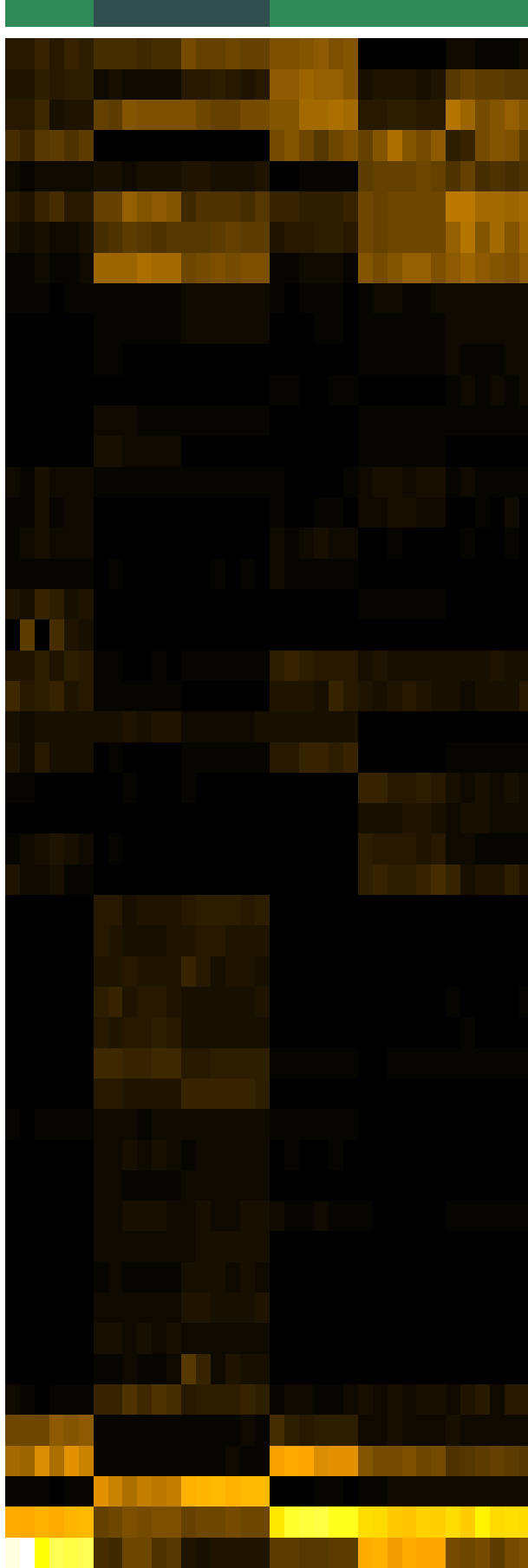


Group003
 Saliva -PMA
 Baseline -PMA



- Streptococcus salivarius
- Veillonella atypica
- Fusobacterium periodonticum
- Neisseria flava
- Campylobacter concisus
- Neisseria perflava
- Veillonella rogosae
- Porphyromonas pasteri
- Veillonella denticariosi_dispar_parvula
- Veillonella dispar
- Gemella haemolysans
- Streptococcus oralis
- Veillonella tobetsuensis
- Alloprevotella sp._HMT_473
- Gemella sanguinis
- Aggregatibacter segnis
- Eikenella corrodens
- Streptococcus sanguinis
- Aggregatibacter aphrophilus
- Citrobacter koseri
- Veillonella parvula
- Neisseria subflava
- Granulicatella adiacens
- Streptococcus parasanguinis_parasanguinis_clade_721
- Haemophilus sputorum
- Campylobacter concisus_nov_96.963%
- Neisseria mucosa
- Neisseria elongata
- Alloprevotella sp._HMT_308
- Prevotella nanceiensis
- Leptotrichia sp._HMT_417
- Streptococcus mitis
- Streptococcus infantis_infantis_clade_638
- Schaalia odontolytica
- Prevotella pallens
- Streptococcus parasanguinis_clade_411
- Streptococcus australis
- Streptococcus infantis_clade_431
- Leptotrichia sp._HMT_215
- Oribacterium sinus
- Prevotella histicola
- Prevotella jejuni
- Alloprevotella sp._HMT_473_nov_97.347%
- Lachnospiraceae_[G-2]_bacterium_HMT_096
- Rothia mucilaginosa
- Streptococcus sp._HMT_423
- Veillonella dispar_parvula
- Prevotella melaninogenica
- Haemophilus parainfluenzae
- Neisseria flavescens

Species

- F28914_S028
- F28914_S027
- F28914_S029
- F28914_S025
- F28914_S030
- F28914_S026
- F28914_S004
- F28914_S006
- F28914_S001
- F28914_S005
- F28914_S003
- F28914_S019
- F28914_S017
- F28914_S013
- F28914_S014
- F28914_S016
- F28914_S015
- F28914_S053
- F28914_S049
- F28914_S051
- F28914_S052
- F28914_S050
- F28914_S054
- F28914_S036
- F28914_S040
- F28914_S041
- F28914_S038
- F28914_S037
- F28914_S064
- F28914_S061
- F28914_S066
- F28914_S063
- F28914_S065
- F28914_S062

Samples