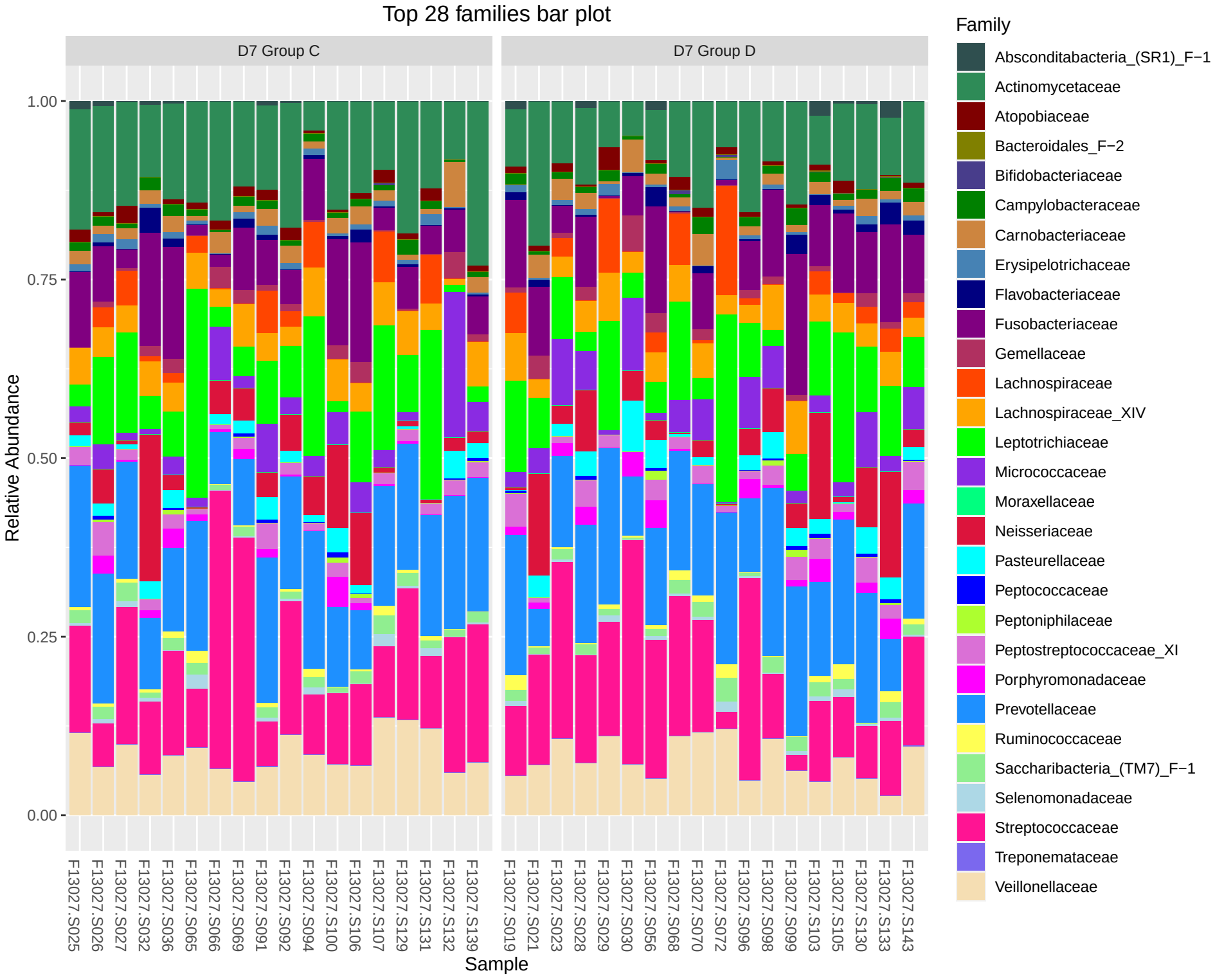
Top 28 families bar plot
D7 Group C
D7 Group D
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F13027.S025
F13027.S026
F13027.S027
F13027.S032
F13027.S036
F13027.S065
F13027.S066
F13027.S069
F13027.S091
F13027.S092
F13027.S094
F13027.S100
F13027.S106
F13027.S107
F13027.S129
F13027.S131
F13027.S132
F13027.S139
F13027.S019
F13027.S021
F13027.S023
F13027.S028
F13027.S029
F13027.S030
F13027.S056
F13027.S068
F13027.S070
F13027.S072
F13027.S096
F13027.S098
F13027.S099
F13027.S103
F13027.S105
F13027.S130
F13027.S133
F13027.S143
Sample
Family
Absconditabacteria_(SR1)_F−1
Actinomycetaceae
Atopobiaceae
Bacteroidales_F−2
Bifidobacteriaceae
Campylobacteraceae
Carnobacteriaceae
Erysipelotrichaceae
Flavobacteriaceae
Fusobacteriaceae
Gemellaceae
Lachnospiraceae
Lachnospiraceae_XIV
Leptotrichiaceae
Micrococcaceae
Moraxellaceae
Neisseriaceae
Pasteurellaceae
Peptococcaceae
Peptoniphilaceae
Peptostreptococcaceae_XI
Porphyromonadaceae
Prevotellaceae
Ruminococcaceae
Saccharibacteria_(TM7)_F−1
Selenomonadaceae
Streptococcaceae
Treponemataceae
Veillonellaceae