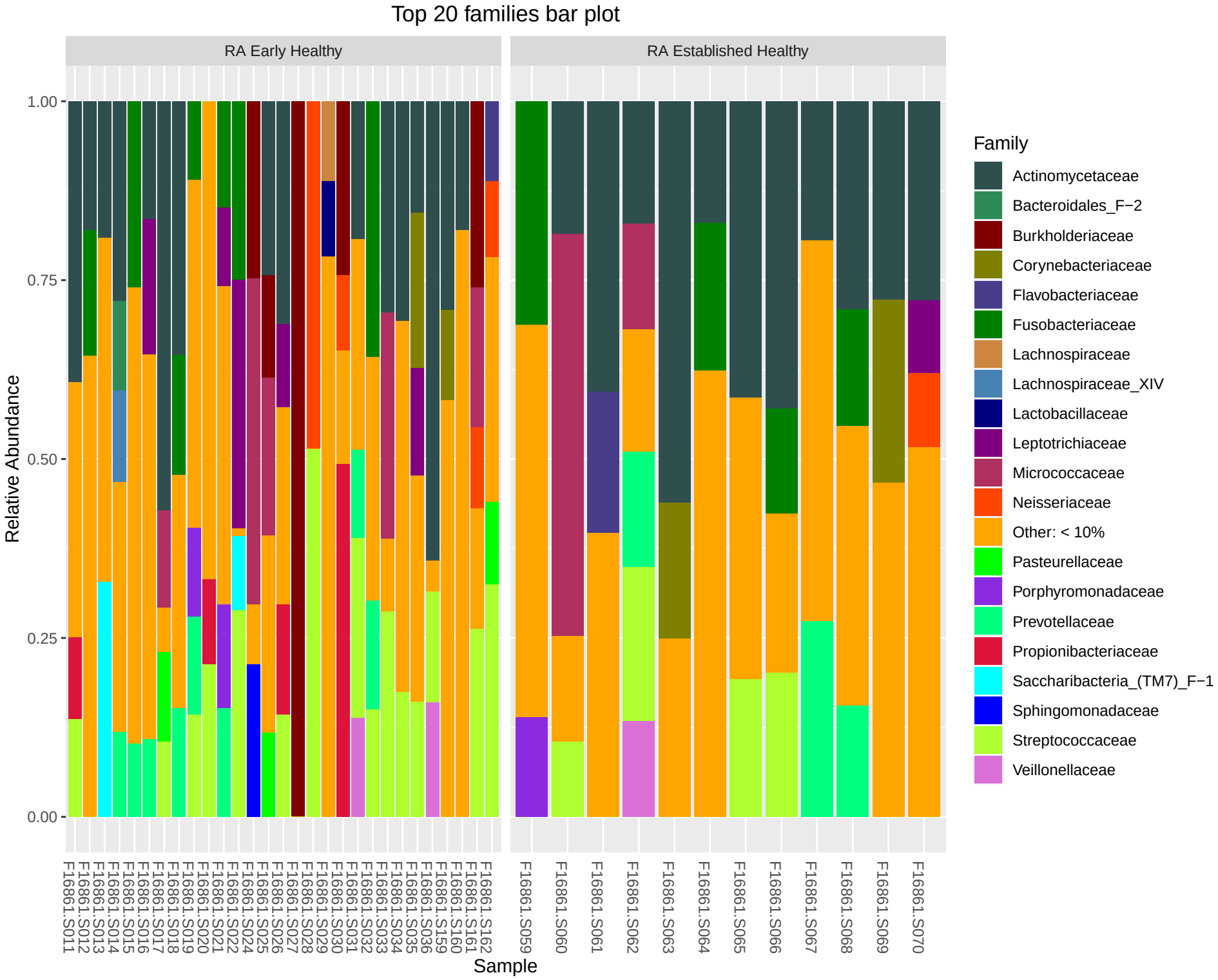
Top 20 families bar plot
RA Early Healthy
RA Established Healthy
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F16861.S011
F16861.S012
F16861.S013
F16861.S014
F16861.S015
F16861.S016
F16861.S017
F16861.S018
F16861.S019
F16861.S020
F16861.S021
F16861.S022
F16861.S024
F16861.S025
F16861.S026
F16861.S027
F16861.S028
F16861.S029
F16861.S030
F16861.S031
F16861.S032
F16861.S033
F16861.S034
F16861.S035
F16861.S036
F16861.S159
F16861.S160
F16861.S161
F16861.S162
F16861.S059
F16861.S060
F16861.S061
F16861.S062
F16861.S063
F16861.S064
F16861.S065
F16861.S066
F16861.S067
F16861.S068
F16861.S069
F16861.S070
Sample
Family
Actinomycetaceae
Bacteroidales_F−2
Burkholderiaceae
Corynebacteriaceae
Flavobacteriaceae
Fusobacteriaceae
Lachnospiraceae
Lachnospiraceae_XIV
Lactobacillaceae
Leptotrichiaceae
Micrococcaceae
Neisseriaceae
Other: < 10%
Pasteurellaceae
Porphyromonadaceae
Prevotellaceae
Propionibacteriaceae
Saccharibacteria_(TM7)_F−1
Sphingomonadaceae
Streptococcaceae
Veillonellaceae