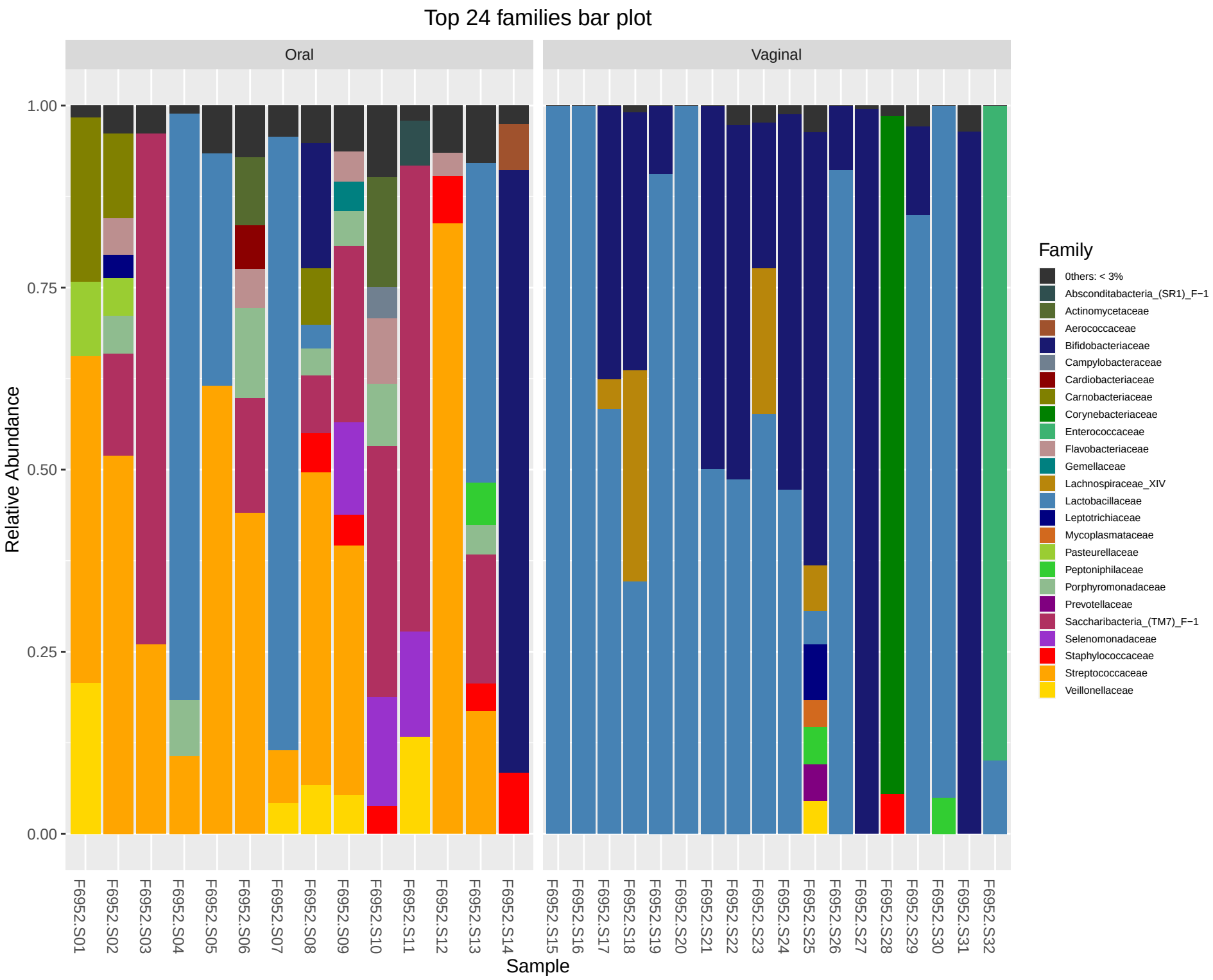
Top 24 families bar plot
Oral
Vaginal
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F6952.S01
F6952.S02
F6952.S03
F6952.S04
F6952.S05
F6952.S06
F6952.S07
F6952.S08
F6952.S09
F6952.S10
F6952.S11
F6952.S12
F6952.S13
F6952.S14
F6952.S15
F6952.S16
F6952.S17
F6952.S18
F6952.S19
F6952.S20
F6952.S21
F6952.S22
F6952.S23
F6952.S24
F6952.S25
F6952.S26
F6952.S27
F6952.S28
F6952.S29
F6952.S30
F6952.S31
F6952.S32
Sample
Family
0thers: < 3%
Absconditabacteria_(SR1)_F−1
Actinomycetaceae
Aerococcaceae
Bifidobacteriaceae
Campylobacteraceae
Cardiobacteriaceae
Carnobacteriaceae
Corynebacteriaceae
Enterococcaceae
Flavobacteriaceae
Gemellaceae
Lachnospiraceae_XIV
Lactobacillaceae
Leptotrichiaceae
Mycoplasmataceae
Pasteurellaceae
Peptoniphilaceae
Porphyromonadaceae
Prevotellaceae
Saccharibacteria_(TM7)_F−1
Selenomonadaceae
Staphylococcaceae
Streptococcaceae
Veillonellaceae