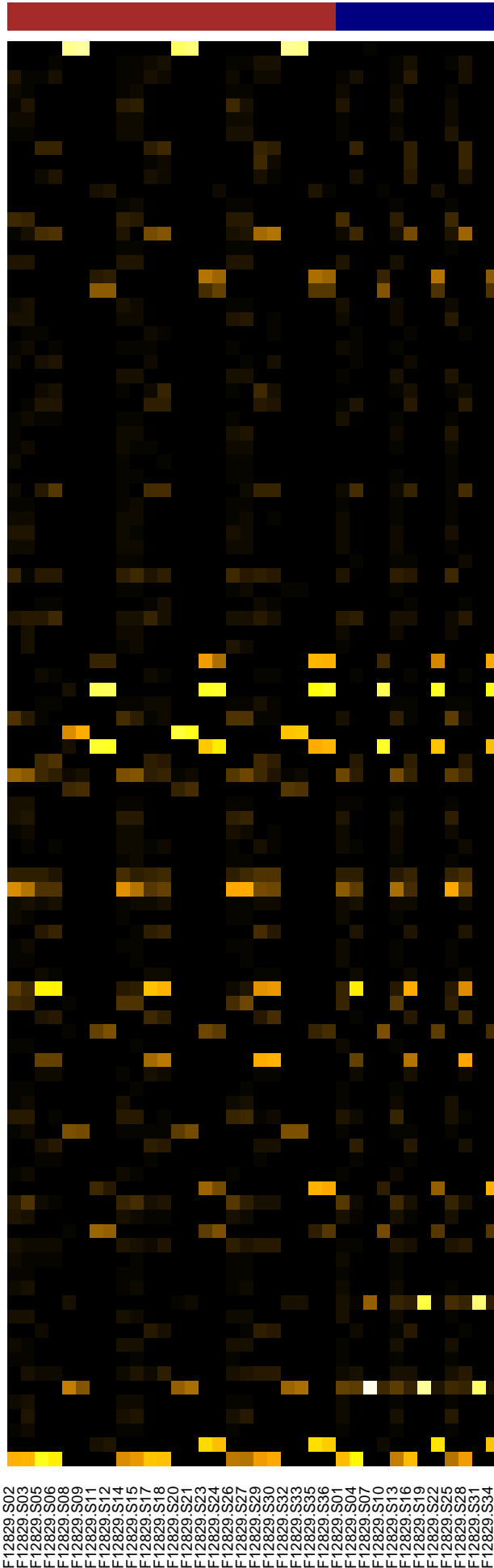




HL  
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- Porphyromonas gingivalis
- Gemella morbillorum
- Capnocytophaga gingivalis
- Campylobacter concisus
- Actinomyces sp.\_HMT\_448
- Lachnoanaerobaculum saburreum
- Peptidiphaga sp.\_HMT\_183
- Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_957
- Streptococcus sobrinus
- Leptotrichia sp.\_HMT\_392
- Escherichia coli
- Streptococcus oralis\_subsp.\_tigurinus\_clade\_070
- Alloprevotella sp.\_HMT\_473
- Streptococcus sanguinis
- Actinomyces gerencseriae
- Leptotrichia sp.\_HMT\_221
- Enterococcus faecalis
- Pseudomonas aeruginosa
- Porphyromonas pasteri
- Actinomyces sp.\_HMT\_175
- Bergeyella sp.\_HMT\_322
- Selenomonas artemidis
- Leptotrichia hongkongensis
- Actinomyces dentalis
- Actinomyces oris
- Neisseria sicca
- Granulicatella elegans
- Schaalia sp.\_HMT\_178
- Prevotella loescheii
- Actinomyces naeslundii
- Actinomyces sp.\_HMT\_414
- Capnocytophaga leadbetteri
- Haemophilus haemolyticus
- Kingella oralis
- Schaalia meyeri
- Neisseria cinerea
- Actinomyces viscosus
- Granulicatella adiacens
- Streptococcus mutans
- Actinomyces sp.\_HMT\_170
- Prevotella melaninogenica
- Capnocytophaga granulosa
- Streptococcus chosunense
- Listeria monocytogenes
- Campylobacter showae
- Bacillus halotolerans
- Streptococcus intermedius
- Streptococcus oralis
- Truepera radiovictrix
- Salmonella enterica
- Lautropia mirabilis
- Fusobacterium nucleatum
- Neisseria subflava
- Prevotella intermedia
- Streptococcus sp.\_HMT\_064
- Gemella haemolysans
- Leptotrichia buccalis
- Actinomyces sp.\_HMT\_897
- Streptococcus gordonii
- Streptococcus mitis
- Prevotella nigrescens
- Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_348
- Leptotrichia sp.\_HMT\_212
- Leptotrichia sp.\_HMT\_215
- Prevotella oulorum
- Ruminococcaceae\_[G-1] bacterium\_HMT\_075
- Veillonella parvula
- Corynebacterium matruchotii
- Abiotrophia defectiva
- Escherichia fergusonii
- Veillonella sp.\_HMT\_780
- Schaalia odontolytica
- Ottowia sp.\_HMT\_894
- Porphyromonas sp.\_HMT\_278
- Streptococcus cristatus
- Leptotrichia wadei
- Prevotella oris
- Capnocytophaga sputigena
- Peptoanaerobacter [Eubacterium] yurii
- Neisseria elongata
- Limosilactobacillus fermentum
- Rothia dentocariosa
- Campylobacter gracilis
- Shigella sonnei
- Schaalia sp.\_HMT\_180
- Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_346
- Aggregatibacter segnis
- Alloprevotella tanneriae
- Allobacillus halotolerans
- Haemophilus parainfluenzae
- Streptococcus salivarius
- Actinomyces massiliensis
- Capnocytophaga endodontalis
- Actinomyces sp.\_HMT\_169
- Frondebacter mangrovi\_nov\_92.771%
- Saccharibacteria\_(TM7)\_[G-6] bacterium\_HMT\_870\_nov\_97.093%
- Actinomyces sp.\_HMT\_175\_nov\_97.830%
- Veillonella sp.\_HMT\_780\_nov\_97.519%
- Staphylococcus argenteus\_aureus\_roterodami
- Veillonella dispar\_parvula

Species

F-12829\_S02  
 F-12829\_S03  
 F-12829\_S05  
 F-12829\_S06  
 F-12829\_S08  
 F-12829\_S09  
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 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