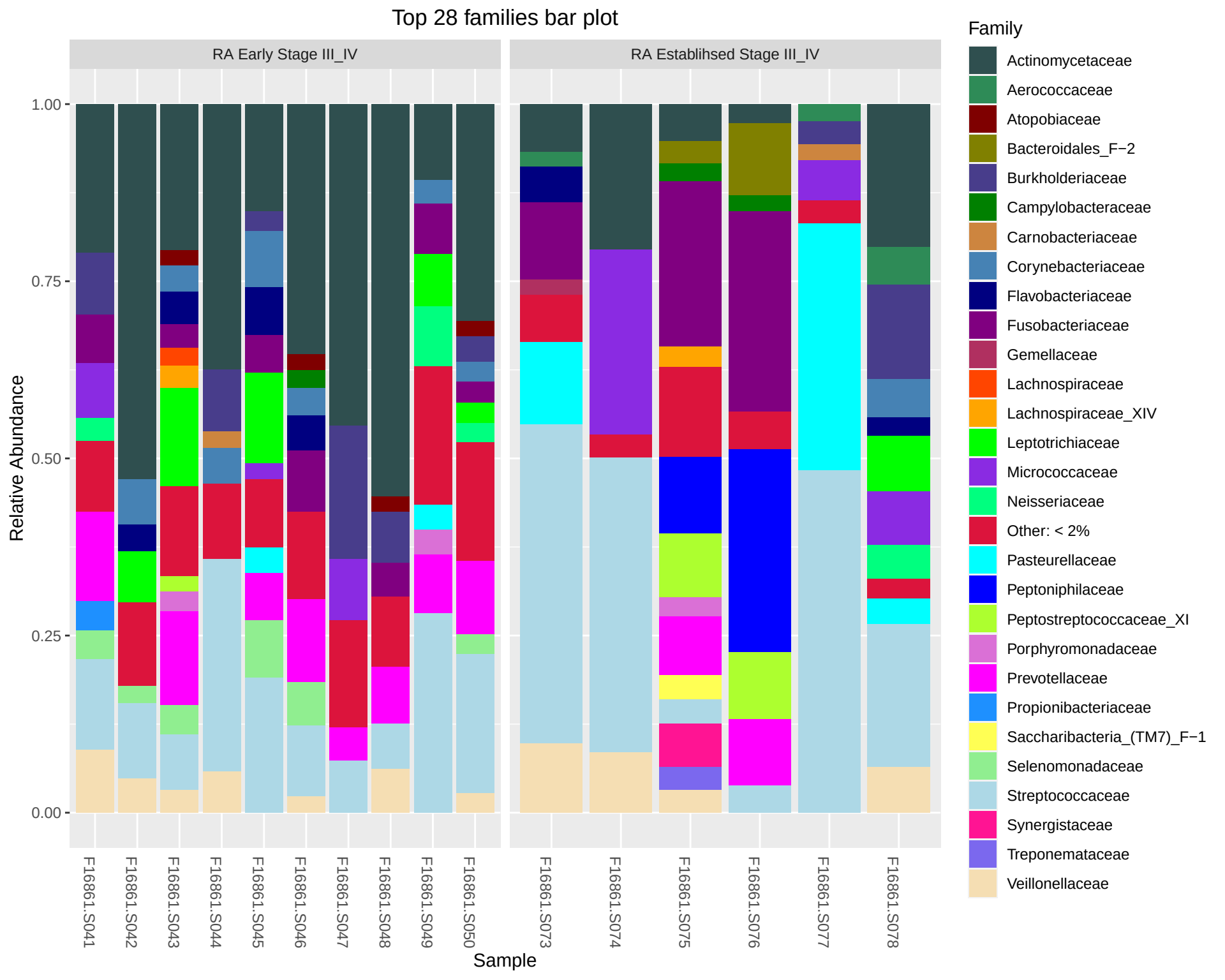
Top 28 families bar plot
RA Early Stage III_IV
RA Establihsed Stage III_IV
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F16861.S041
F16861.S042
F16861.S043
F16861.S044
F16861.S045
F16861.S046
F16861.S047
F16861.S048
F16861.S049
F16861.S050
F16861.S073
F16861.S074
F16861.S075
F16861.S076
F16861.S077
F16861.S078
Sample
Family
Actinomycetaceae
Aerococcaceae
Atopobiaceae
Bacteroidales_F−2
Burkholderiaceae
Campylobacteraceae
Carnobacteriaceae
Corynebacteriaceae
Flavobacteriaceae
Fusobacteriaceae
Gemellaceae
Lachnospiraceae
Lachnospiraceae_XIV
Leptotrichiaceae
Micrococcaceae
Neisseriaceae
Other: < 2%
Pasteurellaceae
Peptoniphilaceae
Peptostreptococcaceae_XI
Porphyromonadaceae
Prevotellaceae
Propionibacteriaceae
Saccharibacteria_(TM7)_F−1
Selenomonadaceae
Streptococcaceae
Synergistaceae
Treponemataceae
Veillonellaceae