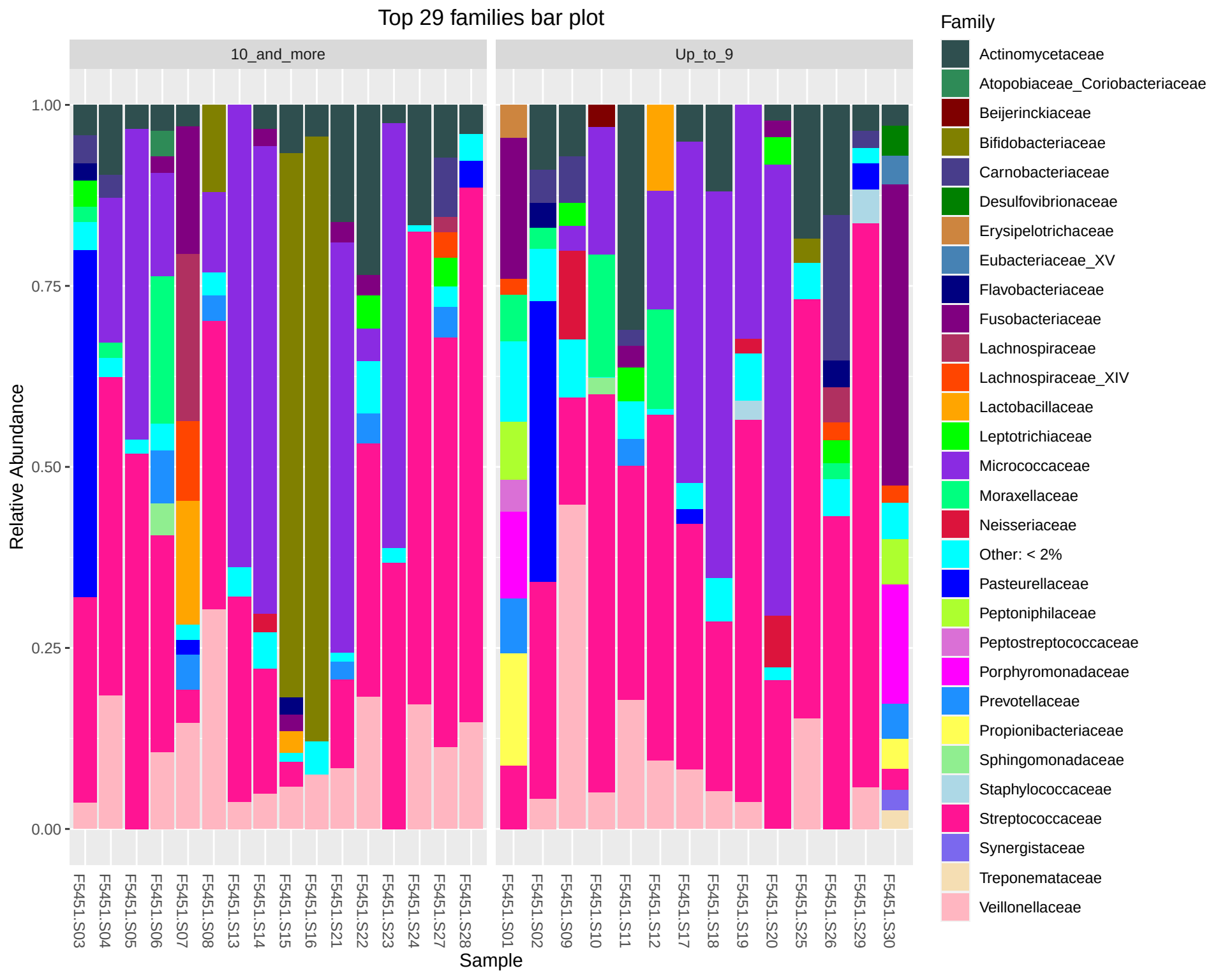
Top 29 families bar plot
10_and_more
Up_to_9
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F5451.S03
F5451.S04
F5451.S05
F5451.S06
F5451.S07
F5451.S08
F5451.S13
F5451.S14
F5451.S15
F5451.S16
F5451.S21
F5451.S22
F5451.S23
F5451.S24
F5451.S27
F5451.S28
F5451.S01
F5451.S02
F5451.S09
F5451.S10
F5451.S11
F5451.S12
F5451.S17
F5451.S18
F5451.S19
F5451.S20
F5451.S25
F5451.S26
F5451.S29
F5451.S30
Sample
Family
Actinomycetaceae
Atopobiaceae_Coriobacteriaceae
Beijerinckiaceae
Bifidobacteriaceae
Carnobacteriaceae
Desulfovibrionaceae
Erysipelotrichaceae
Eubacteriaceae_XV
Flavobacteriaceae
Fusobacteriaceae
Lachnospiraceae
Lachnospiraceae_XIV
Lactobacillaceae
Leptotrichiaceae
Micrococcaceae
Moraxellaceae
Neisseriaceae
Other: < 2%
Pasteurellaceae
Peptoniphilaceae
Peptostreptococcaceae
Porphyromonadaceae
Prevotellaceae
Propionibacteriaceae
Sphingomonadaceae
Staphylococcaceae
Streptococcaceae
Synergistaceae
Treponemataceae
Veillonellaceae