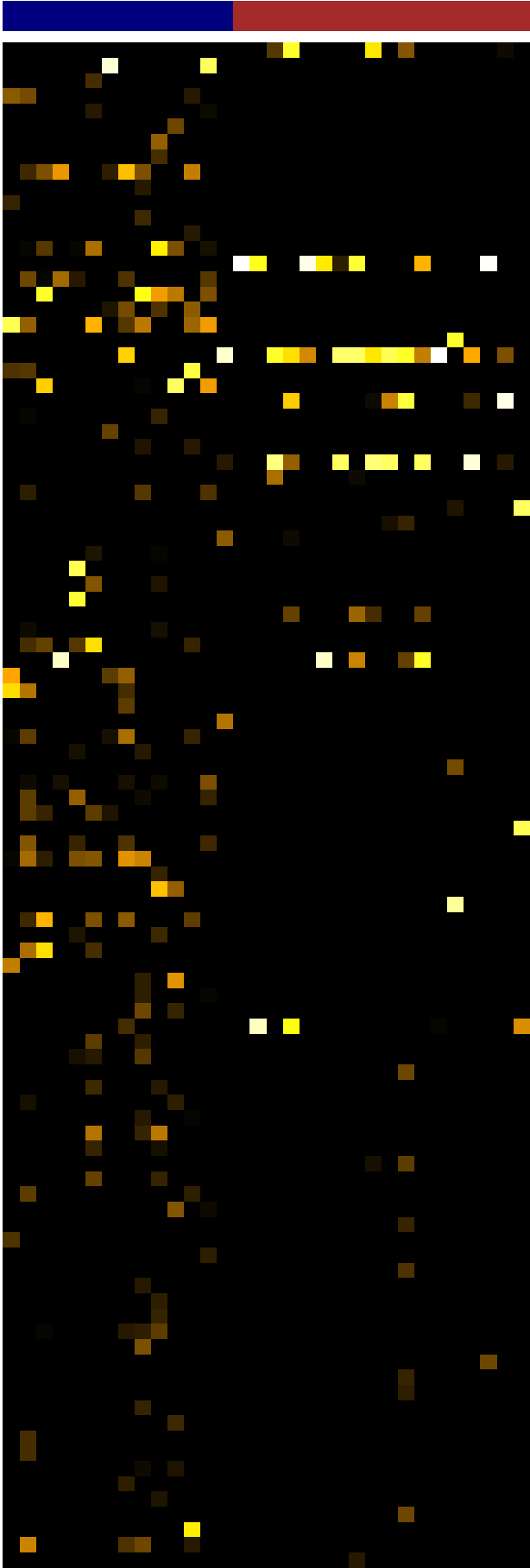




Group
Oral
Vaginal



- Lachnospiraceae_[G-9] bacterium_HMT_924
- Lactobacillus fermentum
- Actinomyces sp._HMT_169
- Haemophilus parainfluenzae
- Bacteroidales_[G-2] sp._Oral_Taxon_274
- Selenomonas sp._HMT_134
- Actinomyces sp._HMT_175
- Capnocytophaga granulosa
- Streptococcus sp._str._C300
- Capnocytophaga gingivalis
- Haemophilus haemolyticus
- Selenomonas sp._Oral_Taxon_E83
- Lautropia mirabilis
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348
- Lactobacillus jensenii
- Porphyromonas endodontalis
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346
- Gemella haemolysans
- Streptococcus mitis
- Corynebacterium mucifaciens
- Gardnerella vaginalis
- Streptococcus chosunense
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349
- Gardnerella swidsinskii
- Bergeyella sp._HMT_900
- Streptococcus sp._HMT_064
- Capnocytophaga leadbetteri
- Lactobacillus iners
- Gardnerella leopoldii
- Gemella moribillum
- Enterococcus faecalis
- Prevotella bivia
- Aerococcus christensenii_Oral_Taxon_D44
- Prevotella micans
- Streptococcus salivarius
- Cardiobacterium hominis
- Lactobacillus salivarius
- Limosilactobacillus coleohominis
- Parvimonas sp._HMT_110
- Streptococcus sanguinis
- Lactobacillus gasseri
- Veillonella parvula
- Granulicatella elegans
- Granulicatella adiacens
- Gemella sp._Oral_Taxon_B93
- Saccharibacteria_(TM7)_[G-6] bacterium_HMT_870
- Parvimonas sp._Oral_Taxon_110
- Staphylococcus epidermidis
- Parvimonas micra
- Streptococcus oralis_subsp._tigurinus_clade_070
- Capnocytophaga sputigena
- Enterococcus sp._str._F95
- Streptococcus sp._Oral_Taxon_71
- Streptococcus cristatus
- Ruminococcaceae_[G-1] bacterium_HMT_075
- Selenomonas sputigena
- Corynebacterium appendicis
- Streptococcus oralis
- Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356
- Saccharibacteria_(TM7)_[G-8] bacterium_HMT_955
- Veillonella sp._HMT_780
- Selenomonas sp._Oral_Taxon_137
- Lachnospiraceae_[G] sp._Oral_Taxon_B32
- Selenomonas sp._HMT_137
- Lactobacillus crispatus
- Tannerella sp._HMT_286
- Gemella morbillorum
- Sneathia amnii_[Not_Validly_Published]
- Actinomyces naeslundii
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957
- Streptococcus oralis_subsp._dentisani_clade_058
- Porphyromonas pasteri
- Actinomyces johnsonii
- Megasphaera genomosp.
- Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351
- Streptococcus oralis_subsp._tigurinus_clade_071
- Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345
- Sneathia sanguinegens
- Granulicatella paradiacens
- Fretibacterium sp._HMT_359
- Mycoplasma hominis
- Selenomonas sp._HMT_133
- Capnocytophaga ochracea
- Actinomyces oris
- Campylobacter gracilis
- Selenomonas sp._HMT_133_nov_96.947%
- Anaerococcus lactolyticus_nov_97.271%
- Gordonibacter pamelaee_nov_88.867%
- Prevotella colorans_nov_96.212%
- Streptococcus cristatus_nov_97.363%
- Selenomonas sp._Oral_Taxon_137_nov_97.786%
- Granulicatella elegans_nov_97.180%
- Streptobacillus notomitis_nov_93.555%
- Capnocytophaga gingivalis_nov_97.471%
- Parvimonas sp._Oral_Taxon_110_nov_97.466%
- Actinomyces sp._HMT_448_nov_96.346%
- Parvimonas micra_nov_91.456%
- Streptococcus pneumoniae_sp._str._C300_sp._str._M334
- Streptococcus pneumoniae_sp._str._M334
- Staphylococcus argenteus_aureus

Species

- F6952.S01
- F6952.S02
- F6952.S03
- F6952.S04
- F6952.S05
- F6952.S06
- F6952.S07
- F6952.S08
- F6952.S09
- F6952.S10
- F6952.S11
- F6952.S12
- F6952.S13
- F6952.S14
- F6952.S15
- F6952.S16
- F6952.S17
- F6952.S18
- F6952.S19
- F6952.S20
- F6952.S21
- F6952.S22
- F6952.S23
- F6952.S24
- F6952.S25
- F6952.S26
- F6952.S27
- F6952.S28
- F6952.S29
- F6952.S30
- F6952.S31
- F6952.S32

Samples