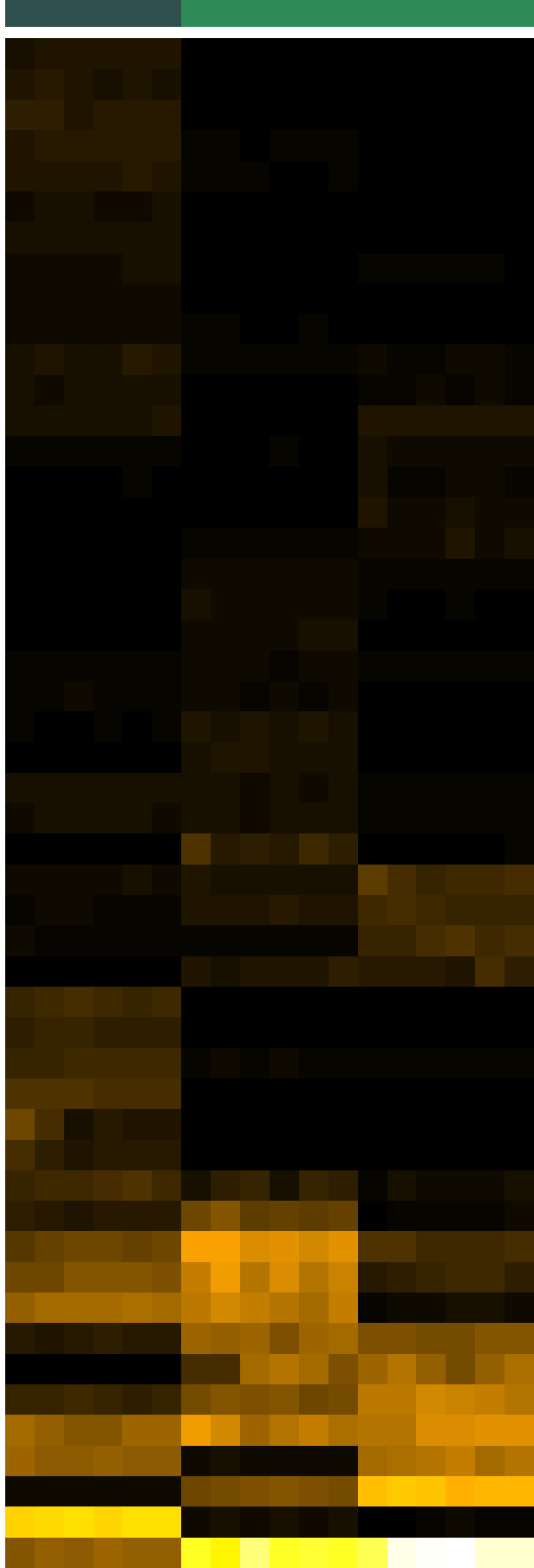


Group002
 Saliva P2 -PMA
 Baseline P2 -PMA



- Oribacterium sinus
- Prevotella histicola
- Prevotella jejunii
- Streptococcus mitis
- Streptococcus infantis_infantis_clade_638
- Streptococcus infantis_clade_431
- Alloprevotella sp._HMT_473_nov_97.347%
- Streptococcus australis
- Streptococcus sp._HMT_066
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_352
- Leptotrichia sp._HMT_215
- Streptococcus parasanguinis_clade_411
- Granulicatella adiacens
- Streptococcus sanguinis
- Fusobacterium nucleatum
- Streptococcus gordonii
- Eikenella corrodens
- Streptococcus oralis
- Neisseria mucosa
- Gemella haemolysans
- Gemella sanguinis
- Veillonella tobetsuensis
- Haemophilus sputorum
- Campylobacter concisus_nov_96.963%
- Veillonella dispar
- Veillonella denticariosi_dispar_parvula
- Neisseria elongata
- Streptococcus sp._HMT_423
- Veillonella parvula
- Streptococcus parasanguinis_parasanguinis_clade_721
- Neisseria subflava
- Alloprevotella sp._HMT_308
- Prevotella nanceiensis
- Schaalia odontolytica
- Prevotella pallens
- Lachnospiraceae_[G-2] bacterium_HMT_096
- Leptotrichia sp._HMT_417
- Rothia mucilaginos
- Campylobacter concisus
- Neisseria perflava
- Veillonella rogosae
- Porphyromonas pasteri
- Neisseria flavescens
- Neisseria flava
- Veillonella atypica
- Fusobacterium periodonticum
- Streptococcus salivarius
- Veillonella dispar_parvula
- Prevotella melaninogenica
- Haemophilus parainfluenzae

Species

- F28914.S018
- F28914.S017
- F28914.S013
- F28914.S014
- F28914.S016
- F28914.S015
- F28914.S064
- F28914.S061
- F28914.S066
- F28914.S063
- F28914.S065
- F28914.S062
- F28914.S053
- F28914.S049
- F28914.S051
- F28914.S052
- F28914.S050
- F28914.S054

Samples