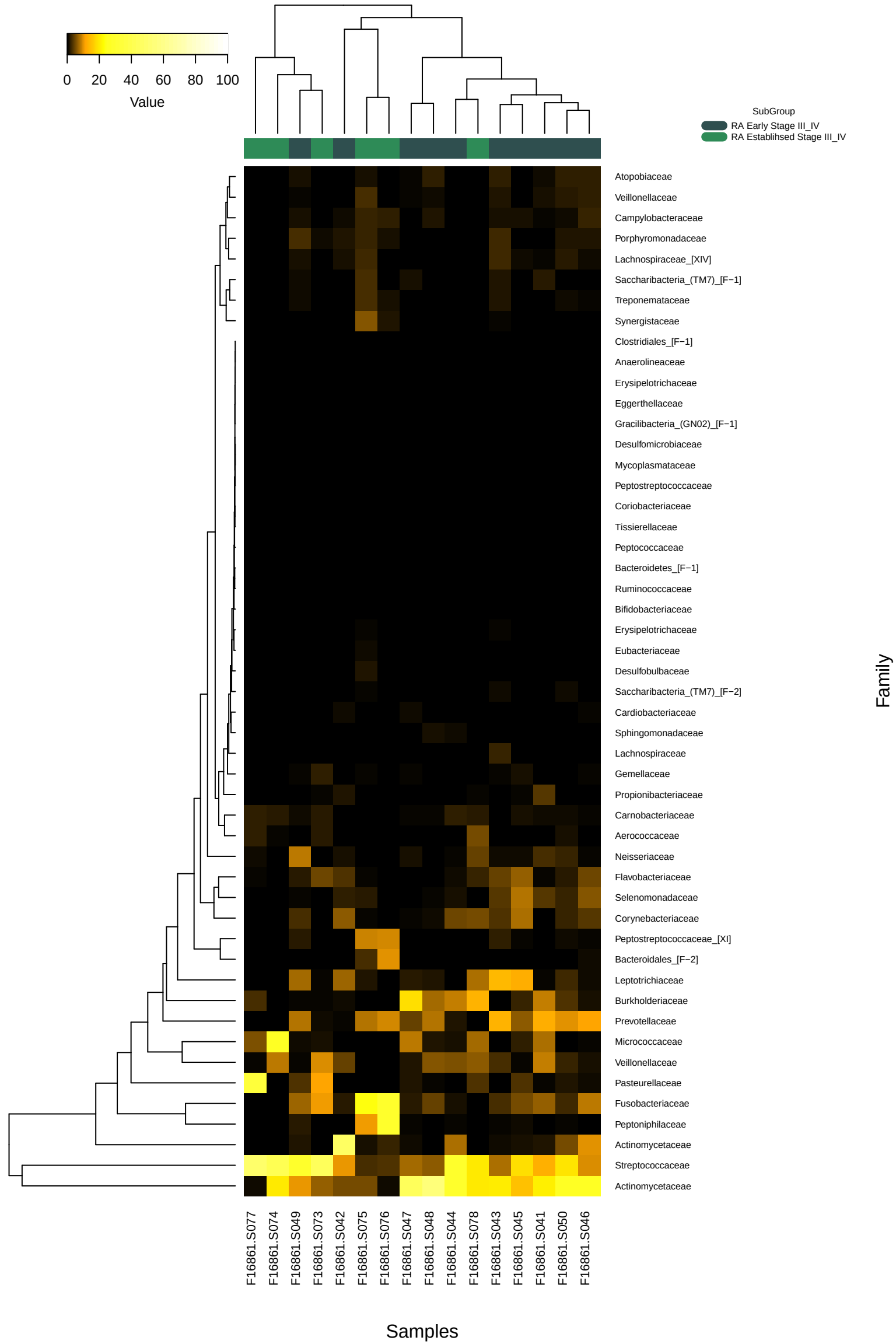
0
20
40
60
80
100
Value
SubGroup
RA Early Stage III_IV
RA Establihsed Stage III_IV
Atopobiaceae
Veillonellaceae
Campylobacteraceae
Porphyromonadaceae
Lachnospiraceae_[XIV]
Saccharibacteria_(TM7)_[F−1]
Treponemataceae
Synergistaceae
Clostridiales_[F−1]
Anaerolineaceae
Erysipelotrichaceae
Eggerthellaceae
Gracilibacteria_(GN02)_[F−1]
Desulfomicrobiaceae
Mycoplasmataceae
Peptostreptococcaceae
Coriobacteriaceae
Tissierellaceae
Peptococcaceae
Bacteroidetes_[F−1]
Ruminococcaceae
Bifidobacteriaceae
Erysipelotrichaceae
Eubacteriaceae
Desulfobulbaceae
Saccharibacteria_(TM7)_[F−2]
Cardiobacteriaceae
Sphingomonadaceae
Lachnospiraceae
Gemellaceae
Propionibacteriaceae
Carnobacteriaceae
Aerococcaceae
Neisseriaceae
Flavobacteriaceae
Selenomonadaceae
Corynebacteriaceae
Peptostreptococcaceae_[XI]
Bacteroidales_[F−2]
Leptotrichiaceae
Burkholderiaceae
Prevotellaceae
Micrococcaceae
Veillonellaceae
Pasteurellaceae
Fusobacteriaceae
Peptoniphilaceae
Actinomycetaceae
Streptococcaceae
Actinomycetaceae
Family
F16861.S077
F16861.S074
F16861.S049
F16861.S073
F16861.S042
F16861.S075
F16861.S076
F16861.S047
F16861.S048
F16861.S044
F16861.S078
F16861.S043
F16861.S045
F16861.S041
F16861.S050
F16861.S046
Samples