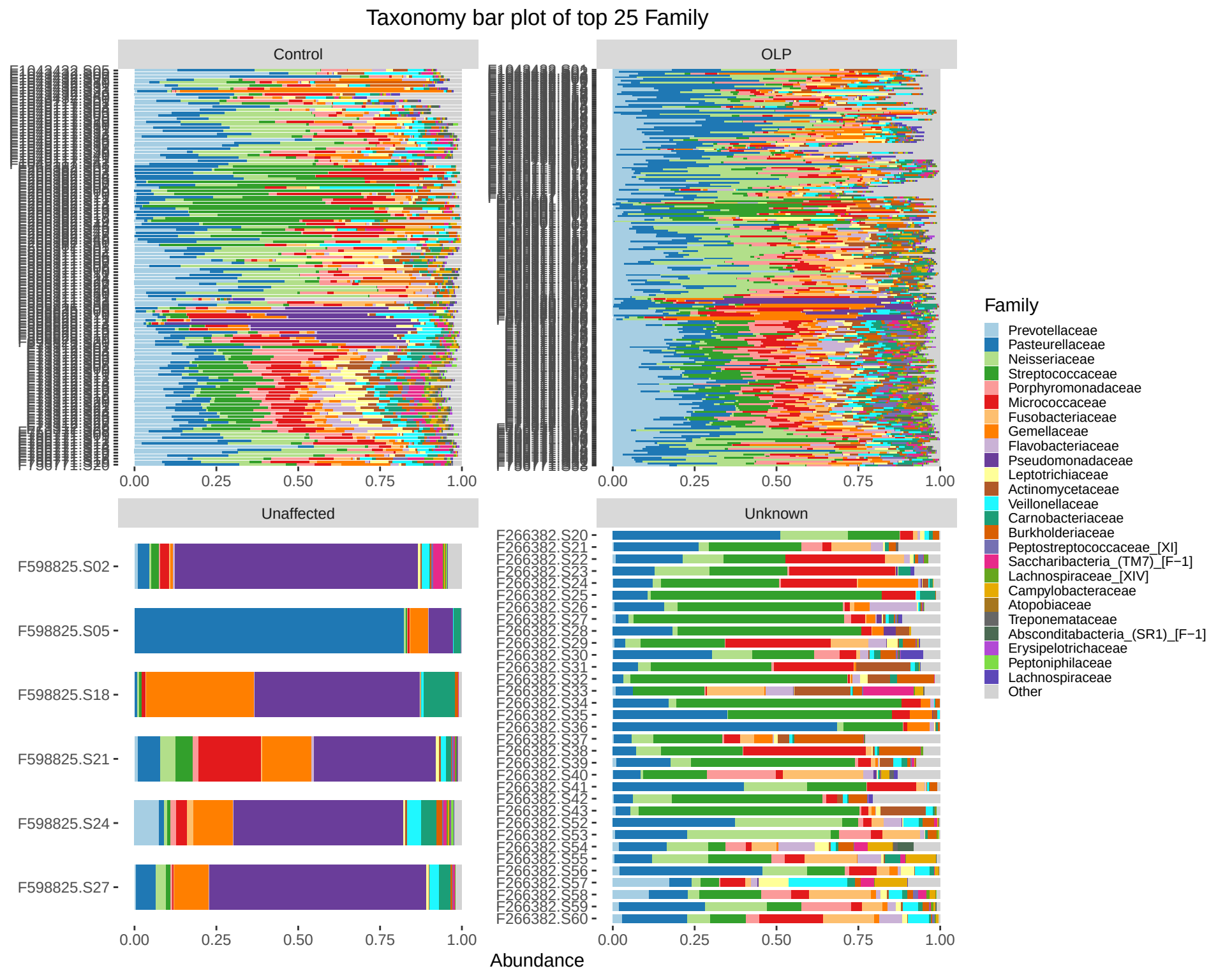
Taxonomy bar plot of top 25 Family
Control
OLP
Unaffected
Unknown
F598825.S02
F598825.S05
F598825.S18
F598825.S21
F598825.S24
F598825.S27
F266382.S20
F266382.S21
F266382.S22
F266382.S23
F266382.S24
F266382.S25
F266382.S26
F266382.S27
F266382.S28
F266382.S29
F266382.S30
F266382.S31
F266382.S32
F266382.S33
F266382.S34
F266382.S35
F266382.S36
F266382.S37
F266382.S38
F266382.S39
F266382.S40
F266382.S41
F266382.S42
F266382.S43
F266382.S52
F266382.S53
F266382.S54
F266382.S55
F266382.S56
F266382.S57
F266382.S58
F266382.S59
F266382.S60
0.00
0.25
0.50
0.75
1.00
Abundance
Family
Prevotellaceae
Pasteurellaceae
Neisseriaceae
Streptococcaceae
Porphyromonadaceae
Micrococcaceae
Fusobacteriaceae
Gemellaceae
Flavobacteriaceae
Pseudomonadaceae
Leptotrichiaceae
Actinomycetaceae
Veillonellaceae
Carnobacteriaceae
Burkholderiaceae
Peptostreptococcaceae_[XI]
Saccharibacteria_(TM7)_[F−1]
Lachnospiraceae_[XIV]
Campylobacteraceae
Atopobiaceae
Treponemataceae
Absconditabacteria_(SR1)_[F−1]
Erysipelotrichaceae
Peptoniphilaceae
Lachnospiraceae
Other