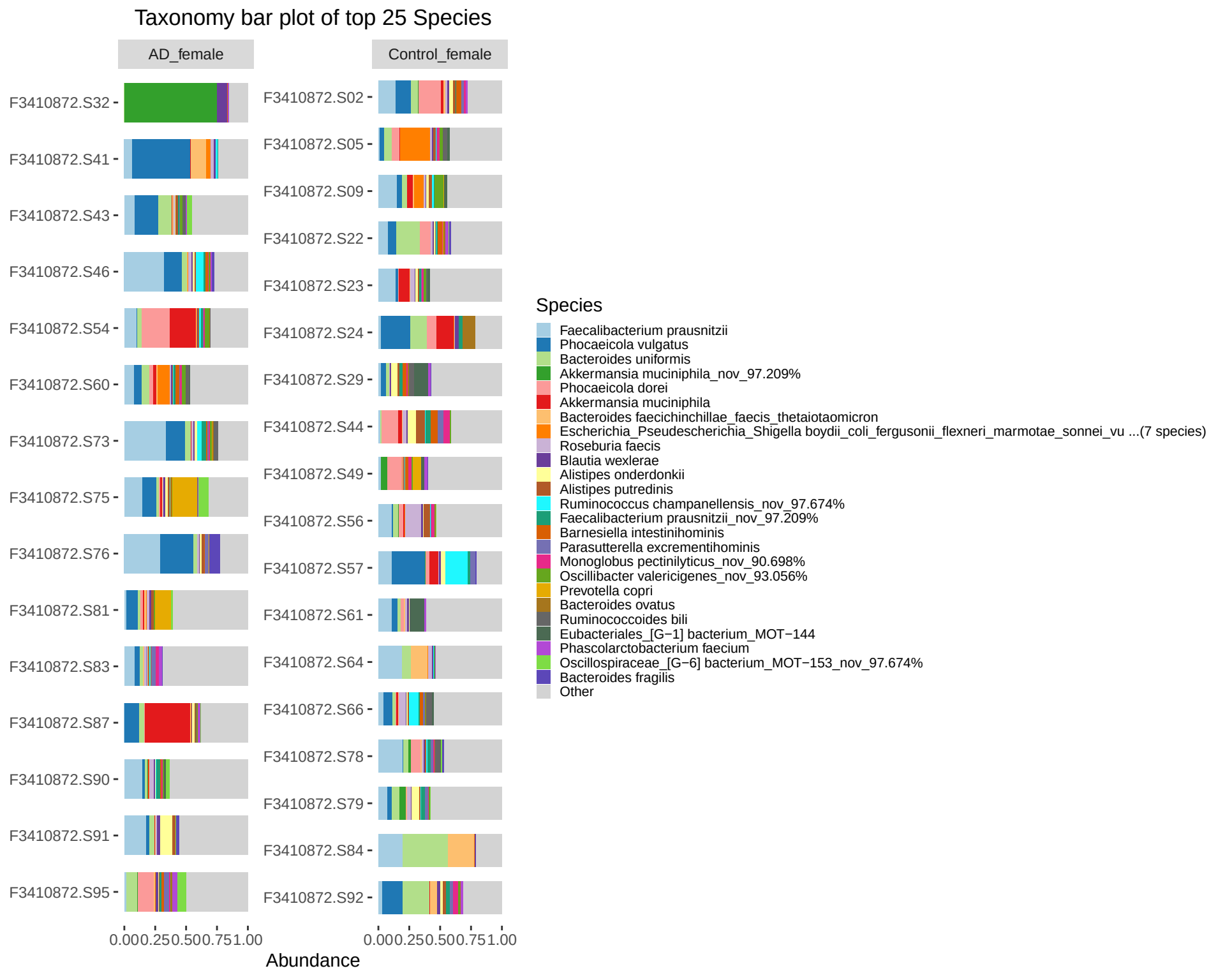
Taxonomy bar plot of top 25 Species
AD_female
Control_female
F3410872.S32
F3410872.S41
F3410872.S43
F3410872.S46
F3410872.S54
F3410872.S60
F3410872.S73
F3410872.S75
F3410872.S76
F3410872.S81
F3410872.S83
F3410872.S87
F3410872.S90
F3410872.S91
F3410872.S95
F3410872.S02
F3410872.S05
F3410872.S09
F3410872.S22
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
0.00 0.25 0.50 0.75 1.00
Abundance
Species
Faecalibacterium prausnitzii
Phocaeicola vulgatus
Bacteroides uniformis
Akkermansia muciniphila_nov_97.209%
Phocaeicola dorei
Akkermansia muciniphila
Bacteroides faecichinchillae_faecis_thetaiotaomicron
Escherichia_Pseudescherichia_Shigella boydii_coli_fergusonii_flexneri_marmotae_sonnei_vu ...(7 species)
Roseburia faecis
Blautia wexlerae
Alistipes onderdonkii
Alistipes putredinis
Ruminococcus champanellensis_nov_97.674%
Faecalibacterium prausnitzii_nov_97.209%
Barnesiella intestinihominis
Parasutterella excrementihominis
Monoglobus pectinilyticus_nov_90.698%
Oscillibacter valericigenes_nov_93.056%
Prevotella copri
Bacteroides ovatus
Ruminococcoides bili
Eubacteriales_[G−1] bacterium_MOT−144
Phascolarctobacterium faecium
Oscillospiraceae_[G−6] bacterium_MOT−153_nov_97.674%
Bacteroides fragilis
Other