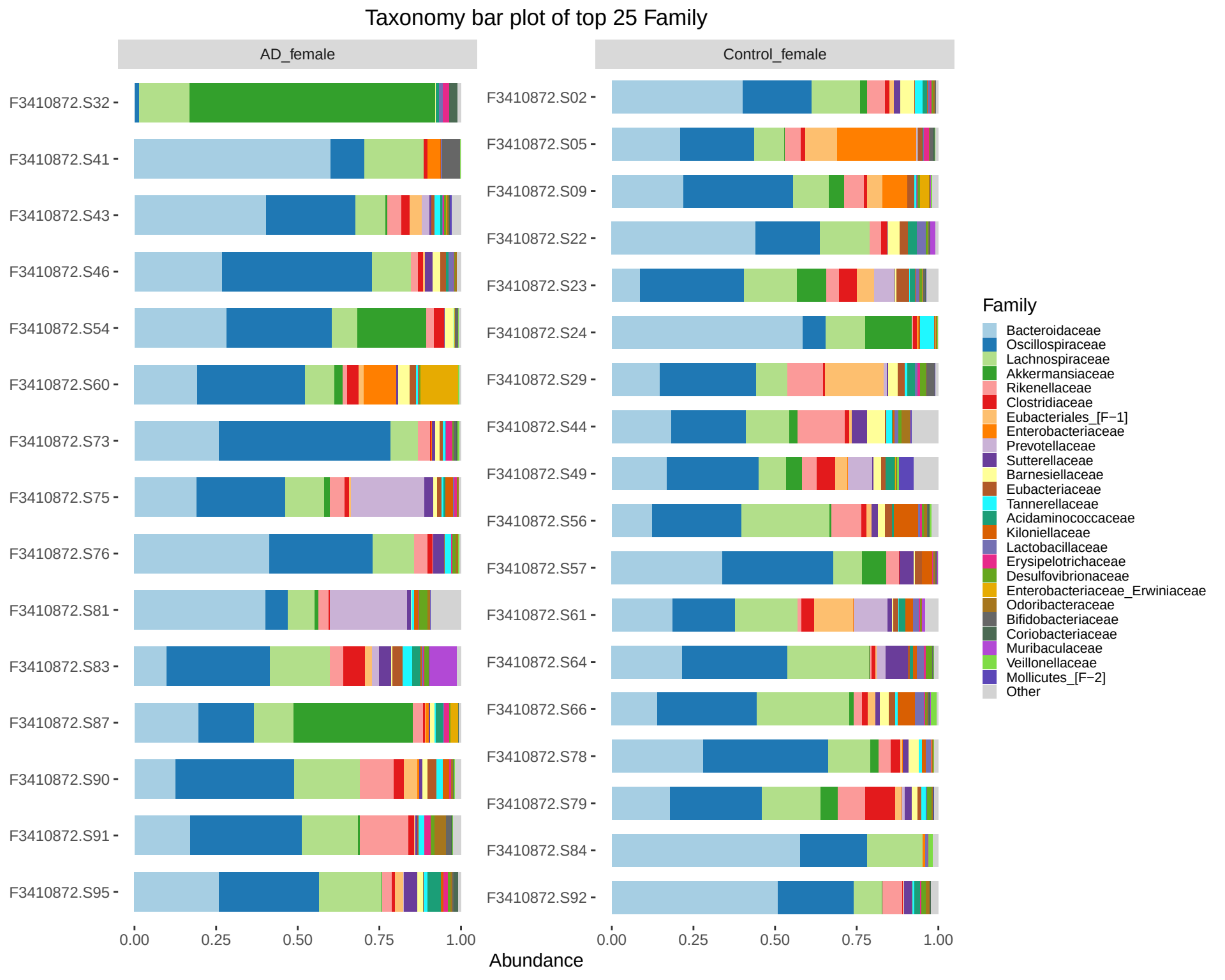
Taxonomy bar plot of top 25 Family
AD_female
Control_female
F3410872.S32
F3410872.S41
F3410872.S43
F3410872.S46
F3410872.S54
F3410872.S60
F3410872.S73
F3410872.S75
F3410872.S76
F3410872.S81
F3410872.S83
F3410872.S87
F3410872.S90
F3410872.S91
F3410872.S95
F3410872.S02
F3410872.S05
F3410872.S09
F3410872.S22
F3410872.S23
F3410872.S24
F3410872.S29
F3410872.S44
F3410872.S49
F3410872.S56
F3410872.S57
F3410872.S61
F3410872.S64
F3410872.S66
F3410872.S78
F3410872.S79
F3410872.S84
F3410872.S92
0.00
0.25
0.50
0.75
1.00
Abundance
Family
Bacteroidaceae
Oscillospiraceae
Lachnospiraceae
Akkermansiaceae
Rikenellaceae
Clostridiaceae
Eubacteriales_[F−1]
Enterobacteriaceae
Prevotellaceae
Sutterellaceae
Barnesiellaceae
Eubacteriaceae
Tannerellaceae
Acidaminococcaceae
Kiloniellaceae
Lactobacillaceae
Erysipelotrichaceae
Desulfovibrionaceae
Enterobacteriaceae_Erwiniaceae
Odoribacteraceae
Bifidobacteriaceae
Coriobacteriaceae
Muribaculaceae
Veillonellaceae
Mollicutes_[F−2]
Other