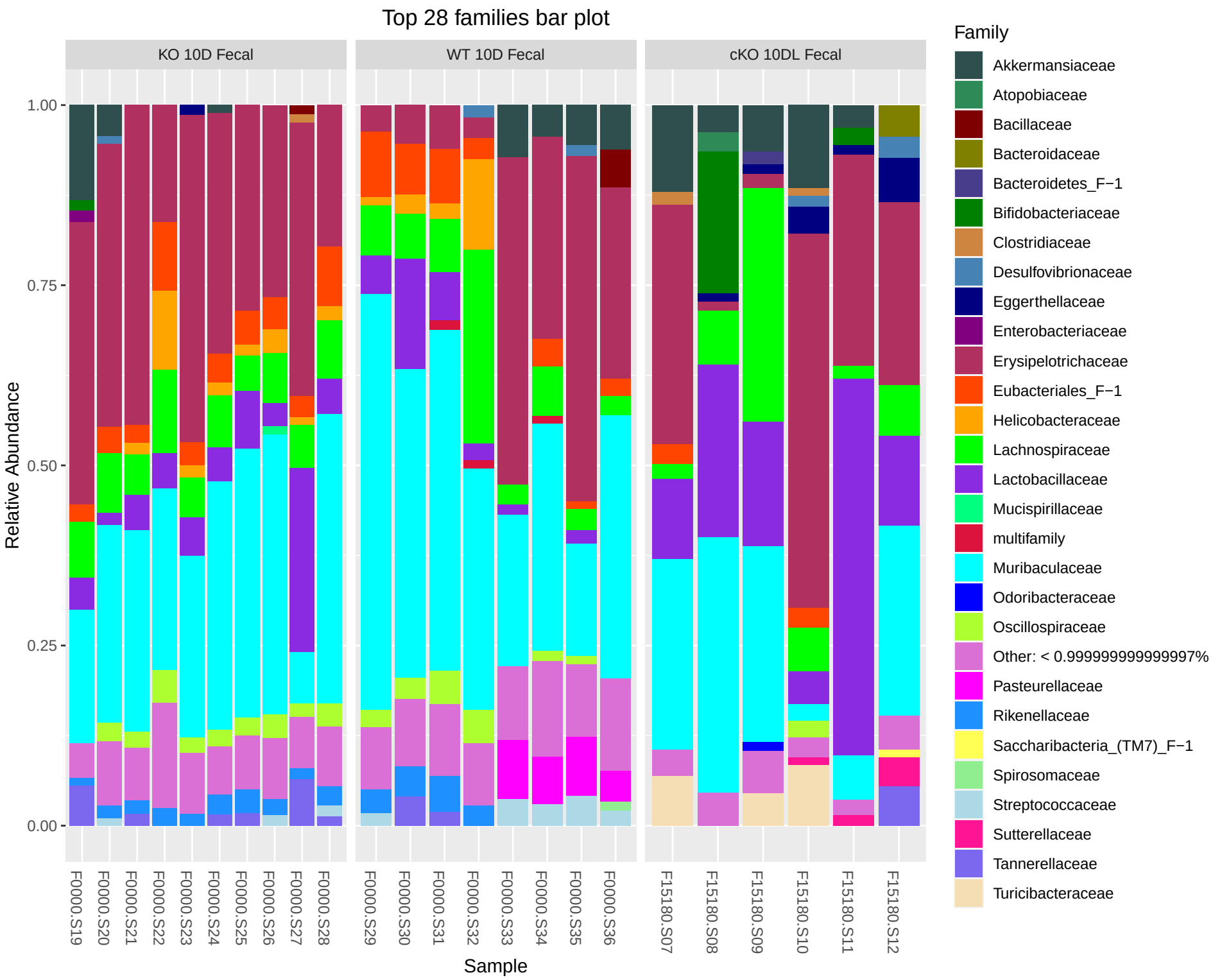
Top 28 families bar plot
KO 10D Fecal
WT 10D Fecal
cKO 10DL Fecal
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F0000.S19
F0000.S20
F0000.S21
F0000.S22
F0000.S23
F0000.S24
F0000.S25
F0000.S26
F0000.S27
F0000.S28
F0000.S29
F0000.S30
F0000.S31
F0000.S32
F0000.S33
F0000.S34
F0000.S35
F0000.S36
F15180.S07
F15180.S08
F15180.S09
F15180.S10
F15180.S11
F15180.S12
Sample
Family
Akkermansiaceae
Atopobiaceae
Bacillaceae
Bacteroidaceae
Bacteroidetes_F−1
Bifidobacteriaceae
Clostridiaceae
Desulfovibrionaceae
Eggerthellaceae
Enterobacteriaceae
Erysipelotrichaceae
Eubacteriales_F−1
Helicobacteraceae
Lachnospiraceae
Lactobacillaceae
Mucispirillaceae
multifamily
Muribaculaceae
Odoribacteraceae
Oscillospiraceae
Other: < 0.999999999999997%
Pasteurellaceae
Rikenellaceae
Saccharibacteria_(TM7)_F−1
Spirosomaceae
Streptococcaceae
Sutterellaceae
Tannerellaceae
Turicibacteraceae