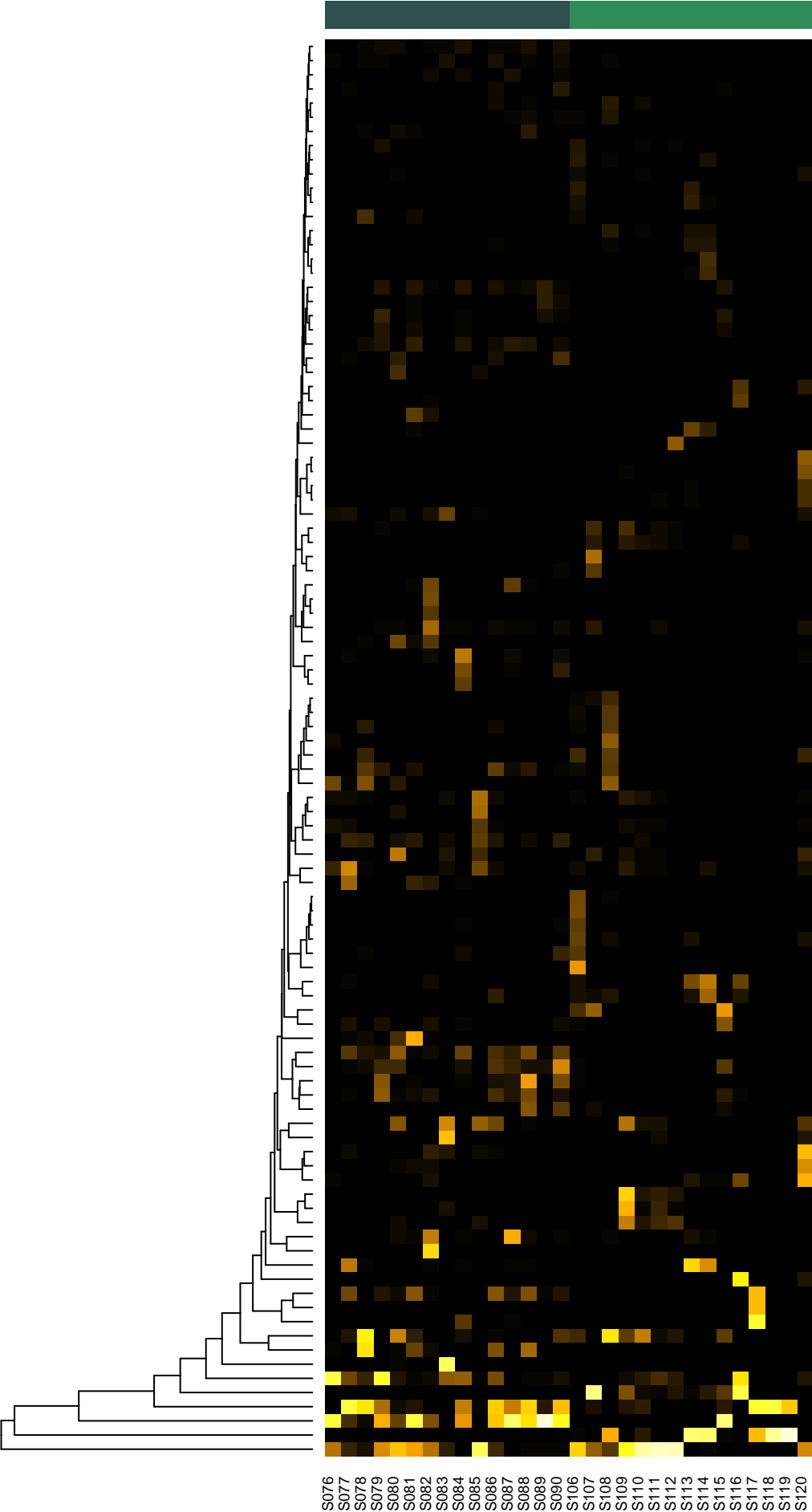




Description2
 PT_NP_PP
 PT_VP_HPT



- Schaalia odontolytica
- Aerococcus urinaeequi_viridans
- Stomatobaculum sp._HMT_373
- Lachnoanaerobaculum sp._HMT_089
- Peptoanaerobacter_Peptostreptococcaceae_[XII][G-7] [Eubacterium]
- Bacteroidales_[G-2] bacterium_HMT_274
- Schaalia meyeri
- Gemella morbillorum
- Catonella morbi_sp._HMT_164
- Neisseria elongata
- Leptotrichia hongkongensis
- Selenomonas sp._HMT_146
- Fusobacterium nucleatum
- Eikenella corrodens
- Lachnospiraceae_[G-3] bacterium_HMT_100
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_869
- Campylobacter concisus
- Peptostreptococcaceae_[XII][G-4] bacterium_HMT_369
- Veillonellaceae_[G-1] bacterium_HMT_145
- Fretibacterium fastidiosum
- Fretibacterium sp._HMT_360
- Mogibacterium timidum
- Schnuerera ultunensis_nov_92.840%
- Pseudomonas alcaliphila_chengduensis_oleovorans_toyotomiensis
- Veillonella parvula
- Actinomyces naeslundii
- Actinomyces sp._HMT_448
- Veillonellaceae_[G-1] bacterium_HMT_155
- Massilia atriviolacea
- Granulicatella elegans
- Bacillus albus_anthraxis_cereus_fungorum_gaemokensis_homini ...
- Gemella haemolysans
- Abiotrophia defectiva
- Granulicatella adiacens
- Rothia aeria
- Sphingomonas aerolata_aurantiaca_faeni_ginsenosidivorax_olei_pa
- Erysipelotrichaceae_[G-1] bacterium_HMT_905
- Veillonellaceae_[G-1] bacterium_HMT_135
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349
- Limosilactobacillus mucosae
- Olsenella sp._HMT_807
- Staphylococcus capitis_caprae_epidermidis
- Rothia mucilaginoso
- Peptostreptococcus stomatis
- Schaalia cardiffensis
- Pseudacidobacterium ailaui_nov_94.321%
- Fusobacterium nucleatum_nucleatum_subsp._animalis
- Cardiobacterium valvarum
- Fusobacterium naviforme_nucleatum_sp._HMT_204
- Fusobacterium nucleatum_nucleatum_subsp._animalis_simiae
- Fusobacterium canifelinum_nucleatum_nucleatum_subsp._polydora
- Fusobacterium nucleatum_nucleatum_subsp._vincentii
- Johnsonella ignava
- Streptococcus lactarius
- Paracoccus aestuarii_beibuensis_hibisci_marinus_pueri
- Streptococcus cristatus_cristatus_clade_578_downii_gwanguense
- Streptococcus intermedius
- Haemophilus parainfluenzae
- Streptococcus cristatus_cristatus_clade_578
- Streptococcus anginosus
- Prevotella intermedia
- Leptotrichia sp._HMT_463
- Porphyromonas gingivalis
- Leptotrichia buccalis
- Fusobacterium nucleatum_subsp._vincentii
- Aggregatibacter sp._HMT_458
- Selenomonas sputigena
- Selenomonas noxia
- Campylobacter rectus_showae
- Johnsonella sp._HMT_166
- Peptoniphilaceae_[G-1] bacterium_HMT_113
- Streptococcus anginosus_constellatus
- Filifactor alocis
- Tannerella forsythia
- Peptostreptococcaceae_[XII][G-5] [Eubacterium]_saphenum
- Porphyromonas endodontalis
- Streptococcus sanguinis
- Neisseria flava_macacae_mucosa_sicca
- Streptococcus gordonii
- Streptococcus parasanguinis_clade_411
- Veillonella dispar
- Pseudomonas azotoformans_lactis_paralactis
- Lautropia mirabilis
- Acinetobacter_Prolinoborus fasciculus_lwoffii
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346
- Olsenella profusa
- Oribacterium sp._HMT_078
- Leptotrichia sp._HMT_498
- Peptostreptococcaceae_[XII][G-9] [Eubacterium]_brachy
- Aridibacter famidurans_nov_96.552%
- Pseudoramibacter alactolyticus
- Fusobacterium sp._HMT_203
- Parvimonas_Peptostreptococcus Candidatus massiliensis_sp._HMT
- Aggregatibacter paraphrophilus
- Enterococcus faecalis
- Janthinobacterium aquaticum_lividum_rivuli
- Parvimonas micra
- Desulfobulbus sp._HMT_041
- Campylobacter gracilis
- Streptococcus cristatus_downii_gwanguense_infantis_infantis_cla .

Species

S076 S077 S078 S079 S080 S081 S082 S083 S084 S085 S086 S087 S088 S089 S090 S096 S107 S108 S109 S110 S111 S112 S113 S114 S115 S116 S117 S118 S119 S120

Samples