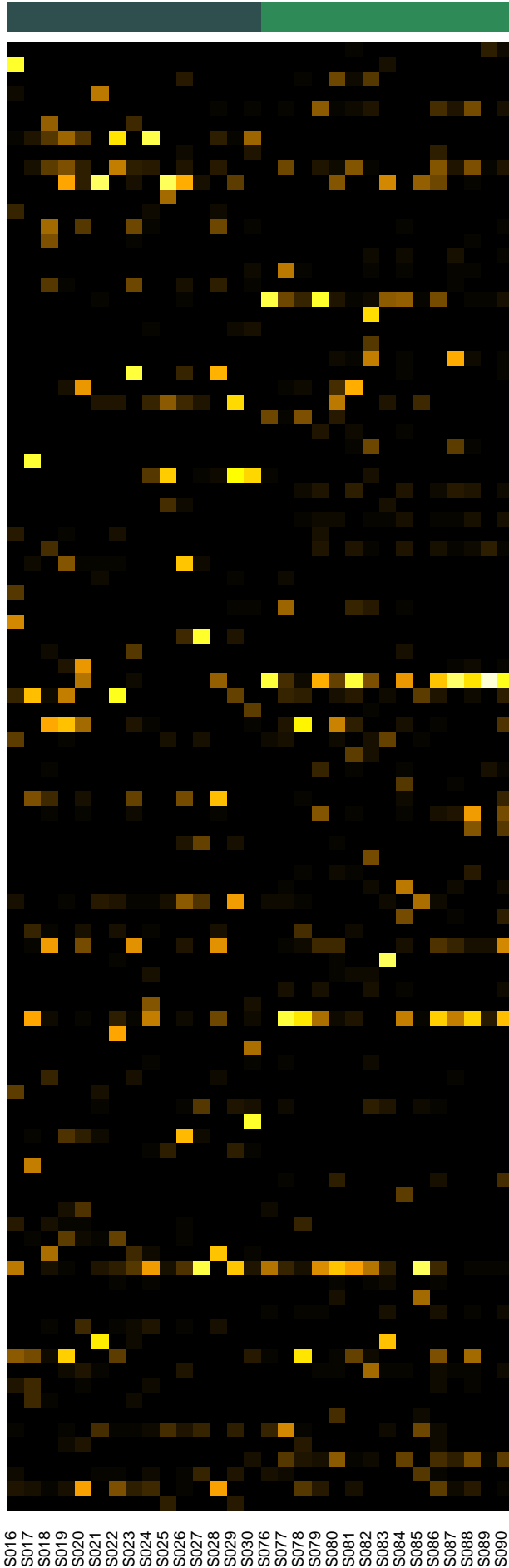




Description2
 PT_NP_HPT
 PT_NP_PP



- Veillonellaceae_[G-1] bacterium_HMT_145
- Neisseria subflava
- Rothia mucilaginosa
- Neisseria oralis
- Peptostreptococcaceae_[XII][G-5] [Eubacterium]_saphenum
- Mycoplasma faucium
- Campylobacter gracilis
- Selenomonas noxia
- Peptostreptococcaceae_[XII][G-9] [Eubacterium]_brachy
- Streptococcus sanguinis
- Abiotrophia defectiva
- Aggregatibacter aphrophilus
- Treponema socranskii
- Bacteroidetes_[G-3] bacterium_HMT_280
- Stomatobaculum sp._HMT_373
- Oribacterium sp._HMT_078
- Lachnospiraceae_[G-8] bacterium_HMT_500
- Enterococcus faecalis
- Olsenella profusa
- Veillonella parvula
- Olsenella sp._HMT_807
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346
- Porphyromonas gingivalis
- Peptoniphilaceae_[G-1] bacterium_HMT_113
- Haemophilus parainfluenzae
- Johnsonella ignava
- Fretibacterium sp._HMT_360
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349
- Pseudomonas koreensis
- Veillonella dispar
- Mogibacterium timidum
- Lautropia mirabilis
- Schaalia odontolytica
- Gemella morbillorum
- Peptostreptococcaceae_[XII][G-4] bacterium_HMT_369
- Pseudomonas helmanticensis
- Streptococcus sinensis
- Aggregatibacter sp._HMT_898
- Streptococcus anginosus
- Aggregatibacter sp._HMT_513
- Haemophilus haemolyticus
- Treponema denticola
- Bacteroidales_[G-2] bacterium_HMT_274
- Desulfobulbus sp._HMT_041
- Streptococcus intermedius
- Dialister invisus
- Fusobacterium sp._HMT_203
- Granulicatella adiacens
- Actinomyces sp._HMT_448
- Fretibacterium fastidiosum
- Pseudoramibacter alactolyticus
- Fusobacterium nucleatum_subsp._vincentii
- Tannerella forsythia
- Porphyromonas endodontalis
- Actinomyces oris
- Limosilactobacillus mucosae
- Schaalia meyeri
- Peptostreptococcus stomatis
- Streptococcus lactarius
- Schaalia cardiffensis
- Fusobacterium nucleatum
- Filifactor alocis
- Aggregatibacter paraphrophilus
- Streptococcus parasanguinis_clade_411
- Johnsonella sp._HMT_166
- Veillonella atypica
- Parvimonas micra
- Methylobacterium radiotolerans
- Prevotella denticola
- Selenomonas sputigena
- Treponema maltophilum
- Cardiobacterium hominis
- Streptococcus gordonii
- Stomatobaculum longum
- Stenotrophomonas pavanii_nov_97.902%
- Veillonella atypica_nov_90.000%
- Pradoshia eiseniae_nov_93.764%
- Schnuerera ultunensis_nov_92.840%
- Pseudacidobacterium ailaui_nov_94.321%
- Fusobacterium nucleatum_nucleatum_subsp._animalis_simiae
- Fusobacterium canifelinum_nucleatum_nucleatum_subsp._polymorphum
- Moraxella catarrhalis_nonliquefaciens
- Campylobacter rectus_showae
- Streptococcus cristatus_downii_gwangjuense_infantis_infantis_clade_578
- Streptococcus salivarius_vestibularis
- Paracoccus aestuarii_beibuensis_hibisci_marinus_pueri
- Aerococcus urinaeequi_viridans
- Fusobacterium nucleatum_nucleatum_subsp._animalis
- Neisseria flava_macacae_mucosa_sicca
- Parvimonas_Peptostreptococcus Candidatus massiliensis_sp._HMT_164
- Staphylococcus capitis_caprae_epidermidis
- Peptoanaerobacter_Peptostreptococcaceae_[XII][G-7] [Eubacterium]_saphenum
- Catonella morbi_sp._HMT_164
- Pseudomonas alcaliphila_chengduensis_oleovorans_toyotomiensis
- Streptococcus cristatus_cristatus_clade_578
- Fusobacterium naviforme_nucleatum_sp._HMT_204
- Streptococcus anginosus_constellatus
- Streptococcus cristatus_cristatus_clade_578_downii_gwangjuense_infantis_infantis_clade_578
- Fusobacterium nucleatum_nucleatum_subsp._vincentii
- Streptococcus multispecies_sppn14_4_nov_90.531%

Species

Samples