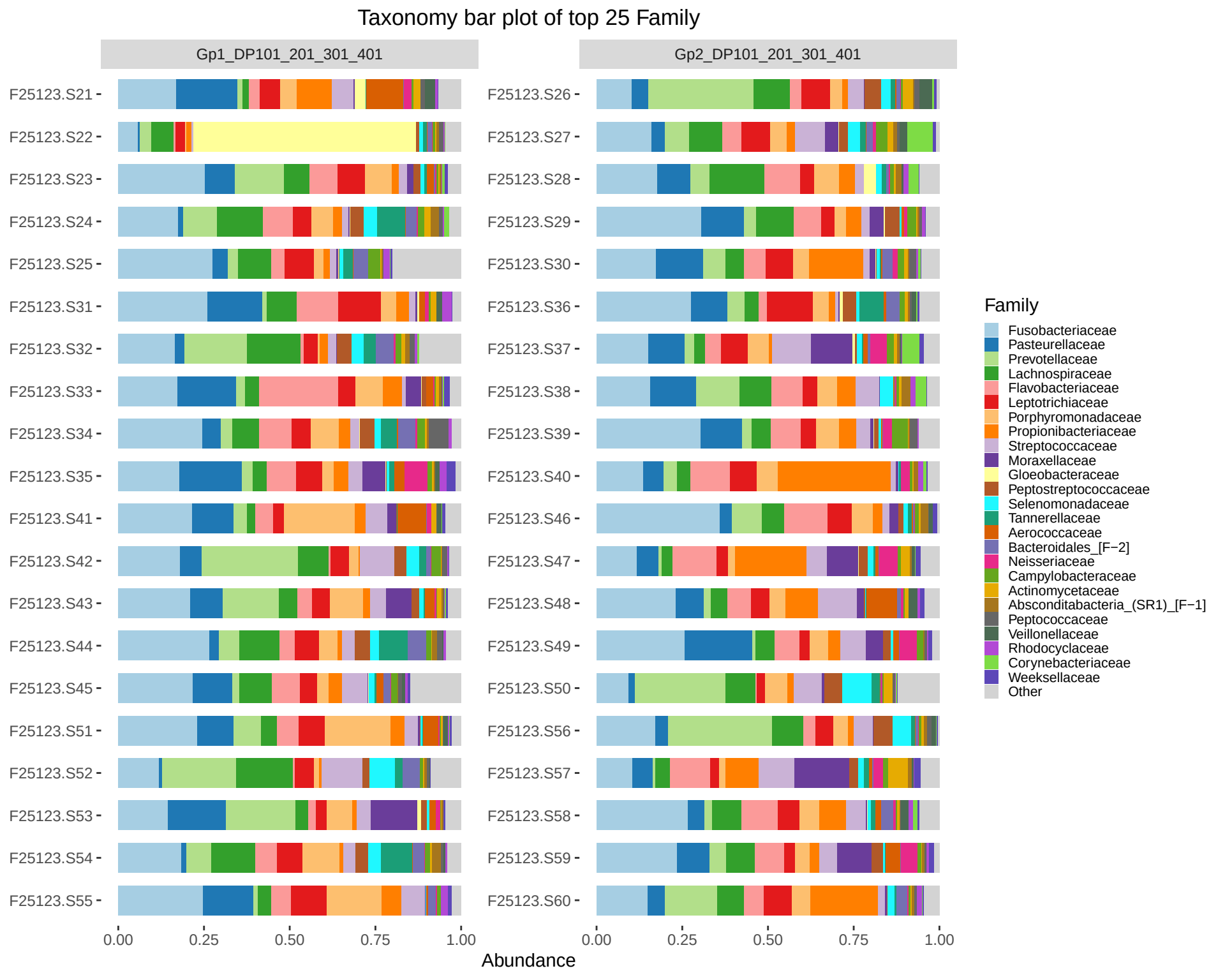
Taxonomy bar plot of top 25 Family
Gp1_DP101_201_301_401
Gp2_DP101_201_301_401
F25123.S21
F25123.S22
F25123.S23
F25123.S24
F25123.S25
F25123.S31
F25123.S32
F25123.S33
F25123.S34
F25123.S35
F25123.S41
F25123.S42
F25123.S43
F25123.S44
F25123.S45
F25123.S51
F25123.S52
F25123.S53
F25123.S54
F25123.S55
F25123.S26
F25123.S27
F25123.S28
F25123.S29
F25123.S30
F25123.S36
F25123.S37
F25123.S38
F25123.S39
F25123.S40
F25123.S46
F25123.S47
F25123.S48
F25123.S49
F25123.S50
F25123.S56
F25123.S57
F25123.S58
F25123.S59
F25123.S60
0.00
0.25
0.50
0.75
1.00
Abundance
Family
Fusobacteriaceae
Pasteurellaceae
Prevotellaceae
Lachnospiraceae
Flavobacteriaceae
Leptotrichiaceae
Porphyromonadaceae
Propionibacteriaceae
Streptococcaceae
Moraxellaceae
Gloeobacteraceae
Peptostreptococcaceae
Selenomonadaceae
Tannerellaceae
Aerococcaceae
Bacteroidales_[F−2]
Neisseriaceae
Campylobacteraceae
Actinomycetaceae
Absconditabacteria_(SR1)_[F−1]
Peptococcaceae
Veillonellaceae
Rhodocyclaceae
Corynebacteriaceae
Weeksellaceae
Other