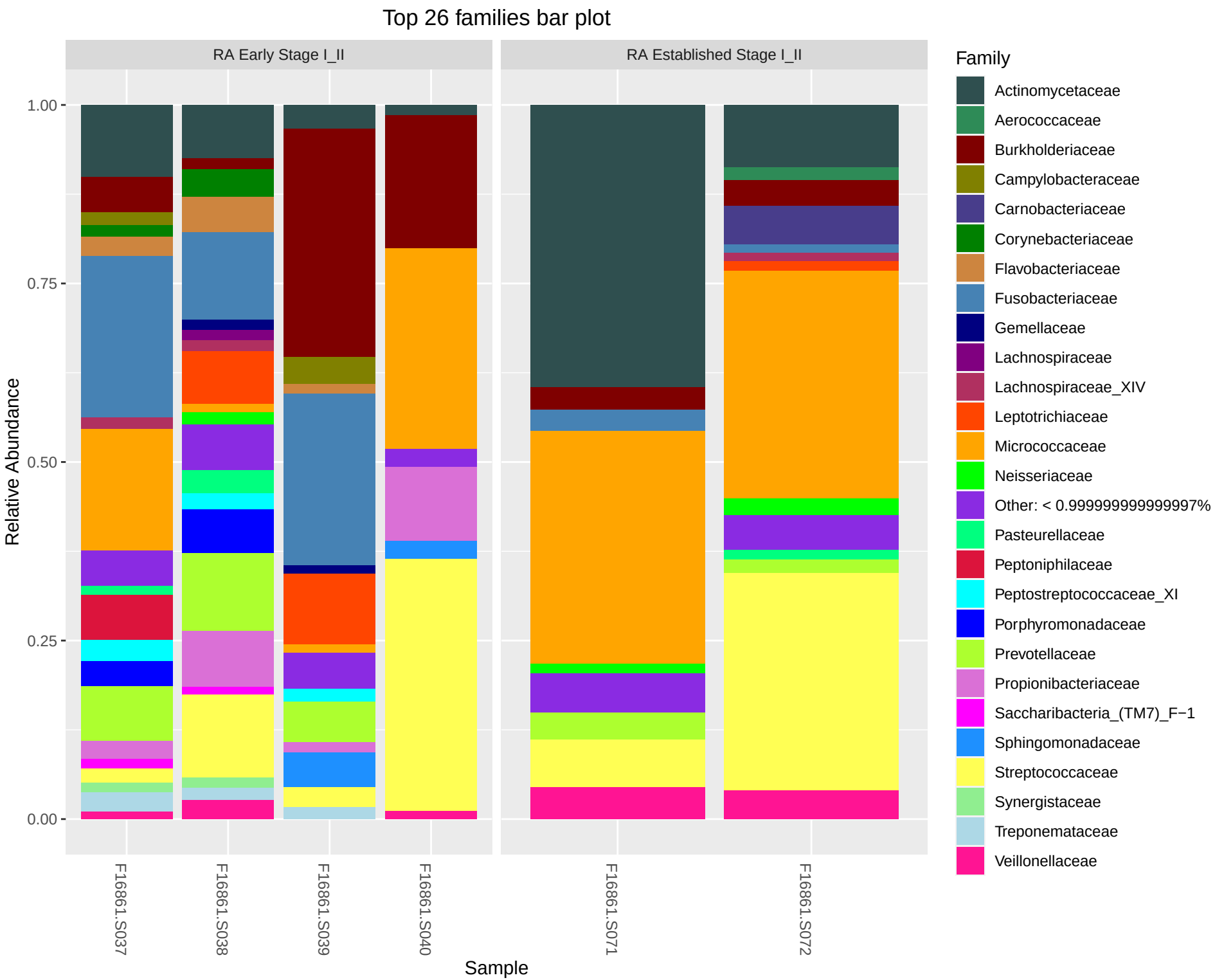
Top 26 families bar plot
RA Early Stage I_II
RA Established Stage I_II
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F16861.S037
F16861.S038
F16861.S039
F16861.S040
F16861.S071
F16861.S072
Sample
Family
Actinomycetaceae
Aerococcaceae
Burkholderiaceae
Campylobacteraceae
Carnobacteriaceae
Corynebacteriaceae
Flavobacteriaceae
Fusobacteriaceae
Gemellaceae
Lachnospiraceae
Lachnospiraceae_XIV
Leptotrichiaceae
Micrococcaceae
Neisseriaceae
Other: < 0.999999999999997%
Pasteurellaceae
Peptoniphilaceae
Peptostreptococcaceae_XI
Porphyromonadaceae
Prevotellaceae
Propionibacteriaceae
Saccharibacteria_(TM7)_F−1
Sphingomonadaceae
Streptococcaceae
Synergistaceae
Treponemataceae
Veillonellaceae