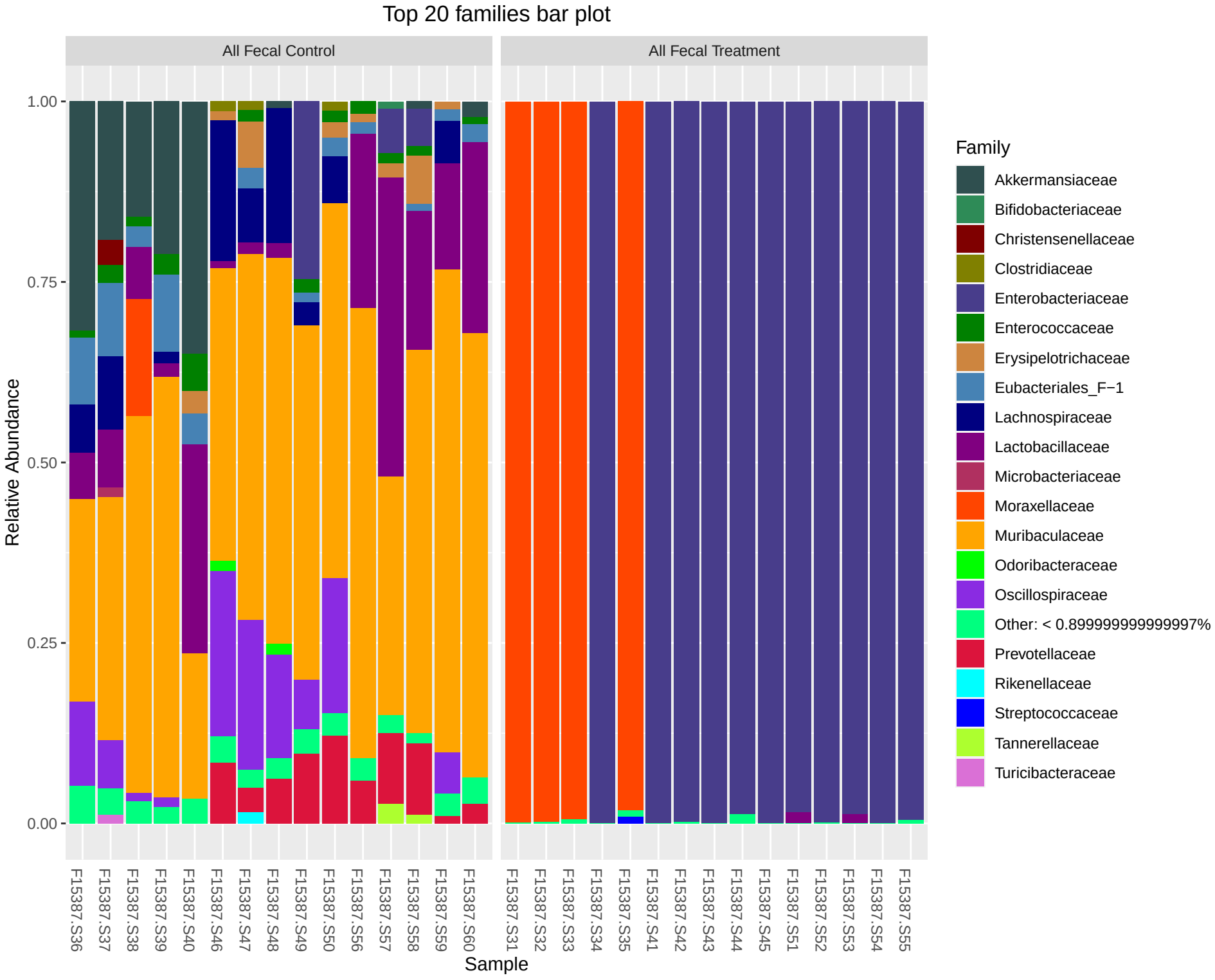
Top 20 families bar plot
All Fecal Control
All Fecal Treatment
Relative Abundance
1.00
0.75
0.50
0.25
0.00
F15387.S36
F15387.S37
F15387.S38
F15387.S39
F15387.S40
F15387.S46
F15387.S47
F15387.S48
F15387.S49
F15387.S50
F15387.S56
F15387.S57
F15387.S58
F15387.S59
F15387.S60
F15387.S31
F15387.S32
F15387.S33
F15387.S34
F15387.S35
F15387.S41
F15387.S42
F15387.S43
F15387.S44
F15387.S45
F15387.S51
F15387.S52
F15387.S53
F15387.S54
F15387.S55
Sample
Family
Akkermansiaceae
Bifidobacteriaceae
Christensenellaceae
Clostridiaceae
Enterobacteriaceae
Enterococcaceae
Erysipelotrichaceae
Eubacteriales_F−1
Lachnospiraceae
Lactobacillaceae
Microbacteriaceae
Moraxellaceae
Muribaculaceae
Odoribacteraceae
Oscillospiraceae
Other: < 0.899999999999997%
Prevotellaceae
Rikenellaceae
Streptococcaceae
Tannerellaceae
Turicibacteraceae