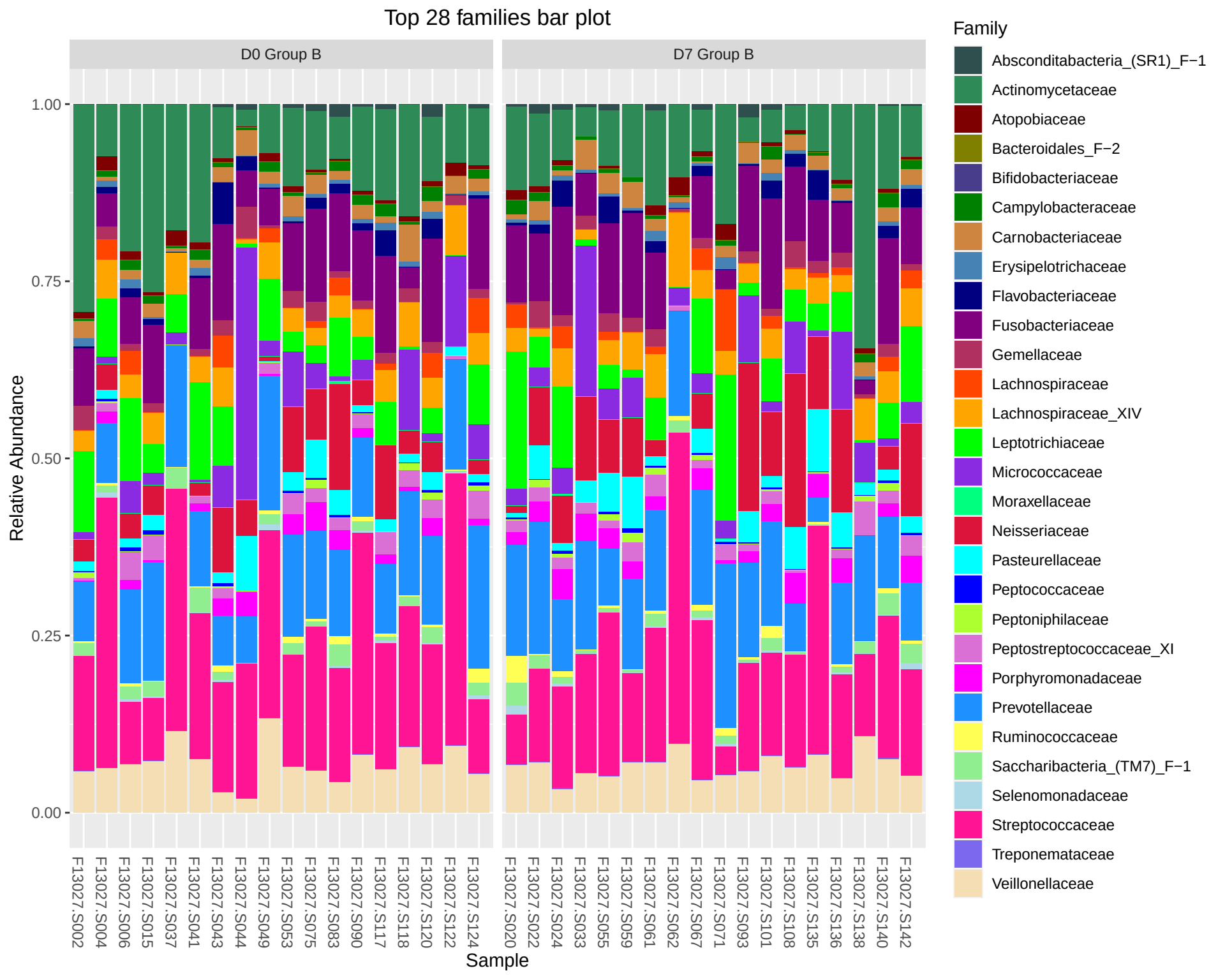
Top 28 families bar plot
D0 Group B
D7 Group B
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F13027.S002
F13027.S004
F13027.S006
F13027.S015
F13027.S037
F13027.S041
F13027.S043
F13027.S044
F13027.S049
F13027.S053
F13027.S075
F13027.S083
F13027.S090
F13027.S117
F13027.S118
F13027.S120
F13027.S122
F13027.S124
F13027.S020
F13027.S022
F13027.S024
F13027.S033
F13027.S055
F13027.S059
F13027.S061
F13027.S062
F13027.S067
F13027.S071
F13027.S093
F13027.S101
F13027.S108
F13027.S135
F13027.S136
F13027.S138
F13027.S140
F13027.S142
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