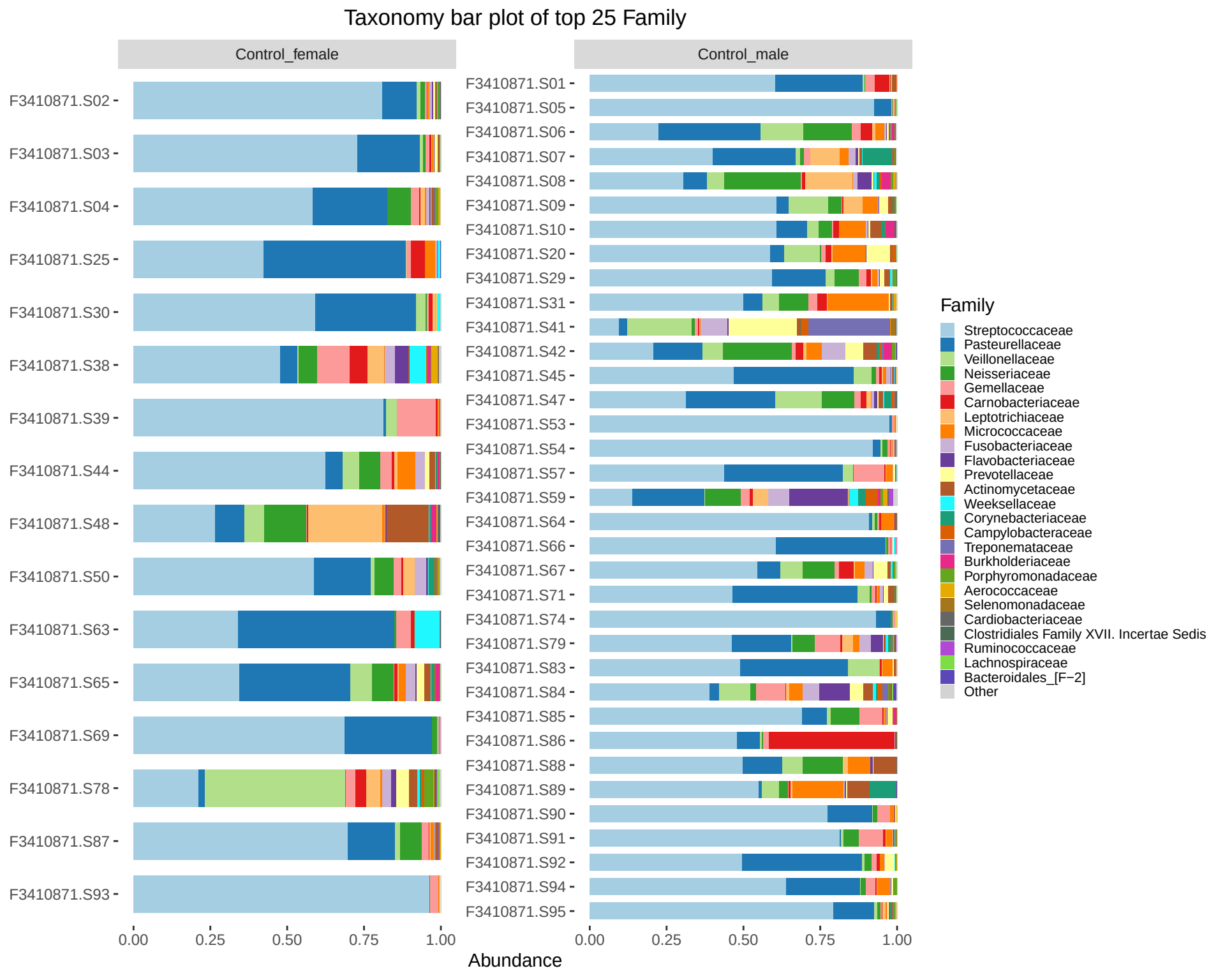
Taxonomy bar plot of top 25 Family
Control_female
Control_male
F3410871.S02
F3410871.S03
F3410871.S04
F3410871.S25
F3410871.S30
F3410871.S38
F3410871.S39
F3410871.S44
F3410871.S48
F3410871.S50
F3410871.S63
F3410871.S65
F3410871.S69
F3410871.S78
F3410871.S87
F3410871.S93
F3410871.S01
F3410871.S05
F3410871.S06
F3410871.S07
F3410871.S08
F3410871.S09
F3410871.S10
F3410871.S20
F3410871.S29
F3410871.S31
F3410871.S41
F3410871.S42
F3410871.S45
F3410871.S47
F3410871.S53
F3410871.S54
F3410871.S57
F3410871.S59
F3410871.S64
F3410871.S66
F3410871.S67
F3410871.S71
F3410871.S74
F3410871.S79
F3410871.S83
F3410871.S84
F3410871.S85
F3410871.S86
F3410871.S88
F3410871.S89
F3410871.S90
F3410871.S91
F3410871.S92
F3410871.S94
F3410871.S95
0.00
0.25
0.50
0.75
1.00
Abundance
Family
Streptococcaceae
Pasteurellaceae
Veillonellaceae
Neisseriaceae
Gemellaceae
Carnobacteriaceae
Leptotrichiaceae
Micrococcaceae
Fusobacteriaceae
Flavobacteriaceae
Prevotellaceae
Actinomycetaceae
Weeksellaceae
Corynebacteriaceae
Campylobacteraceae
Treponemataceae
Burkholderiaceae
Porphyromonadaceae
Aerococcaceae
Selenomonadaceae
Cardiobacteriaceae
Clostridiales Family XVII. Incertae Sedis
Ruminococcaceae
Lachnospiraceae
Bacteroidales_[F−2]
Other