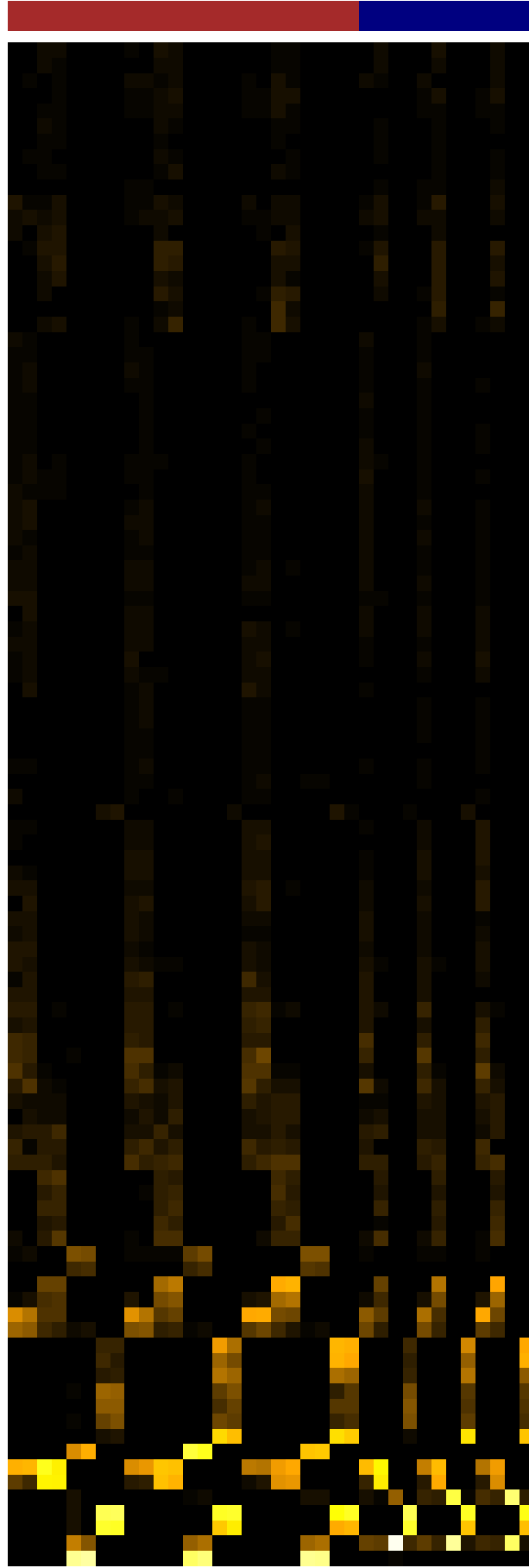




HL
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- Ottowia sp._HMT_894
- Ruminococcaceae_[G-1] bacterium_HMT_075
- Leptotrichia buccalis
- Gemella morbillorum
- Streptococcus intermedius
- Campylobacter showae
- Peptoanaerobacter [Eubacterium] yurii
- Bergeyella sp._HMT_322
- Actinomyces sp._HMT_170
- Actinomyces viscosus
- Capnocytophaga gingivalis
- Prevotella nigrescens
- Leptotrichia hongkongensis
- Neisseria sicca
- Capnocytophaga sputigena
- Leptotrichia sp._HMT_392
- Streptococcus salivarius
- Streptococcus sobrinus
- Actinomyces oris
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348
- Neisseria elongata
- Veillonella sp._HMT_780_nov_97.519%
- Veillonella sp._HMT_780
- Porphyromonas sp._HMT_278
- Capnocytophaga endodontalis
- Prevotella oulorum
- Selenomonas artemidis
- Granulicatella elegans
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346
- Alloprevotella tannerae
- Saccharibacteria_(TM7)_[G-6] bacterium_HMT_870_nov_97.093%
- Campylobacter concisus
- Leptotrichia sp._HMT_215
- Kingella oralis
- Neisseria cinerea
- Prevotella intermedia
- Capnocytophaga granulosa
- Gemella haemolysans
- Lachnoanaerobaculum saburreum
- Streptococcus cristatus
- Prevotella loescheii
- Streptococcus chosunense
- Actinomyces sp._HMT_897
- Streptococcus oralis_subsp_tigurinus_clade_070
- Actinomyces sp._HMT_414
- Actinomyces gerencseriae
- Haemophilus haemolyticus
- Streptococcus mutans
- Actinomyces naeslundii
- Escherichia coli
- Peptidiphaga sp._HMT_183
- Schaalia sp._HMT_178
- Actinomyces dentalis
- Actinomyces massiliensis
- Actinomyces sp._HMT_175
- Actinomyces sp._HMT_175_nov_97.830%
- Haemophilus parainfluenzae
- Porphyromonas pasteri
- Schaalia meyeri
- Campylobacter gracilis
- Actinomyces sp._HMT_448
- Leptotrichia sp._HMT_221
- Leptotrichia wadei
- Streptococcus sp._HMT_064
- Alloprevotella sp._HMT_473
- Corynebacterium matruchotii
- Streptococcus oralis
- Rothia dentocariosa
- Schaalia sp._HMT_180
- Actinomyces sp._HMT_169
- Prevotella melaninogenica
- Granulicatella adiacens
- Streptococcus gordonii
- Lautropia mirabilis
- Leptotrichia sp._HMT_212
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957
- Abiotrophia defectiva
- Capnocytophaga leadbetteri
- Prevotella oris
- Neisseria subflava
- Schaalia odontolytica
- Streptococcus sanguinis
- Streptococcus mitis
- Fusobacterium nucleatum
- Listeria monocytogenes
- Limosilactobacillus fermentum
- Enterococcus faecalis
- Shigella sonnei
- Pseudomonas aeruginosa
- Escherichia fergusonii
- Staphylococcus argenteus_aureus_roterodami
- Trueperia radiovictrix
- Veillonella dispar_parvula
- Veillonella parvula
- Allobacillus halotolerans
- Bacillus halotolerans
- Salmonella enterica
- Frondbacter mangrovi_nov_92.771%
- Porphyromonas gingivalis

Species



F-12829_S02
 F-12829_S03
 F-12829_S05
 F-12829_S06
 F-12829_S08
 F-12829_S09
 F-12829_S11
 F-12829_S12
 F-12829_S14
 F-12829_S15
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 F-12829_S01
 F-12829_S04
 F-12829_S07
 F-12829_S10
 F-12829_S13
 F-12829_S16
 F-12829_S19
 F-12829_S22
 F-12829_S25
 F-12829_S28
 F-12829_S31
 F-12829_S34

Samples