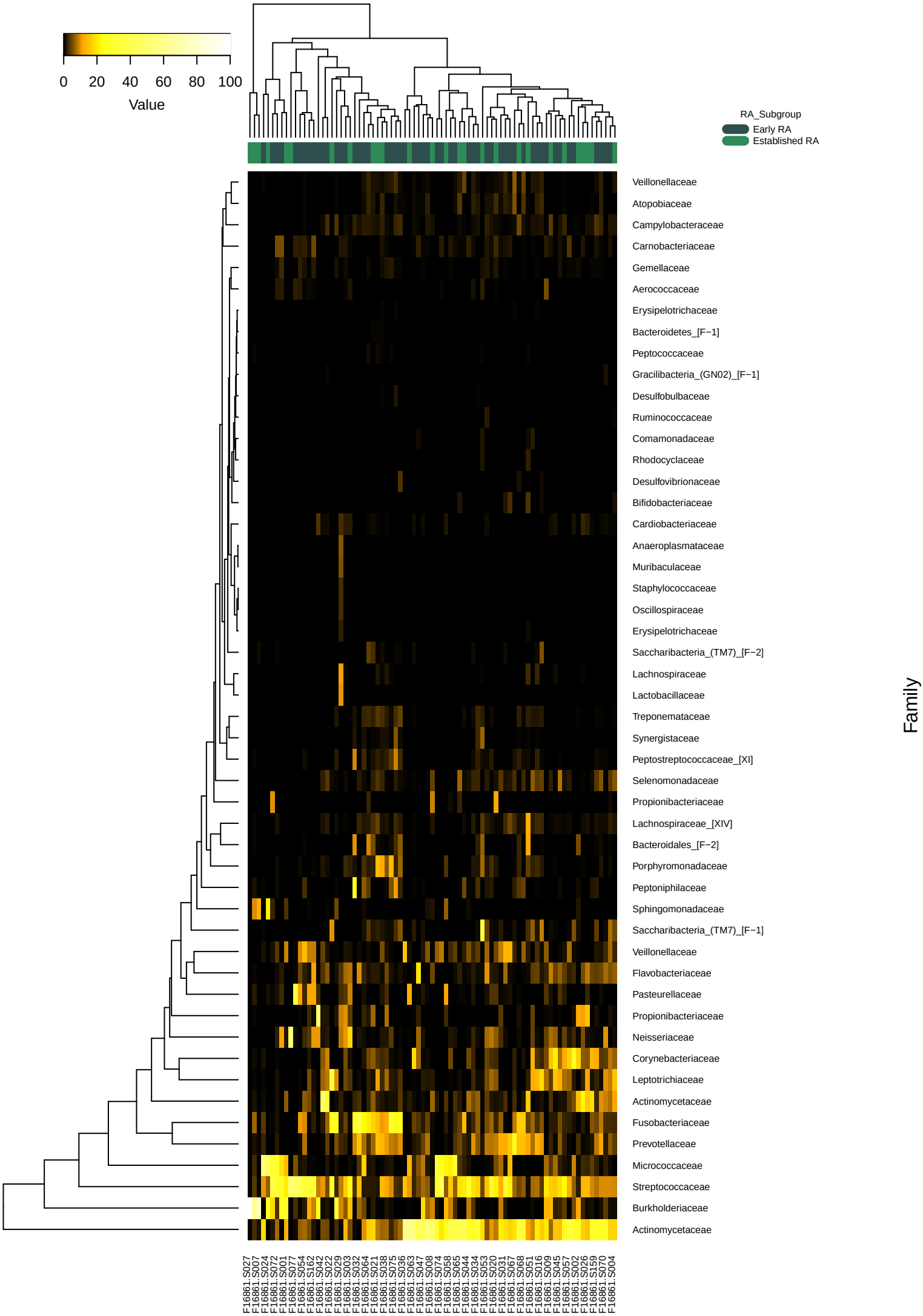
0
20
40
60
80
100
Value
RA_Subgroup
Early RA
Established RA
Veillonellaceae
Atopobiaceae
Campylobacteraceae
Carnobacteriaceae
Gemellaceae
Aerococcaceae
Erysipelotrichaceae
Bacteroidetes_[F−1]
Peptococcaceae
Gracilibacteria_(GN02)_[F−1]
Desulfobulbaceae
Ruminococcaceae
Comamonadaceae
Rhodocyclaceae
Desulfovibrionaceae
Bifidobacteriaceae
Cardiobacteriaceae
Anaeroplasmataceae
Muribaculaceae
Staphylococcaceae
Oscillospiraceae
Erysipelotrichaceae
Saccharibacteria_(TM7)_[F−2]
Lachnospiraceae
Lactobacillaceae
Treponemataceae
Synergistaceae
Peptostreptococcaceae_[XI]
Selenomonadaceae
Propionibacteriaceae
Lachnospiraceae_[XIV]
Bacteroidales_[F−2]
Porphyromonadaceae
Peptoniphilaceae
Sphingomonadaceae
Saccharibacteria_(TM7)_[F−1]
Veillonellaceae
Flavobacteriaceae
Pasteurellaceae
Propionibacteriaceae
Neisseriaceae
Corynebacteriaceae
Leptotrichiaceae
Actinomycetaceae
Fusobacteriaceae
Prevotellaceae
Micrococcaceae
Streptococcaceae
Burkholderiaceae
Actinomycetaceae
Family
F16861.S027
F16861.S007
F16861.S024
F16861.S072
F16861.S001
F16861.S077
F16861.S054
F16861.S162
F16861.S042
F16861.S022
F16861.S029
F16861.S003
F16861.S032
F16861.S064
F16861.S021
F16861.S038
F16861.S075
F16861.S036
F16861.S063
F16861.S047
F16861.S008
F16861.S074
F16861.S058
F16861.S065
F16861.S044
F16861.S034
F16861.S053
F16861.S020
F16861.S031
F16861.S067
F16861.S068
F16861.S051
F16861.S016
F16861.S009
F16861.S045
F16861.S057
F16861.S002
F16861.S026
F16861.S159
F16861.S070
F16861.S004
Samples