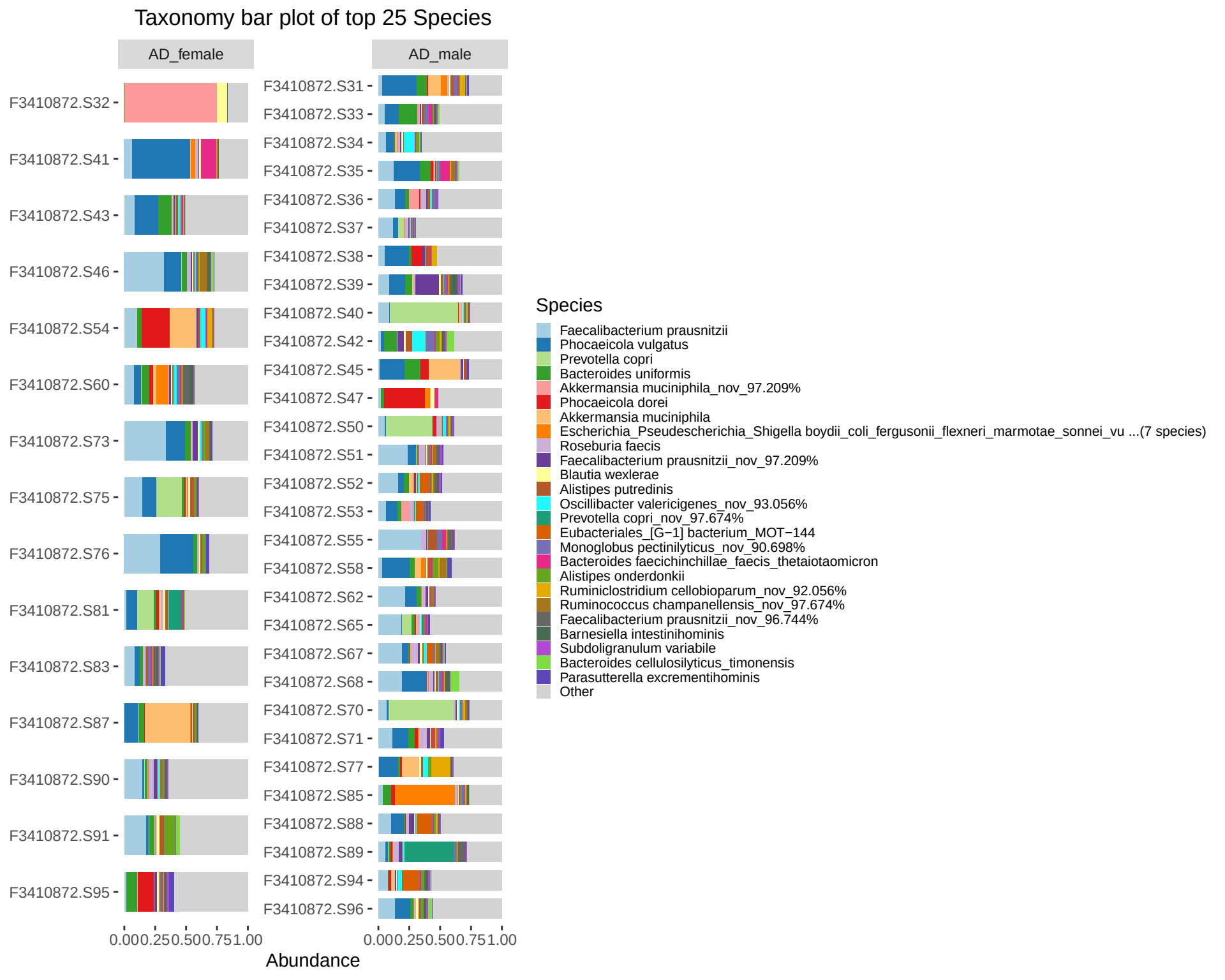
Taxonomy bar plot of top 25 Species
AD_female
AD_male
F3410872.S32
F3410872.S41
F3410872.S43
F3410872.S46
F3410872.S54
F3410872.S60
F3410872.S73
F3410872.S75
F3410872.S76
F3410872.S81
F3410872.S83
F3410872.S87
F3410872.S90
F3410872.S91
F3410872.S95
F3410872.S31
F3410872.S33
F3410872.S34
F3410872.S35
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
0.00 0.25 0.50 0.75 1.00
Abundance
Species
Faecalibacterium prausnitzii
Phocaeicola vulgatus
Prevotella copri
Bacteroides uniformis
Akkermansia muciniphila_nov_97.209%
Phocaeicola dorei
Akkermansia muciniphila
Escherichia_Pseudescherichia_Shigella boydii_coli_fergusonii_flexneri_marmotae_sonnei_vu ...(7 species)
Roseburia faecis
Faecalibacterium prausnitzii_nov_97.209%
Blautia wexlerae
Alistipes putredinis
Oscillibacter valericigenes_nov_93.056%
Prevotella copri_nov_97.674%
Eubacteriales_[G−1] bacterium_MOT−144
Monoglobus pectinilyticus_nov_90.698%
Bacteroides faecichinchillae_faecis_thetaiotaomicron
Alistipes onderdonkii
Ruminiclostridium cellobioparum_nov_92.056%
Ruminococcus champanellensis_nov_97.674%
Faecalibacterium prausnitzii_nov_96.744%
Barnesiella intestinihominis
Subdoligranulum variabile
Bacteroides cellulosilyticus_timonensis
Parasutterella excrementihominis
Other