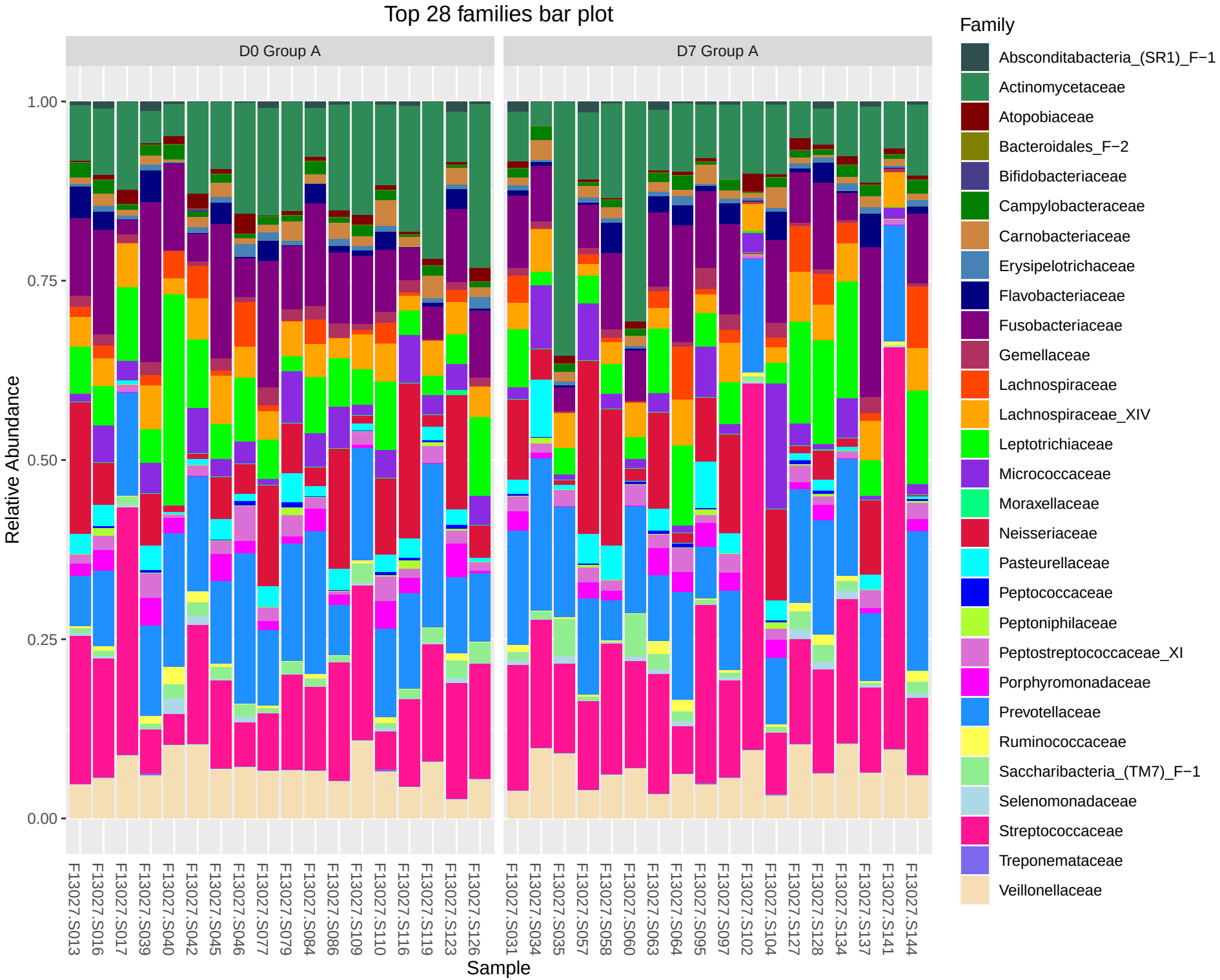
Top 28 families bar plot
D0 Group A
D7 Group A
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F13027.S013
F13027.S016
F13027.S017
F13027.S039
F13027.S040
F13027.S042
F13027.S045
F13027.S046
F13027.S077
F13027.S079
F13027.S084
F13027.S086
F13027.S109
F13027.S110
F13027.S116
F13027.S119
F13027.S123
F13027.S126
F13027.S031
F13027.S034
F13027.S035
F13027.S057
F13027.S058
F13027.S060
F13027.S063
F13027.S064
F13027.S095
F13027.S097
F13027.S102
F13027.S104
F13027.S127
F13027.S128
F13027.S134
F13027.S137
F13027.S141
F13027.S144
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