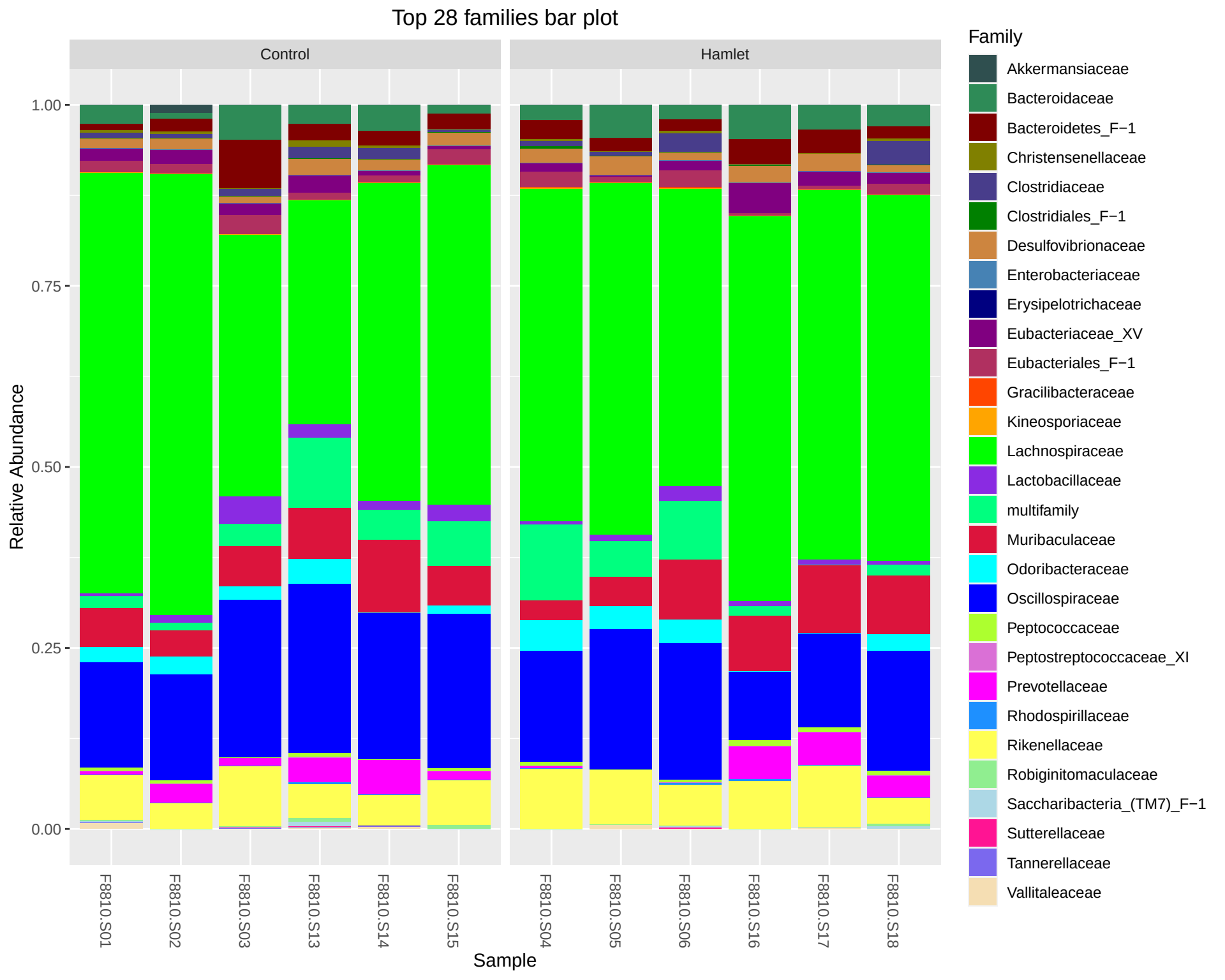
Top 28 families bar plot
Control
Hamlet
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F8810.S01
F8810.S02
F8810.S03
F8810.S13
F8810.S14
F8810.S15
F8810.S04
F8810.S05
F8810.S06
F8810.S16
F8810.S17
F8810.S18
Sample
Family
Akkermansiaceae
Bacteroidaceae
Bacteroidetes_F−1
Christensenellaceae
Clostridiaceae
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