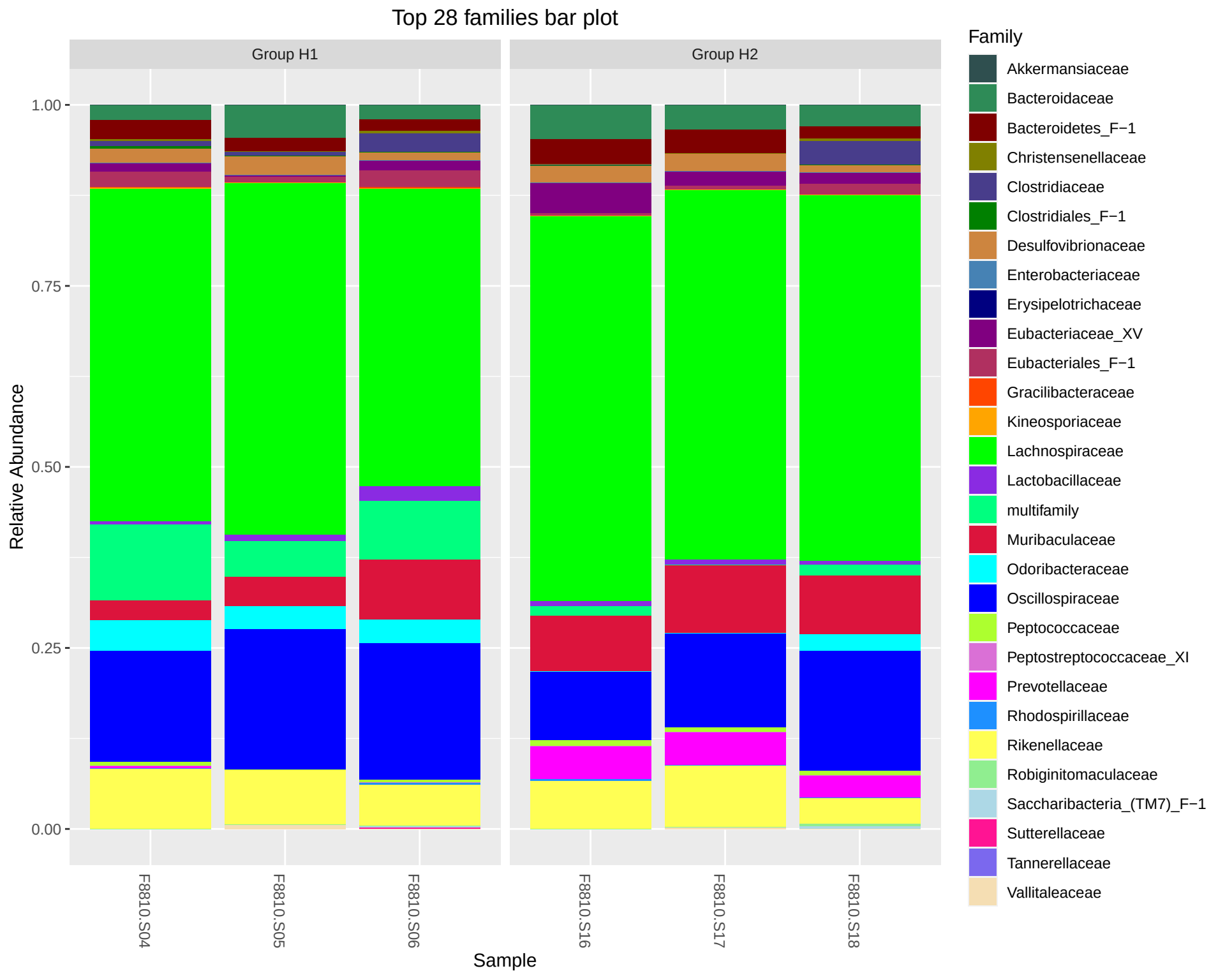
Top 28 families bar plot
Group H1
Group H2
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F8810.S04
F8810.S05
F8810.S06
F8810.S16
F8810.S17
F8810.S18
Sample
Family
Akkermansiaceae
Bacteroidaceae
Bacteroidetes_F−1
Christensenellaceae
Clostridiaceae
Clostridiales_F−1
Desulfovibrionaceae
Enterobacteriaceae
Erysipelotrichaceae
Eubacteriaceae_XV
Eubacteriales_F−1
Gracilibacteraceae
Kineosporiaceae
Lachnospiraceae
Lactobacillaceae
multifamily
Muribaculaceae
Odoribacteraceae
Oscillospiraceae
Peptococcaceae
Peptostreptococcaceae_XI
Prevotellaceae
Rhodospirillaceae
Rikenellaceae
Robiginitomaculaceae
Saccharibacteria_(TM7)_F−1
Sutterellaceae
Tannerellaceae
Vallitaleaceae