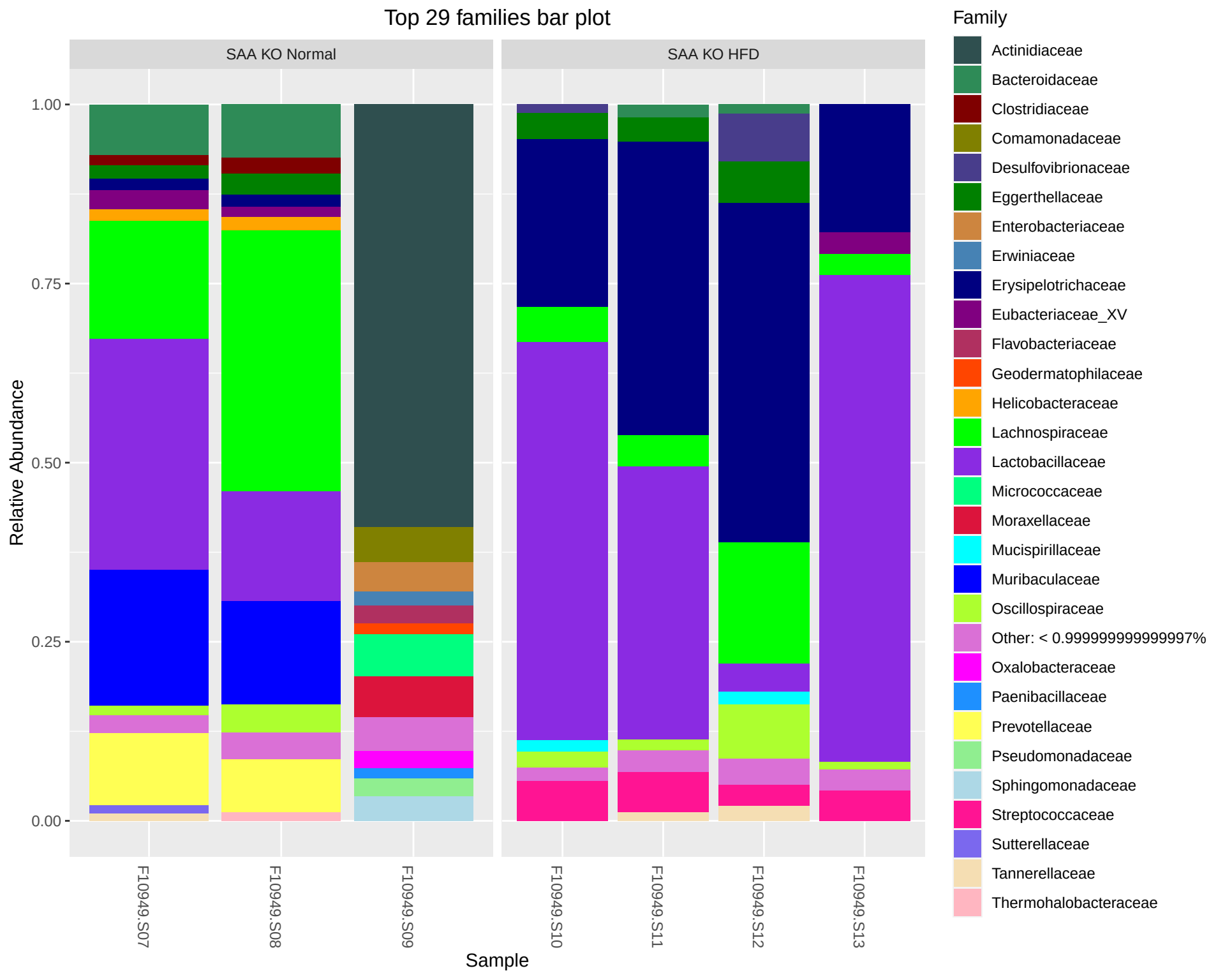
Top 29 families bar plot
SAA KO Normal
SAA KO HFD
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F10949.S07
F10949.S08
F10949.S09
F10949.S10
F10949.S11
F10949.S12
F10949.S13
Sample
Family
Actinidiaceae
Bacteroidaceae
Clostridiaceae
Comamonadaceae
Desulfovibrionaceae
Eggerthellaceae
Enterobacteriaceae
Erwiniaceae
Erysipelotrichaceae
Eubacteriaceae_XV
Flavobacteriaceae
Geodermatophilaceae
Helicobacteraceae
Lachnospiraceae
Lactobacillaceae
Micrococcaceae
Moraxellaceae
Mucispirillaceae
Muribaculaceae
Oscillospiraceae
Other: < 0.999999999999997%
Oxalobacteraceae
Paenibacillaceae
Prevotellaceae
Pseudomonadaceae
Sphingomonadaceae
Streptococcaceae
Sutterellaceae
Tannerellaceae
Thermohalobacteraceae