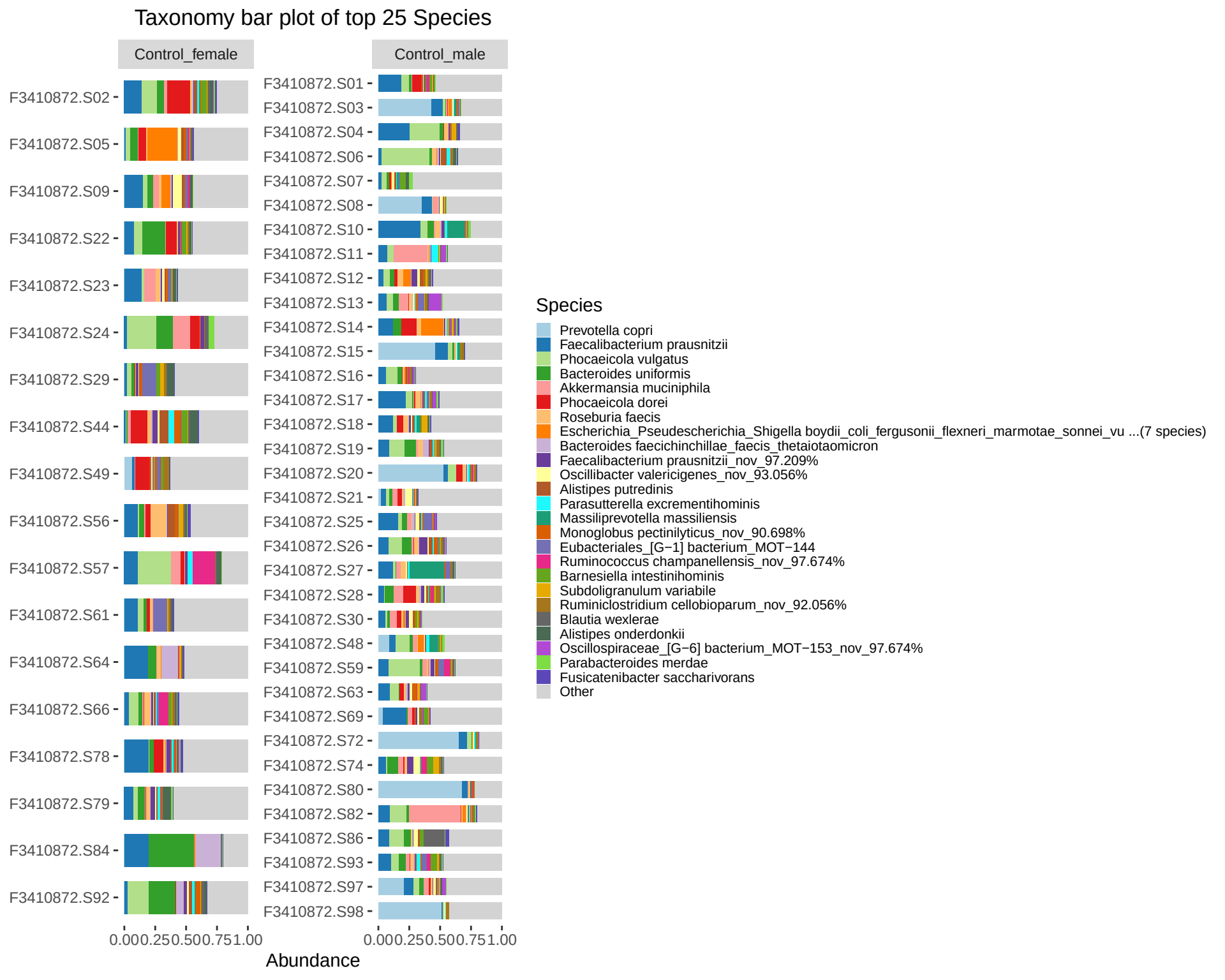
Taxonomy bar plot of top 25 Species
Control_female
Control_male
F3410872.S02
F3410872.S05
F3410872.S09
F3410872.S22
F3410872.S23
F3410872.S24
F3410872.S29
F3410872.S44
F3410872.S49
F3410872.S56
F3410872.S57
F3410872.S61
F3410872.S64
F3410872.S66
F3410872.S78
F3410872.S79
F3410872.S84
F3410872.S92
F3410872.S01
F3410872.S03
F3410872.S04
F3410872.S06
F3410872.S07
F3410872.S08
F3410872.S10
F3410872.S11
F3410872.S12
F3410872.S13
F3410872.S14
F3410872.S15
F3410872.S16
F3410872.S17
F3410872.S18
F3410872.S19
F3410872.S20
F3410872.S21
F3410872.S25
F3410872.S26
F3410872.S27
F3410872.S28
F3410872.S30
F3410872.S48
F3410872.S59
F3410872.S63
F3410872.S69
F3410872.S72
F3410872.S74
F3410872.S80
F3410872.S82
F3410872.S86
F3410872.S93
F3410872.S97
F3410872.S98
0.00 0.25 0.50 0.75 1.00
0.00 0.25 0.50 0.75 1.00
Abundance
Species
Prevotella copri
Faecalibacterium prausnitzii
Phocaeicola vulgatus
Bacteroides uniformis
Akkermansia muciniphila
Phocaeicola dorei
Roseburia faecis
Escherichia_Pseudescherichia_Shigella boydii_coli_fergusonii_flexneri_marmotae_sonnei_vu ...(7 species)
Bacteroides faecichinchillae_faecis_thetaiotaomicron
Faecalibacterium prausnitzii_nov_97.209%
Oscillibacter valericigenes_nov_93.056%
Alistipes putredinis
Parasutterella excrementihominis
Massiliprevotella massiliensis
Monoglobus pectinilyticus_nov_90.698%
Eubacteriales_[G−1] bacterium_MOT−144
Ruminococcus champanellensis_nov_97.674%
Barnesiella intestinihominis
Subdoligranulum variabile
Ruminiclostridium cellobioparum_nov_92.056%
Blautia wexlerae
Alistipes onderdonkii
Oscillospiraceae_[G−6] bacterium_MOT−153_nov_97.674%
Parabacteroides merdae
Fusicatenibacter saccharivorans
Other