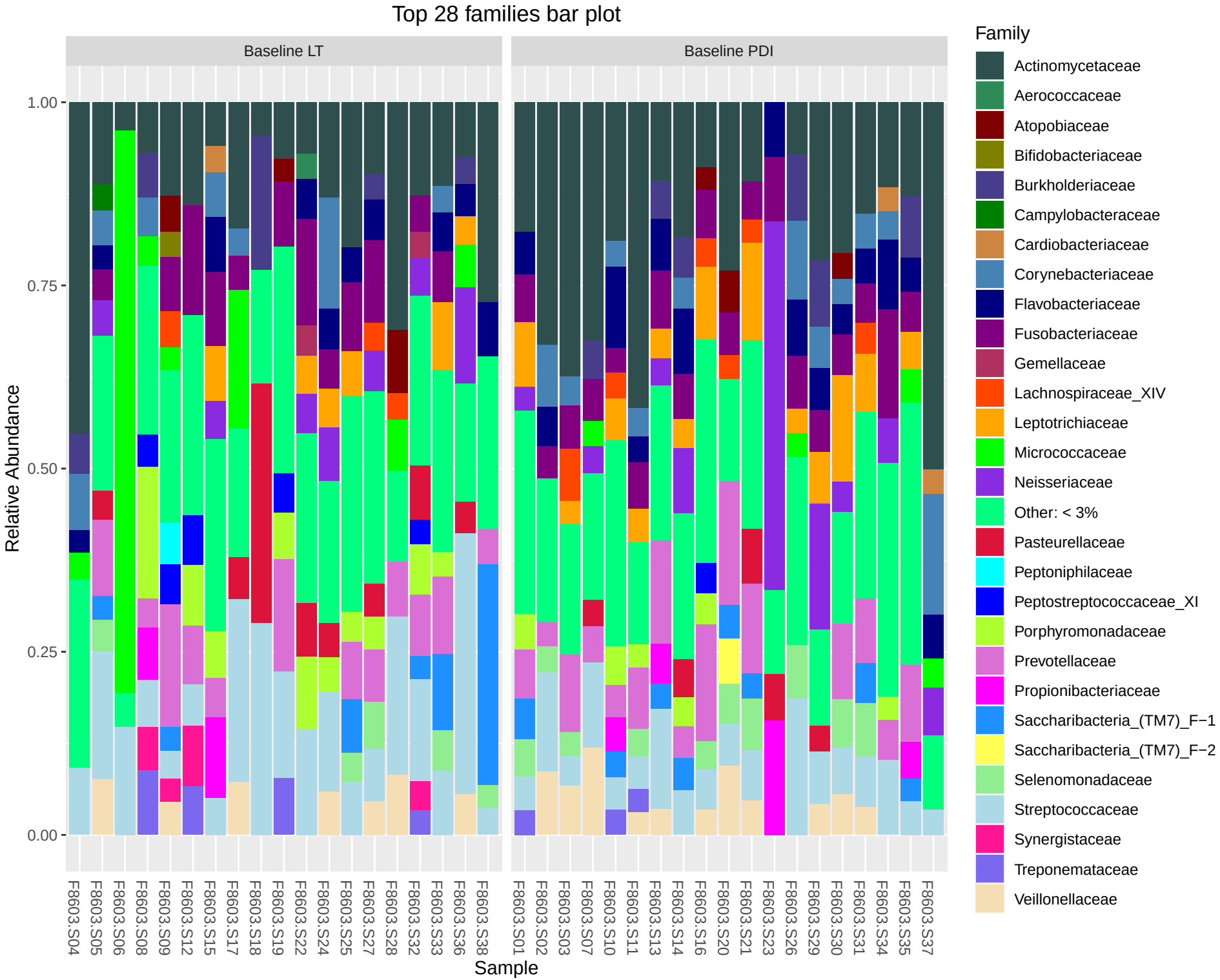
Top 28 families bar plot
Baseline LT
Baseline PDI
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F8603.S04
F8603.S05
F8603.S06
F8603.S08
F8603.S09
F8603.S12
F8603.S15
F8603.S17
F8603.S18
F8603.S19
F8603.S22
F8603.S24
F8603.S25
F8603.S27
F8603.S28
F8603.S32
F8603.S33
F8603.S36
F8603.S38
F8603.S01
F8603.S02
F8603.S03
F8603.S07
F8603.S10
F8603.S11
F8603.S13
F8603.S14
F8603.S16
F8603.S20
F8603.S21
F8603.S23
F8603.S26
F8603.S29
F8603.S30
F8603.S31
F8603.S34
F8603.S35
F8603.S37
Sample
Family
Actinomycetaceae
Aerococcaceae
Atopobiaceae
Bifidobacteriaceae
Burkholderiaceae
Campylobacteraceae
Cardiobacteriaceae
Corynebacteriaceae
Flavobacteriaceae
Fusobacteriaceae
Gemellaceae
Lachnospiraceae_XIV
Leptotrichiaceae
Micrococcaceae
Neisseriaceae
Other: < 3%
Pasteurellaceae
Peptoniphilaceae
Peptostreptococcaceae_XI
Porphyromonadaceae
Prevotellaceae
Propionibacteriaceae
Saccharibacteria_(TM7)_F−1
Saccharibacteria_(TM7)_F−2
Selenomonadaceae
Streptococcaceae
Synergistaceae
Treponemataceae
Veillonellaceae