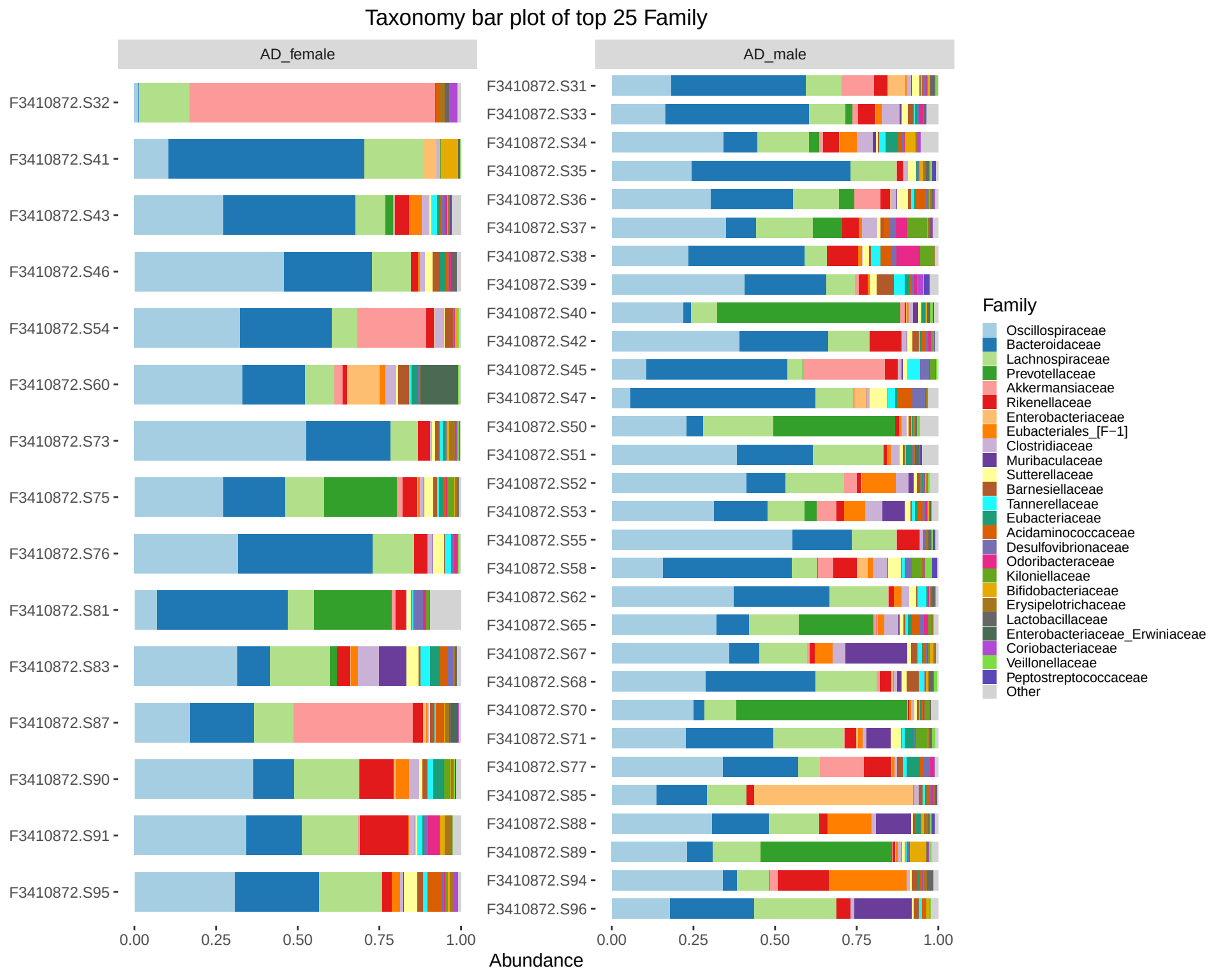
Taxonomy bar plot of top 25 Family
AD_female
AD_male
F3410872.S32
F3410872.S41
F3410872.S43
F3410872.S46
F3410872.S54
F3410872.S60
F3410872.S73
F3410872.S75
F3410872.S76
F3410872.S81
F3410872.S83
F3410872.S87
F3410872.S90
F3410872.S91
F3410872.S95
F3410872.S31
F3410872.S33
F3410872.S34
F3410872.S35
F3410872.S36
F3410872.S37
F3410872.S38
F3410872.S39
F3410872.S40
F3410872.S42
F3410872.S45
F3410872.S47
F3410872.S50
F3410872.S51
F3410872.S52
F3410872.S53
F3410872.S55
F3410872.S58
F3410872.S62
F3410872.S65
F3410872.S67
F3410872.S68
F3410872.S70
F3410872.S71
F3410872.S77
F3410872.S85
F3410872.S88
F3410872.S89
F3410872.S94
F3410872.S96
0.00
0.25
0.50
0.75
1.00
Abundance
Family
Oscillospiraceae
Bacteroidaceae
Lachnospiraceae
Prevotellaceae
Akkermansiaceae
Rikenellaceae
Enterobacteriaceae
Eubacteriales_[F−1]
Clostridiaceae
Muribaculaceae
Sutterellaceae
Barnesiellaceae
Tannerellaceae
Eubacteriaceae
Acidaminococcaceae
Desulfovibrionaceae
Odoribacteraceae
Kiloniellaceae
Bifidobacteriaceae
Erysipelotrichaceae
Lactobacillaceae
Enterobacteriaceae_Erwiniaceae
Coriobacteriaceae
Veillonellaceae
Peptostreptococcaceae
Other