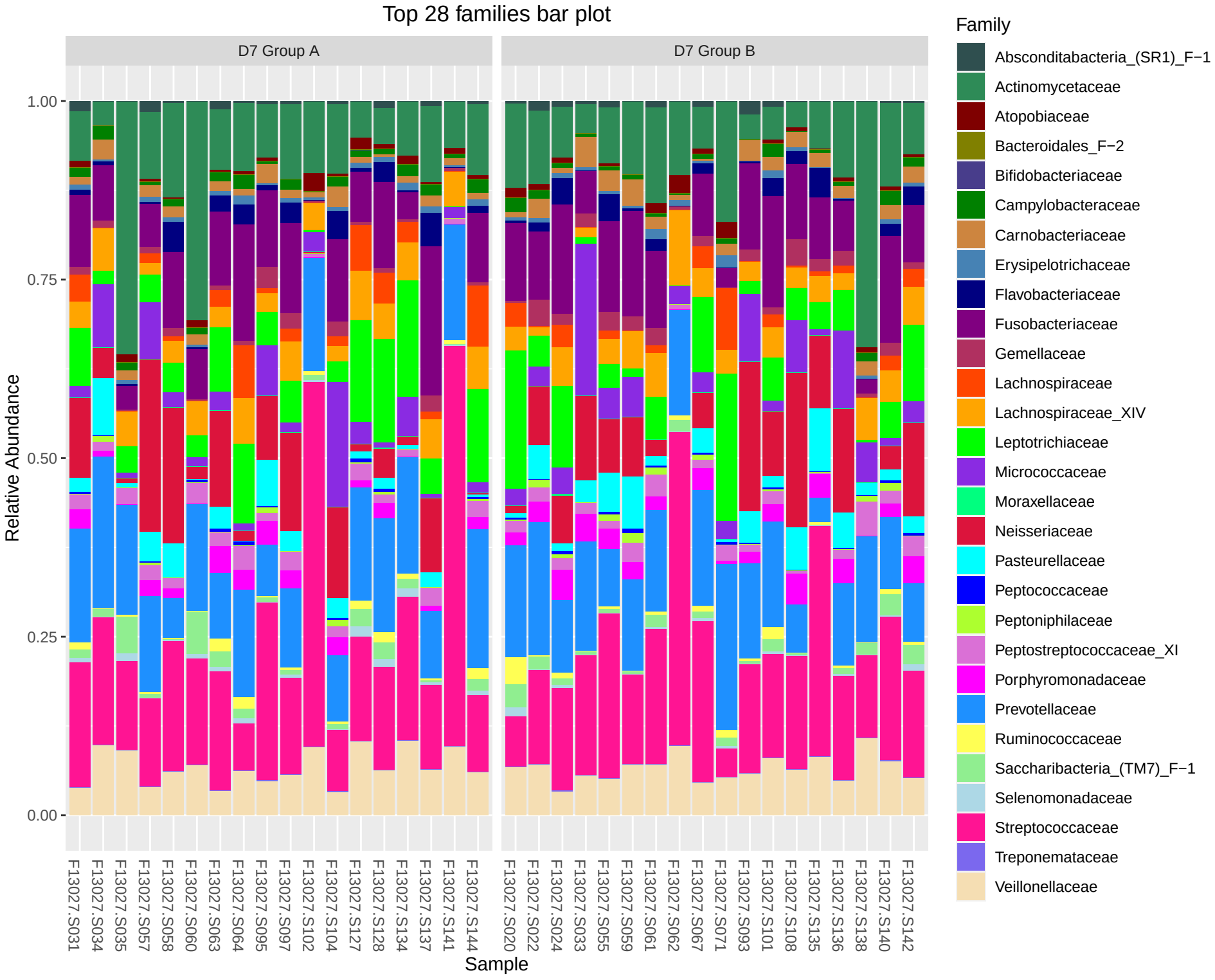
Top 28 families bar plot
D7 Group A
D7 Group B
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F13027.S031
F13027.S034
F13027.S035
F13027.S057
F13027.S058
F13027.S060
F13027.S063
F13027.S064
F13027.S095
F13027.S097
F13027.S102
F13027.S104
F13027.S127
F13027.S128
F13027.S134
F13027.S137
F13027.S141
F13027.S144
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