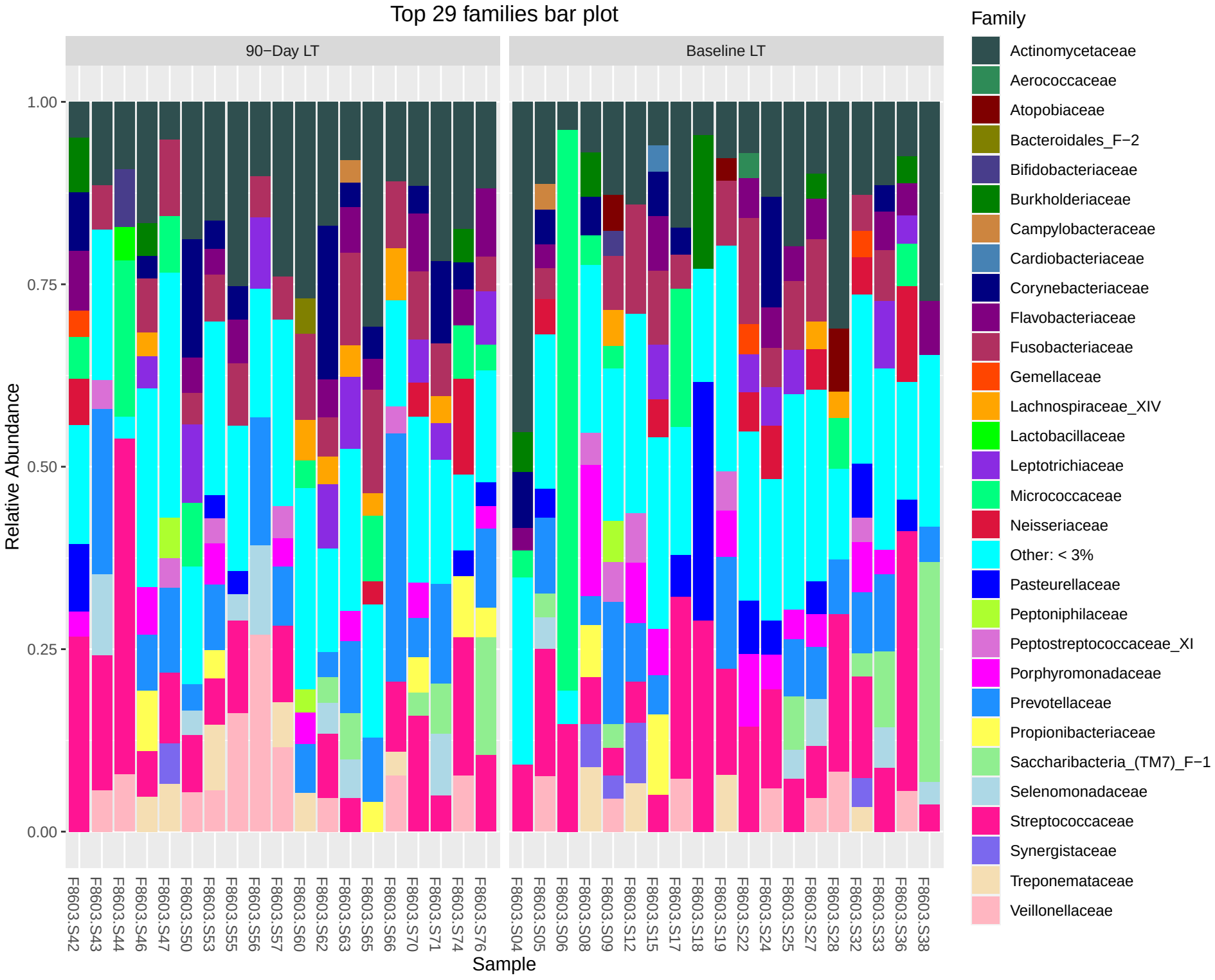
Top 29 families bar plot
90−Day LT
Baseline LT
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F8603.S42
F8603.S43
F8603.S44
F8603.S46
F8603.S47
F8603.S50
F8603.S53
F8603.S55
F8603.S56
F8603.S57
F8603.S60
F8603.S62
F8603.S63
F8603.S65
F8603.S66
F8603.S70
F8603.S71
F8603.S74
F8603.S76
F8603.S04
F8603.S05
F8603.S06
F8603.S08
F8603.S09
F8603.S12
F8603.S15
F8603.S17
F8603.S18
F8603.S19
F8603.S22
F8603.S24
F8603.S25
F8603.S27
F8603.S28
F8603.S32
F8603.S33
F8603.S36
F8603.S38
Sample
Family
Actinomycetaceae
Aerococcaceae
Atopobiaceae
Bacteroidales_F−2
Bifidobacteriaceae
Burkholderiaceae
Campylobacteraceae
Cardiobacteriaceae
Corynebacteriaceae
Flavobacteriaceae
Fusobacteriaceae
Gemellaceae
Lachnospiraceae_XIV
Lactobacillaceae
Leptotrichiaceae
Micrococcaceae
Neisseriaceae
Other: < 3%
Pasteurellaceae
Peptoniphilaceae
Peptostreptococcaceae_XI
Porphyromonadaceae
Prevotellaceae
Propionibacteriaceae
Saccharibacteria_(TM7)_F−1
Selenomonadaceae
Streptococcaceae
Synergistaceae
Treponemataceae
Veillonellaceae