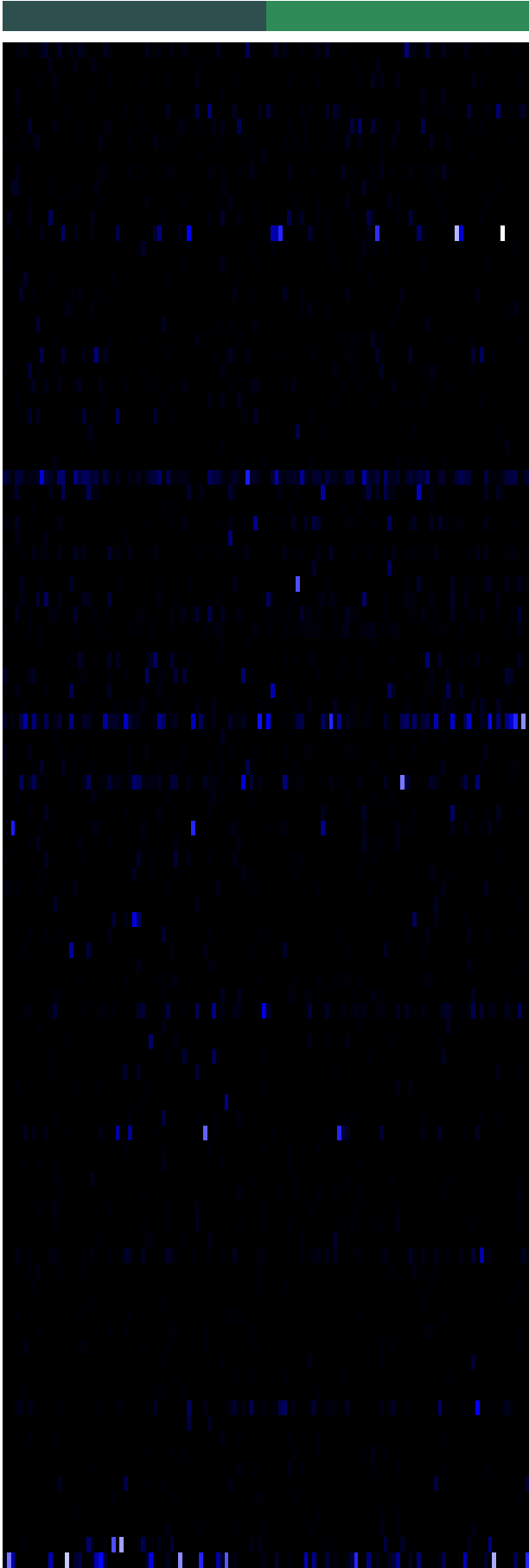
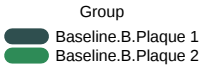
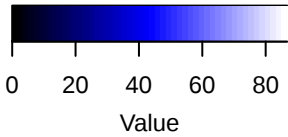




- Streptococcus gordonii
- Lactobacillus vaginalis
- Fretibacterium fastidiosum
- Colibacter massiliensis
- Streptococcus intermedius
- Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356
- Fusobacterium nucleatum
- Peptococcus sp._Oral_Taxon_167
- Dialister invisus
- Veillonella atypica
- Selenomonas noxia
- 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%
- Anaerocolumna xylanovorans_nov_86.91%
- Fusobacterium hwasookii_nov_87.47%
- Streptococcus rifensis_nov_90.02%
- Vagococcus lutrae_nov_87.66%
- Corynebacterium riegelii_nov_90.36%
- Bowdeniella nasicola_nov_86.78%
- 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%
- Bowdeniella nasicola_nov_81.4%
- 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%
- Geosporobacter subterraneus_nov_84.49%
- Tonsilliphilus suis_nov_84.02%
- 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 arenaosum_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%
- Alistipes indistinctus_nov_85.71%
- Selenomonas dianaee_nov_87.22%
- Geosporobacter subterraneus_nov_86.78%
- Cellulomonas gilvus_nov_87.87%
- Mycobacterium multispecies_sppn9_2_nov_83.57%

Species

Samples