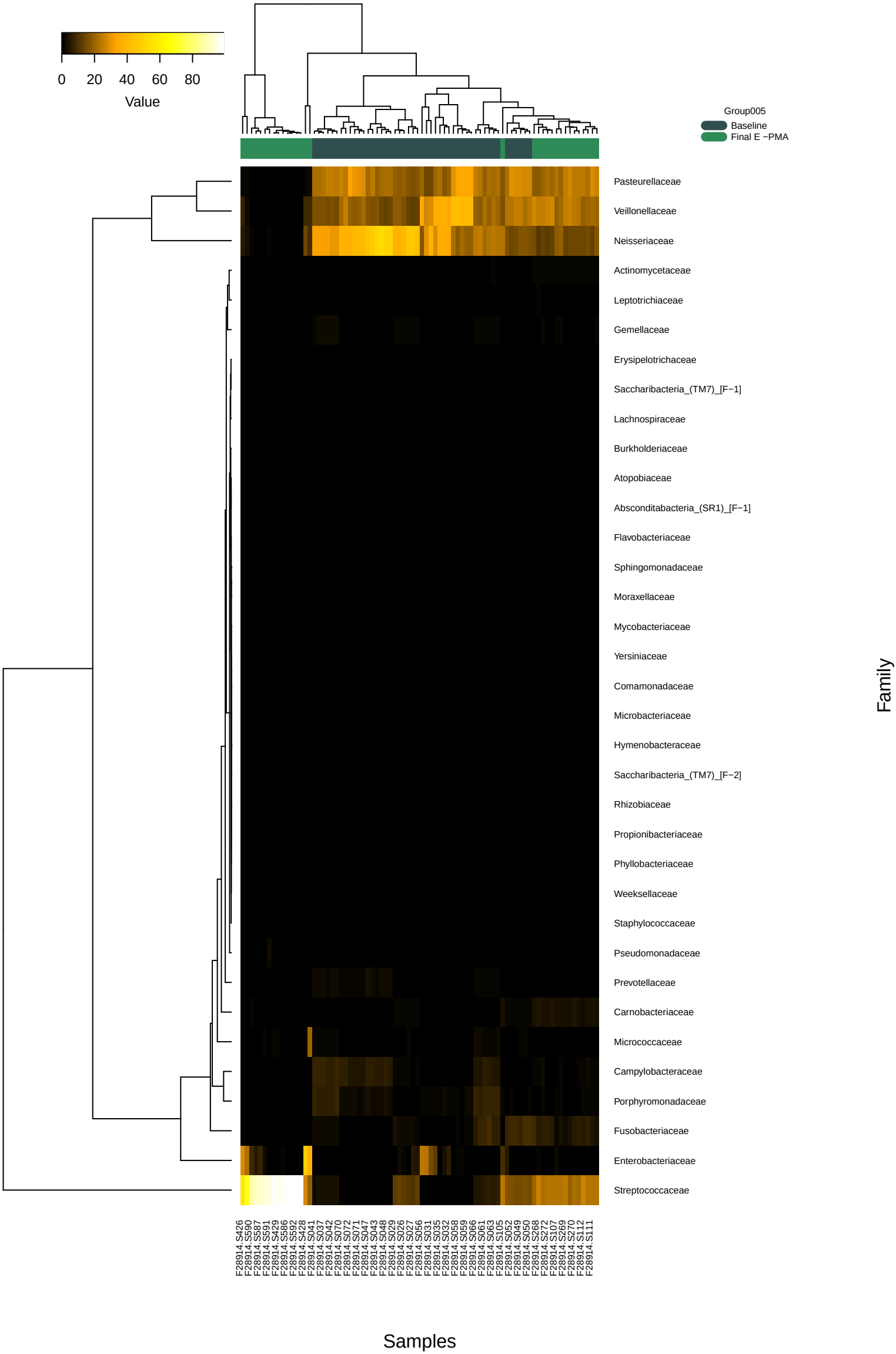
0
20
40
60
80
Value
Group005
Baseline
Final E −PMA
Pasteurellaceae
Veillonellaceae
Neisseriaceae
Actinomycetaceae
Leptotrichiaceae
Gemellaceae
Erysipelotrichaceae
Saccharibacteria_(TM7)_[F−1]
Lachnospiraceae
Burkholderiaceae
Atopobiaceae
Absconditabacteria_(SR1)_[F−1]
Flavobacteriaceae
Sphingomonadaceae
Moraxellaceae
Mycobacteriaceae
Yersiniaceae
Comamonadaceae
Microbacteriaceae
Hymenobacteraceae
Saccharibacteria_(TM7)_[F−2]
Rhizobiaceae
Propionibacteriaceae
Phyllobacteriaceae
Weeksellaceae
Staphylococcaceae
Pseudomonadaceae
Prevotellaceae
Carnobacteriaceae
Micrococcaceae
Campylobacteraceae
Porphyromonadaceae
Fusobacteriaceae
Enterobacteriaceae
Streptococcaceae
Family
F28914.S426
F28914.S590
F28914.S587
F28914.S591
F28914.S429
F28914.S586
F28914.S592
F28914.S428
F28914.S041
F28914.S037
F28914.S042
F28914.S070
F28914.S072
F28914.S071
F28914.S047
F28914.S043
F28914.S048
F28914.S029
F28914.S026
F28914.S027
F28914.S056
F28914.S031
F28914.S035
F28914.S032
F28914.S058
F28914.S059
F28914.S066
F28914.S061
F28914.S063
F28914.S105
F28914.S052
F28914.S049
F28914.S050
F28914.S268
F28914.S272
F28914.S107
F28914.S269
F28914.S270
F28914.S112
F28914.S111
Samples