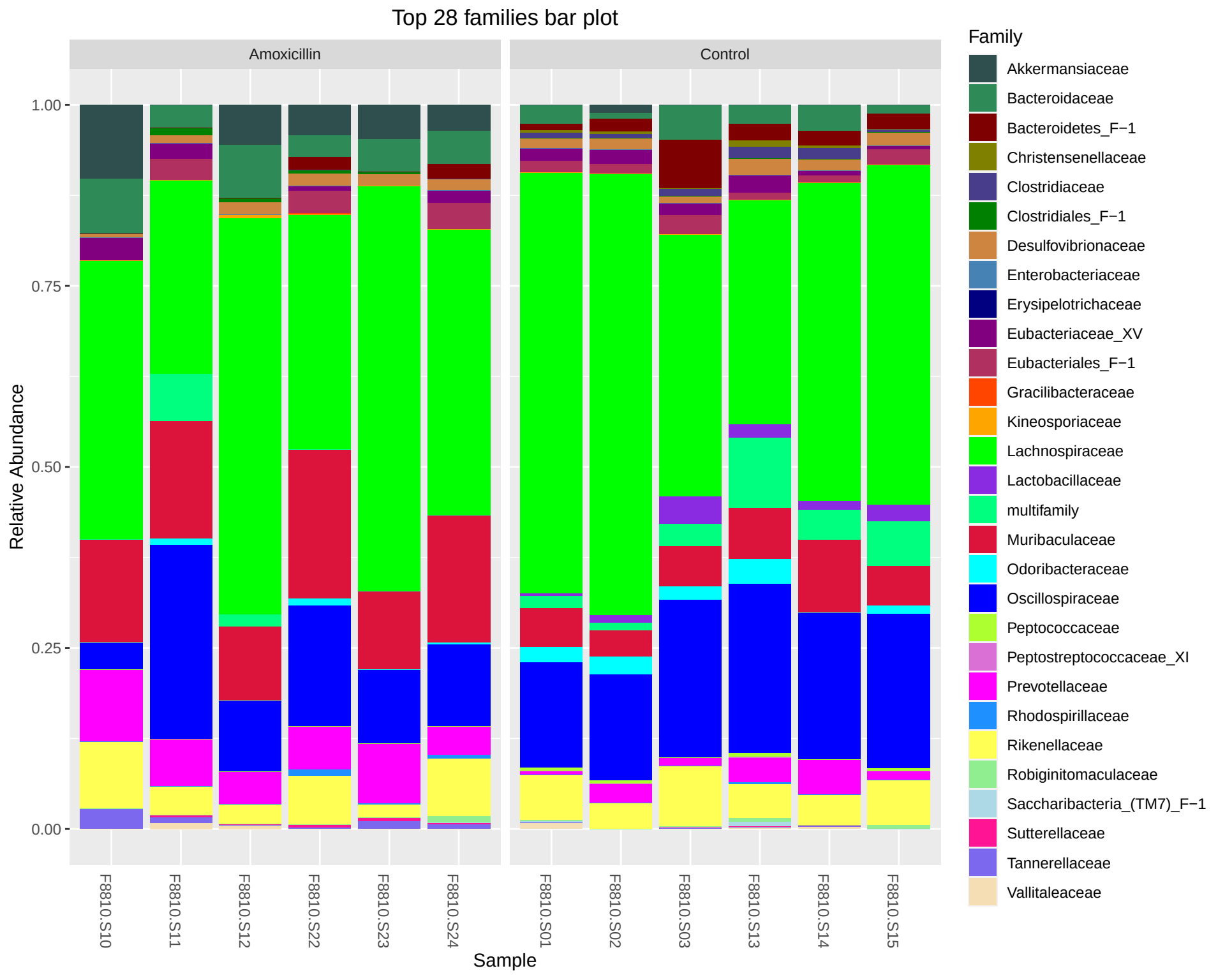
Top 28 families bar plot
Amoxicillin
Control
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F8810.S10
F8810.S11
F8810.S12
F8810.S22
F8810.S23
F8810.S24
F8810.S01
F8810.S02
F8810.S03
F8810.S13
F8810.S14
F8810.S15
Sample
Family
Akkermansiaceae
Bacteroidaceae
Bacteroidetes_F−1
Christensenellaceae
Clostridiaceae
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