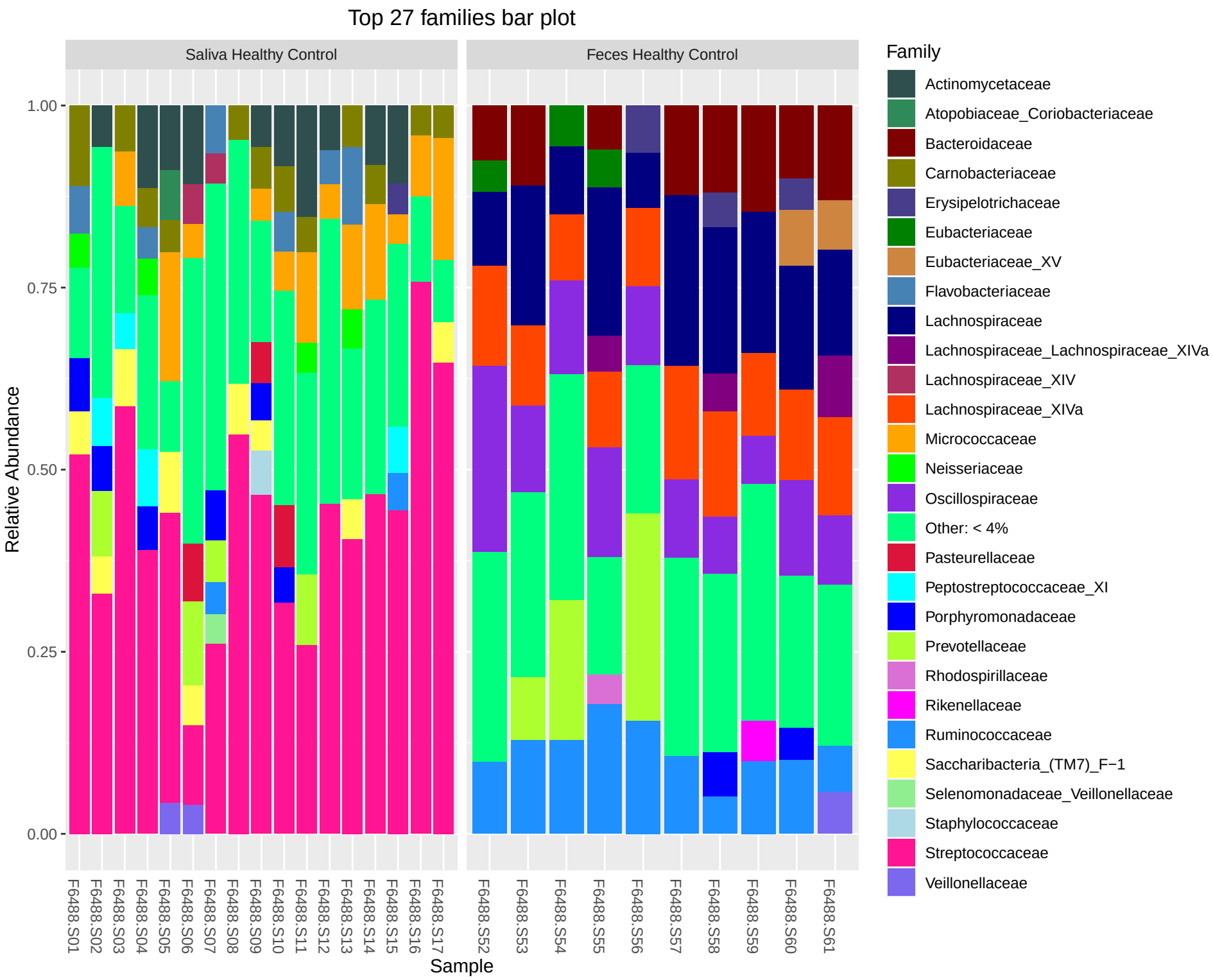
Top 27 families bar plot
Saliva Healthy Control
Feces Healthy Control
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F6488.S01
F6488.S02
F6488.S03
F6488.S04
F6488.S05
F6488.S06
F6488.S07
F6488.S08
F6488.S09
F6488.S10
F6488.S11
F6488.S12
F6488.S13
F6488.S14
F6488.S15
F6488.S16
F6488.S17
F6488.S52
F6488.S53
F6488.S54
F6488.S55
F6488.S56
F6488.S57
F6488.S58
F6488.S59
F6488.S60
F6488.S61
Sample
Family
Actinomycetaceae
Atopobiaceae_Coriobacteriaceae
Bacteroidaceae
Carnobacteriaceae
Erysipelotrichaceae
Eubacteriaceae
Eubacteriaceae_XV
Flavobacteriaceae
Lachnospiraceae
Lachnospiraceae_Lachnospiraceae_XIVa
Lachnospiraceae_XIV
Lachnospiraceae_XIVa
Micrococcaceae
Neisseriaceae
Oscillospiraceae
Other: < 4%
Pasteurellaceae
Peptostreptococcaceae_XI
Porphyromonadaceae
Prevotellaceae
Rhodospirillaceae
Rikenellaceae
Ruminococcaceae
Saccharibacteria_(TM7)_F−1
Selenomonadaceae_Veillonellaceae
Staphylococcaceae
Streptococcaceae
Veillonellaceae