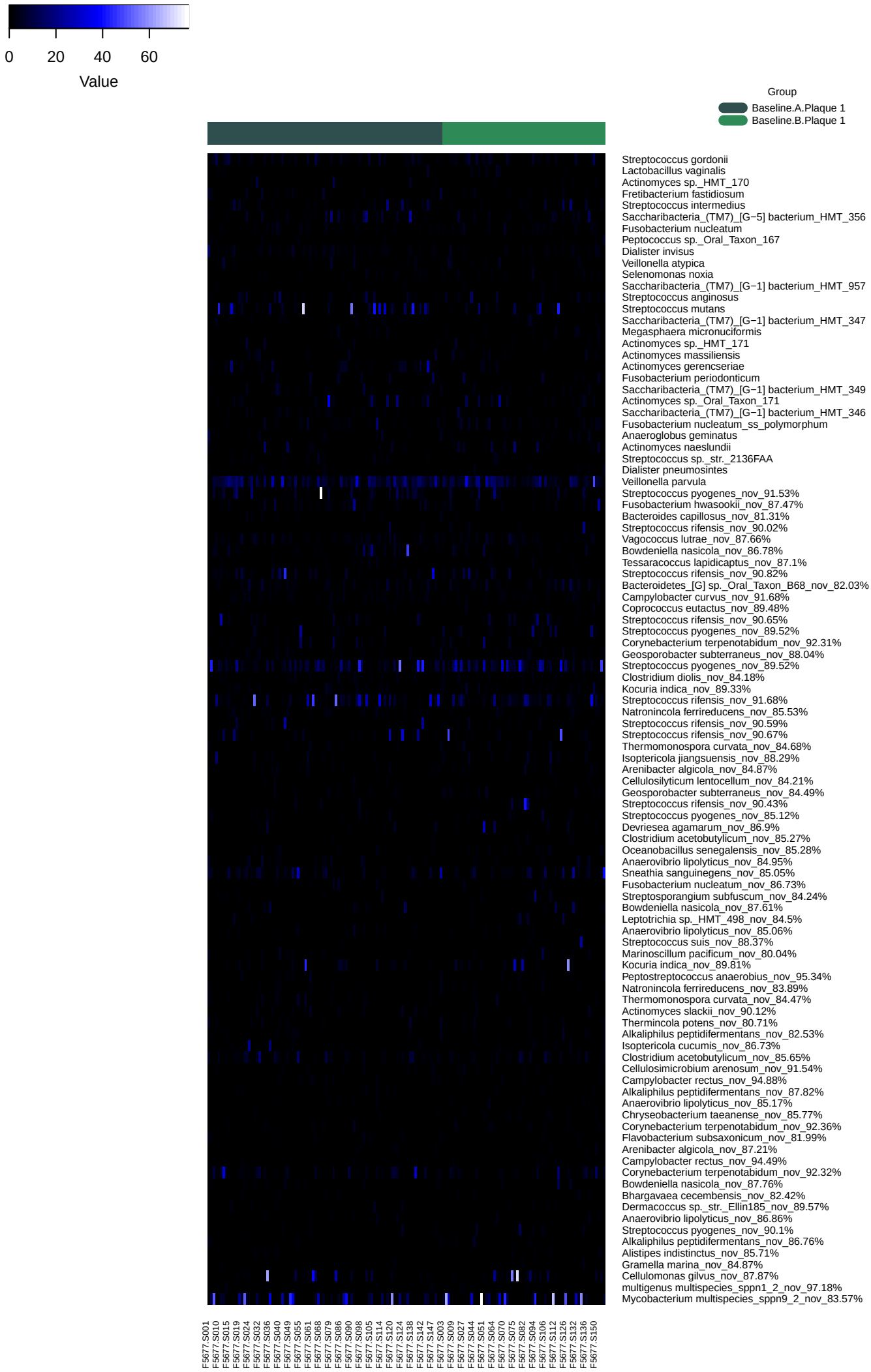
0
20
40
60
Value
Group
Baseline.A.Plaque 1
Baseline.B.Plaque 1
Streptococcus gordonii
Lactobacillus vaginalis
Actinomyces sp._HMT_170
Fretibacterium fastidiosum
Streptococcus intermedius
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Fusobacterium nucleatum
Peptococcus sp._Oral_Taxon_167
Dialister invisus
Veillonella atypica
Selenomonas noxia
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_957
Streptococcus anginosus
Streptococcus mutans
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
Megasphaera micronuciformis
Actinomyces sp._HMT_171
Actinomyces massiliensis
Actinomyces gerencseriae
Fusobacterium periodonticum
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Actinomyces sp._Oral_Taxon_171
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Fusobacterium nucleatum_ss_polymorphum
Anaeroglobus geminatus
Actinomyces naeslundii
Streptococcus sp._str._2136FAA
Dialister pneumosintes
Veillonella parvula
Streptococcus pyogenes_nov_91.53%
Fusobacterium hwasookii_nov_87.47%
Bacteroides capillosus_nov_81.31%
Streptococcus rifensis_nov_90.02%
Vagococcus lutrae_nov_87.66%
Bowdeniella nasicola_nov_86.78%
Tessaracoccus lapidicaptus_nov_87.1%
Streptococcus rifensis_nov_90.82%
Bacteroidetes_[G] sp._Oral_Taxon_B68_nov_82.03%
Campylobacter curvus_nov_91.68%
Coprococcus eutactus_nov_89.48%
Streptococcus rifensis_nov_90.65%
Streptococcus pyogenes_nov_89.52%
Corynebacterium terpenotabidum_nov_92.31%
Geosporobacter subterraneus_nov_88.04%
Streptococcus pyogenes_nov_89.52%
Clostridium diolis_nov_84.18%
Kocuria indica_nov_89.33%
Streptococcus rifensis_nov_91.68%
Natronincola ferrireducens_nov_85.53%
Streptococcus rifensis_nov_90.59%
Streptococcus rifensis_nov_90.67%
Thermomonospora curvata_nov_84.68%
Isoptericola jiangsuensis_nov_88.29%
Arenibacter algicola_nov_84.87%
Cellulosilyticum lentocellum_nov_84.21%
Geosporobacter subterraneus_nov_84.49%
Streptococcus rifensis_nov_90.43%
Streptococcus pyogenes_nov_85.12%
Devriesea agamarum_nov_86.9%
Clostridium acetobutylicum_nov_85.27%
Oceanobacillus senegalensis_nov_85.28%
Anaerovibrio lipolyticus_nov_84.95%
Sneathia sanguinegens_nov_85.05%
Fusobacterium nucleatum_nov_86.73%
Streptosporangium subfuscum_nov_84.24%
Bowdeniella nasicola_nov_87.61%
Leptotrichia sp._HMT_498_nov_84.5%
Anaerovibrio lipolyticus_nov_85.06%
Streptococcus suis_nov_88.37%
Marinoscillum pacificum_nov_80.04%
Kocuria indica_nov_89.81%
Peptostreptococcus anaerobius_nov_95.34%
Natronincola ferrireducens_nov_83.89%
Thermomonospora curvata_nov_84.47%
Actinomyces slackii_nov_90.12%
Thermincola potens_nov_80.71%
Alkaliphilus peptidifermentans_nov_82.53%
Isoptericola cucumis_nov_86.73%
Clostridium acetobutylicum_nov_85.65%
Cellulosimicrobium arenosum_nov_91.54%
Campylobacter rectus_nov_94.88%
Alkaliphilus peptidifermentans_nov_87.82%
Anaerovibrio lipolyticus_nov_85.17%
Chryseobacterium taeanense_nov_85.77%
Corynebacterium terpenotabidum_nov_92.36%
Flavobacterium subsaxonicum_nov_81.99%
Arenibacter algicola_nov_87.21%
Campylobacter rectus_nov_94.49%
Corynebacterium terpenotabidum_nov_92.32%
Bowdeniella nasicola_nov_87.76%
Bhargavaea cecembensis_nov_82.42%
Dermacoccus sp._str._Ellin185_nov_89.57%
Anaerovibrio lipolyticus_nov_86.86%
Streptococcus pyogenes_nov_90.1%
Alkaliphilus peptidifermentans_nov_86.76%
Alistipes indistinctus_nov_85.71%
Gramella marina_nov_84.87%
Cellulomonas gilvus_nov_87.87%
multigenus multispecies_sppn1_2_nov_97.18%
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
F5677.S001
F5677.S010
F5677.S015
F5677.S019
F5677.S024
F5677.S032
F5677.S036
F5677.S040
F5677.S049
F5677.S055
F5677.S061
F5677.S068
F5677.S079
F5677.S086
F5677.S090
F5677.S098
F5677.S105
F5677.S114
F5677.S120
F5677.S124
F5677.S138
F5677.S142
F5677.S147
F5677.S003
F5677.S009
F5677.S027
F5677.S044
F5677.S051
F5677.S064
F5677.S070
F5677.S075
F5677.S082
F5677.S094
F5677.S106
F5677.S112
F5677.S126
F5677.S132
F5677.S136
F5677.S150
Samples