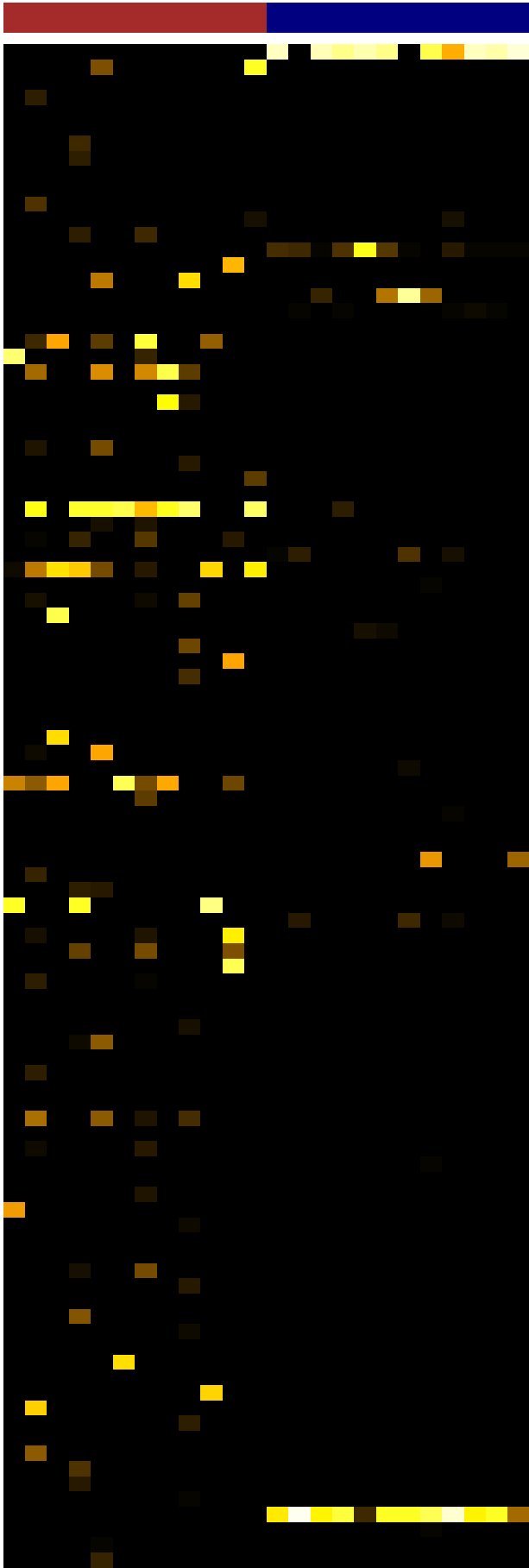




Group1  
All Ligature  
All Brain



- Enterococcus faecalis
- Staphylococcus hominis
- Klebsiella aerogenes
- Veillonella sp.\_HMT\_780
- Parasutterella excrementihominis
- Corynebacterium coyleae
- Glutamicibacter soli
- Tepidiphilus succinatimandens
- Cutibacterium avidum
- Corynebacterium matruchotii
- Neisseria mucosa
- Enterobacter hormaechei
- Bradyrhizobium valentinum
- Ligilactobacillus murinus
- Janibacter cremeus
- Staphylococcus caprae
- Klebsiella pneumoniae
- Streptococcus danieliae
- Varibaculum cambriense
- Staphylococcus capitis
- Corynebacterium tuberculostearicum
- Fusobacterium nucleatum
- Facklamia languida
- Rothia mucilaginosa
- Actinomyces sp.\_HMT\_448
- Pantoea allii
- Streptococcus parasanguinis\_clade\_411
- Streptococcus sp.\_HMT\_066
- Methylophilus leisingeri
- Neisseriaceae\_[G-1]\_bacterium\_HMT\_174
- Cutibacterium acnes
- Weizmannia ginsengihumi
- Bradyrhizobium lupini
- Enterococcus gallinarum
- Acinetobacter lwoffii
- Enterobacter cloacae
- Streptococcus mitis
- Corynebacterium durum
- Bifidobacterium pseudolongum
- Cutibacterium namnetense
- Microbacterium maritipicum
- Corynebacterium ureicelerivorans
- Corynebacterium aurimucosum
- Jeotgalicoccus halotolerans
- Corynebacterium jeikeium
- Triticum aestivum
- Bifidobacterium longum
- Faecalibaculum rodentium
- Bacillus subtilis
- Corynebacterium afermentans
- Salmonella enterica
- Peribacillus frigoritolerans
- Rhodococcus qingshengii
- Streptococcus thoraltensis
- Streptococcus lactarius
- Acinetobacter pittii
- Actinidia eriantha
- Enterococcus casseliflavus
- Staphylococcus warneri
- Bradyrhizobium japonicum
- Staphylococcus pasteurii
- Acinetobacter johnsonii
- Enterobacter mori
- Tannerella serpentiformis
- Mammaliicoccus sciuri
- Shigella flexneri
- Corynebacterium kroppenstedtii
- Corynebacterium pilbarensis
- Mammaliicoccus lentus
- Mediterraneibacter faecis
- Escherichia fergusonii
- Bifidobacterium bifidum
- Streptococcus chosunense
- Enterobacter kobei
- Massilia arenae
- Lawsonella clevelandensis
- Sphingomonas leidyi
- Streptococcus oralis
- Streptococcus mutans
- Acinetobacter septicus
- Bacillus halotolerans
- Abiotrophia defectiva
- Staphylococcus saprophyticus
- Pseudoramibacter alactolyticus
- Corynebacterium pseudodiphtheriticum
- Corynebacterium falsenii
- Lactiplantibacillus plantarum
- Prevotella sp.\_HMT\_314
- Daejeonella oryzae\_nov\_85.551%
- Thermodesulfobium acidiphilum\_nov\_81.729%
- Sporosarcina sp.\_MOT-205\_nov\_96.360%
- Streptococcus danieliae\_nov\_95.594%
- Clostridium saccharoperbutylacetonicum\_nov\_95.800%
- Microbacterium saccharophilum\_nov\_76.981%
- Ralstonia solanacearum\_nov\_95.922%
- Frankia discariae\_nov\_81.481%
- Staphylococcus argenteus\_aureus\_roterodami
- Enterobacter\_Leclercia adacarboxylata\_cloacae
- Terrimonas multispecies\_sppn3\_2\_nov\_90.822%
- Gloeobacter multispecies\_sppn4\_2\_nov\_84.254%

Species

Samples