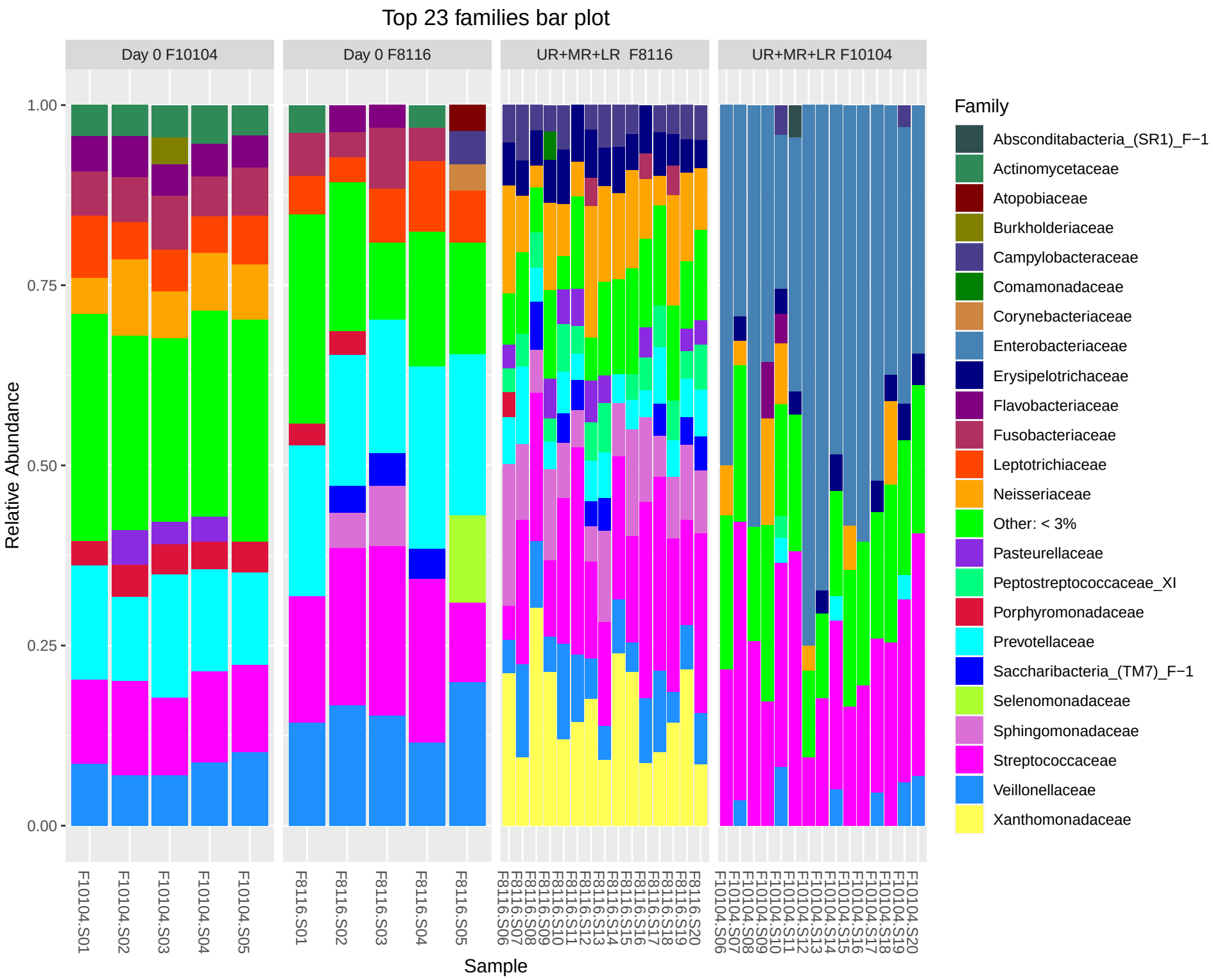
Top 23 families bar plot
Day 0 F10104
Day 0 F8116
UR+MR+LR F8116
UR+MR+LR F10104
Relative Abundance
1.00
0.75
0.50
0.25
0.00
Sample
F10104.S01
F10104.S02
F10104.S03
F10104.S04
F10104.S05
F8116.S01
F8116.S02
F8116.S03
F8116.S04
F8116.S05
F8116.S06
F8116.S07
F8116.S08
F8116.S09
F8116.S10
F8116.S11
F8116.S12
F8116.S13
F8116.S14
F8116.S15
F8116.S16
F8116.S17
F8116.S18
F8116.S19
F8116.S20
F10104.S06
F10104.S07
F10104.S08
F10104.S09
F10104.S10
F10104.S11
F10104.S12
F10104.S13
F10104.S14
F10104.S15
F10104.S16
F10104.S17
F10104.S18
F10104.S19
F10104.S20
Family
Absconditabacteria_(SR1)_F−1
Actinomycetaceae
Atopobiaceae
Burkholderiaceae
Campylobacteraceae
Comamonadaceae
Corynebacteriaceae
Enterobacteriaceae
Erysipelotrichaceae
Flavobacteriaceae
Fusobacteriaceae
Leptotrichiaceae
Neisseriaceae
Other: < 3%
Pasteurellaceae
Peptostreptococcaceae_XI
Porphyromonadaceae
Prevotellaceae
Saccharibacteria_(TM7)_F−1
Selenomonadaceae
Sphingomonadaceae
Streptococcaceae
Veillonellaceae
Xanthomonadaceae