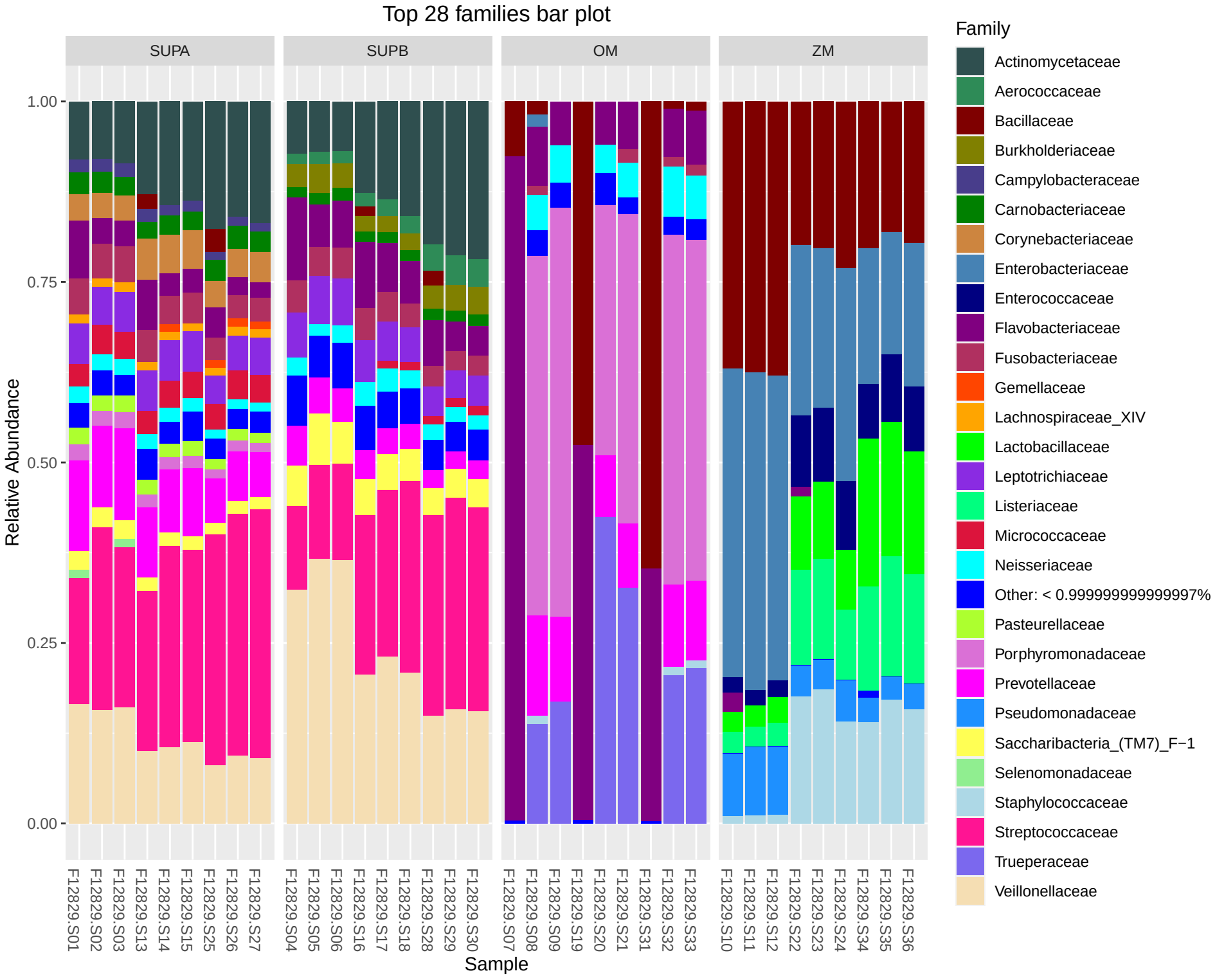
Top 28 families bar plot
SUPA
SUPB
OM
ZM
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F12829.S01
F12829.S02
F12829.S03
F12829.S13
F12829.S14
F12829.S15
F12829.S25
F12829.S26
F12829.S27
F12829.S04
F12829.S05
F12829.S06
F12829.S16
F12829.S17
F12829.S18
F12829.S28
F12829.S29
F12829.S30
F12829.S07
F12829.S08
F12829.S09
F12829.S19
F12829.S20
F12829.S21
F12829.S31
F12829.S32
F12829.S33
F12829.S10
F12829.S11
F12829.S12
F12829.S22
F12829.S23
F12829.S24
F12829.S34
F12829.S35
F12829.S36
Sample
Family
Actinomycetaceae
Aerococcaceae
Bacillaceae
Burkholderiaceae
Campylobacteraceae
Carnobacteriaceae
Corynebacteriaceae
Enterobacteriaceae
Enterococcaceae
Flavobacteriaceae
Fusobacteriaceae
Gemellaceae
Lachnospiraceae_XIV
Lactobacillaceae
Leptotrichiaceae
Listeriaceae
Micrococcaceae
Neisseriaceae
Other: < 0.999999999999997%
Pasteurellaceae
Porphyromonadaceae
Prevotellaceae
Pseudomonadaceae
Saccharibacteria_(TM7)_F−1
Selenomonadaceae
Staphylococcaceae
Streptococcaceae
Trueperaceae
Veillonellaceae