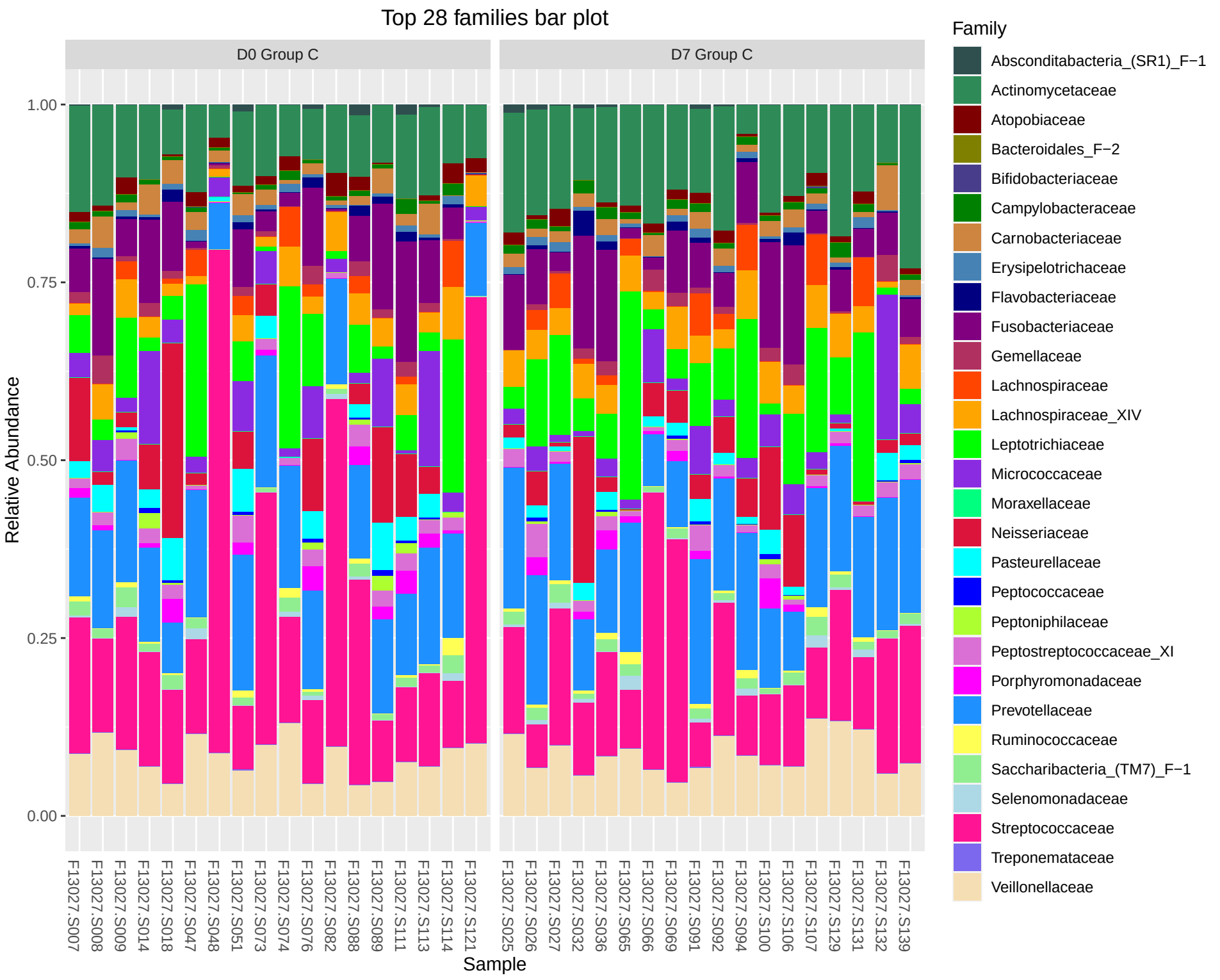
Top 28 families bar plot
D0 Group C
D7 Group C
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F13027.S007
F13027.S008
F13027.S009
F13027.S014
F13027.S018
F13027.S047
F13027.S048
F13027.S051
F13027.S073
F13027.S074
F13027.S076
F13027.S082
F13027.S088
F13027.S089
F13027.S111
F13027.S113
F13027.S114
F13027.S121
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
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