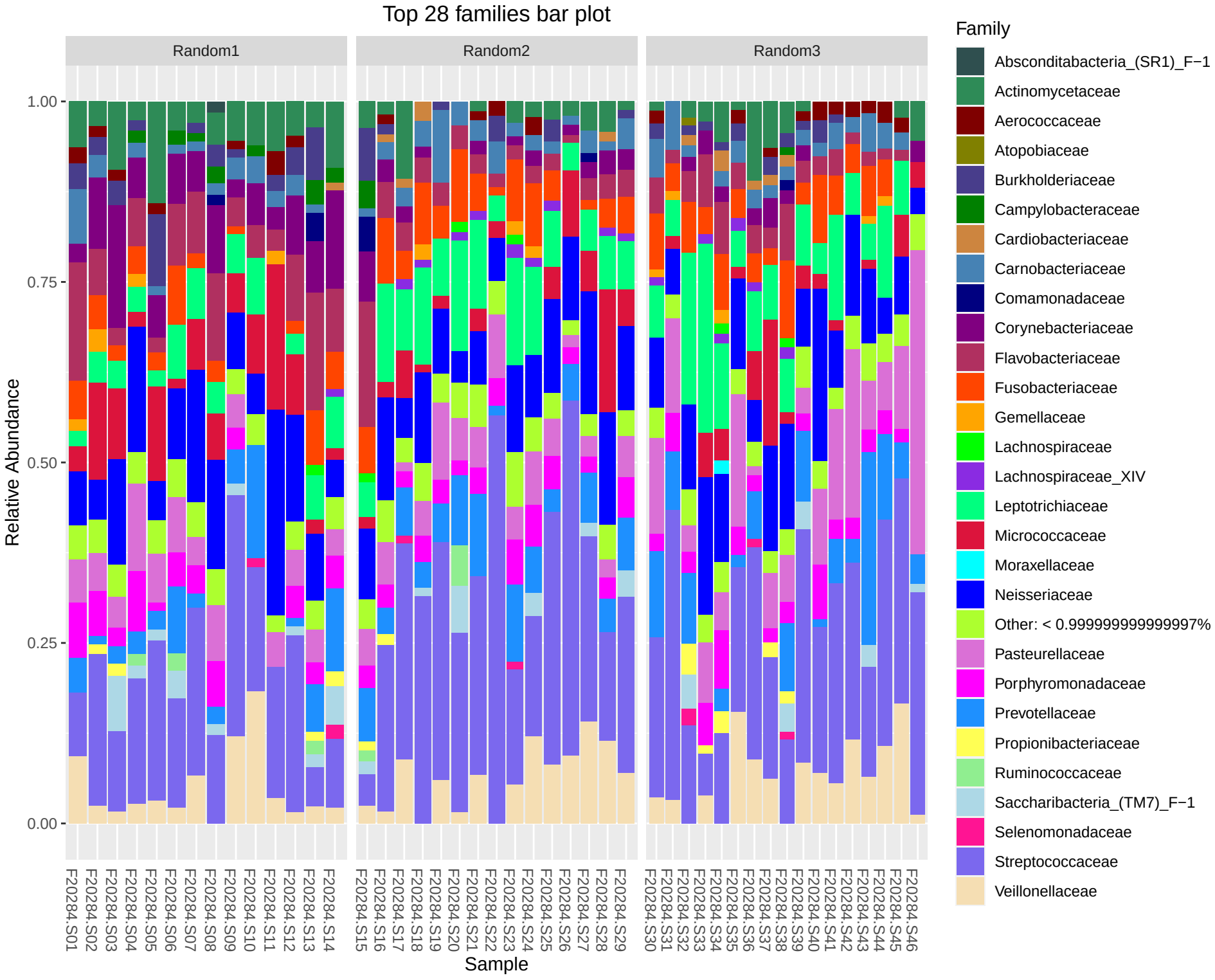
Top 28 families bar plot
Random1
Random2
Random3
Relative Abundance
1.00
0.75
0.50
0.25
0.00
Sample
F20284.S01
F20284.S02
F20284.S03
F20284.S04
F20284.S05
F20284.S06
F20284.S07
F20284.S08
F20284.S09
F20284.S10
F20284.S11
F20284.S12
F20284.S13
F20284.S14
F20284.S15
F20284.S16
F20284.S17
F20284.S18
F20284.S19
F20284.S20
F20284.S21
F20284.S22
F20284.S23
F20284.S24
F20284.S25
F20284.S26
F20284.S27
F20284.S28
F20284.S29
F20284.S30
F20284.S31
F20284.S32
F20284.S33
F20284.S34
F20284.S35
F20284.S36
F20284.S37
F20284.S38
F20284.S39
F20284.S40
F20284.S41
F20284.S42
F20284.S43
F20284.S44
F20284.S45
F20284.S46
Family
Absconditabacteria_(SR1)_F−1
Actinomycetaceae
Aerococcaceae
Atopobiaceae
Burkholderiaceae
Campylobacteraceae
Cardiobacteriaceae
Carnobacteriaceae
Comamonadaceae
Corynebacteriaceae
Flavobacteriaceae
Fusobacteriaceae
Gemellaceae
Lachnospiraceae
Lachnospiraceae_XIV
Leptotrichiaceae
Micrococcaceae
Moraxellaceae
Neisseriaceae
Other: < 0.999999999999997%
Pasteurellaceae
Porphyromonadaceae
Prevotellaceae
Propionibacteriaceae
Ruminococcaceae
Saccharibacteria_(TM7)_F−1
Selenomonadaceae
Streptococcaceae
Veillonellaceae