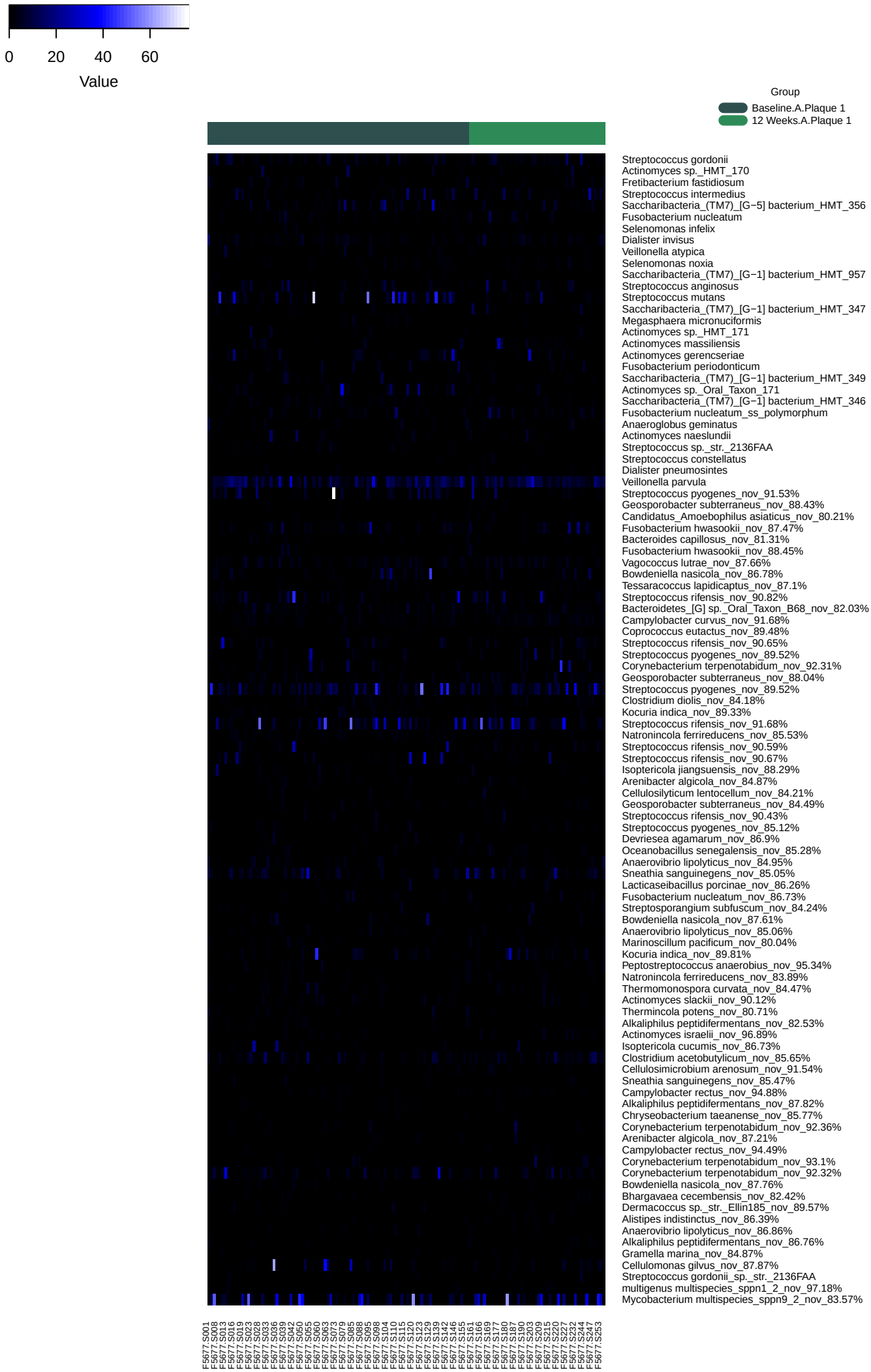
0
20
40
60
Value
Group
Baseline.A.Plaque 1
12 Weeks.A.Plaque 1
Streptococcus gordonii
Actinomyces sp._HMT_170
Fretibacterium fastidiosum
Streptococcus intermedius
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Fusobacterium nucleatum
Selenomonas infelix
Dialister invisus
Veillonella atypica
Selenomonas 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
Streptococcus constellatus
Dialister pneumosintes
Veillonella parvula
Streptococcus pyogenes_nov_91.53%
Geosporobacter subterraneus_nov_88.43%
Candidatus_Amoebophilus asiaticus_nov_80.21%
Fusobacterium hwasookii_nov_87.47%
Bacteroides capillosus_nov_81.31%
Fusobacterium hwasookii_nov_88.45%
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%
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%
Oceanobacillus senegalensis_nov_85.28%
Anaerovibrio lipolyticus_nov_84.95%
Sneathia sanguinegens_nov_85.05%
Lacticaseibacillus porcinae_nov_86.26%
Fusobacterium nucleatum_nov_86.73%
Streptosporangium subfuscum_nov_84.24%
Bowdeniella nasicola_nov_87.61%
Anaerovibrio lipolyticus_nov_85.06%
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%
Actinomyces israelii_nov_96.89%
Isoptericola cucumis_nov_86.73%
Clostridium acetobutylicum_nov_85.65%
Cellulosimicrobium arenosum_nov_91.54%
Sneathia sanguinegens_nov_85.47%
Campylobacter rectus_nov_94.88%
Alkaliphilus peptidifermentans_nov_87.82%
Chryseobacterium taeanense_nov_85.77%
Corynebacterium terpenotabidum_nov_92.36%
Arenibacter algicola_nov_87.21%
Campylobacter rectus_nov_94.49%
Corynebacterium terpenotabidum_nov_93.1%
Corynebacterium terpenotabidum_nov_92.32%
Bowdeniella nasicola_nov_87.76%
Bhargavaea cecembensis_nov_82.42%
Dermacoccus sp._str._Ellin185_nov_89.57%
Alistipes indistinctus_nov_86.39%
Anaerovibrio lipolyticus_nov_86.86%
Alkaliphilus peptidifermentans_nov_86.76%
Gramella marina_nov_84.87%
Cellulomonas gilvus_nov_87.87%
Streptococcus gordonii_sp._str._2136FAA
multigenus multispecies_sppn1_2_nov_97.18%
Mycobacterium multispecies_sppn9_2_nov_83.57%
Species
Samples