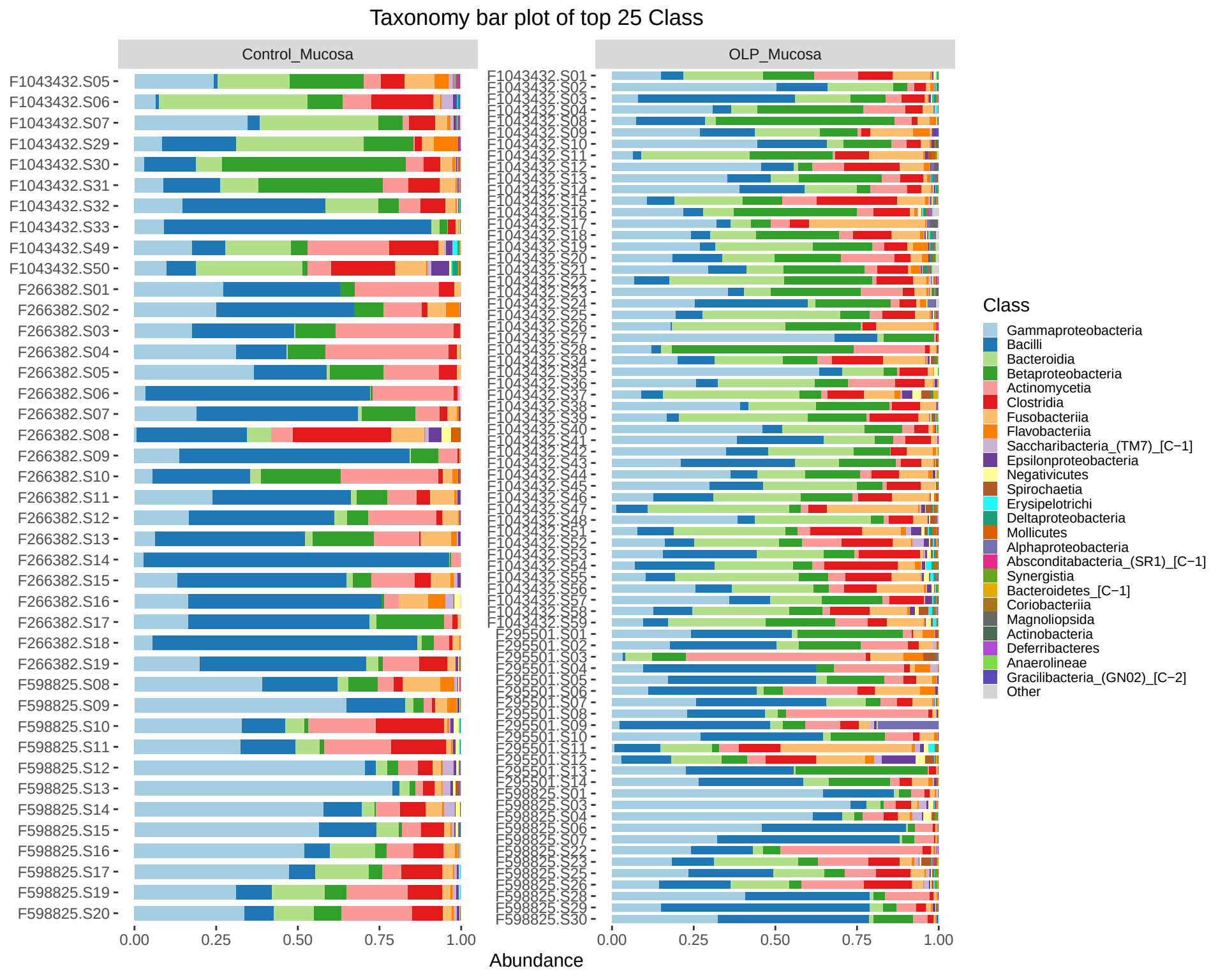
Taxonomy bar plot of top 25 Class
Control_Mucosa
OLP_Mucosa
F1043432.S05
F1043432.S06
F1043432.S07
F1043432.S29
F1043432.S30
F1043432.S31
F1043432.S32
F1043432.S33
F1043432.S49
F1043432.S50
F266382.S01
F266382.S02
F266382.S03
F266382.S04
F266382.S05
F266382.S06
F266382.S07
F266382.S08
F266382.S09
F266382.S10
F266382.S11
F266382.S12
F266382.S13
F266382.S14
F266382.S15
F266382.S16
F266382.S17
F266382.S18
F266382.S19
F598825.S08
F598825.S09
F598825.S10
F598825.S11
F598825.S12
F598825.S13
F598825.S14
F598825.S15
F598825.S16
F598825.S17
F598825.S19
F598825.S20
F1043432.S01
F1043432.S02
F1043432.S03
F1043432.S04
F1043432.S08
F1043432.S09
F1043432.S10
F1043432.S11
F1043432.S12
F1043432.S13
F1043432.S14
F1043432.S15
F1043432.S16
F1043432.S17
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F1043432.S19
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F1043432.S41
F1043432.S42
F1043432.S43
F1043432.S44
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F1043432.S46
F1043432.S47
F1043432.S48
F1043432.S51
F1043432.S52
F1043432.S53
F1043432.S54
F1043432.S55
F1043432.S56
F1043432.S57
F1043432.S58
F1043432.S59
F295501.S01
F295501.S02
F295501.S03
F295501.S04
F295501.S05
F295501.S06
F295501.S07
F295501.S08
F295501.S09
F295501.S10
F295501.S11
F295501.S12
F295501.S13
F295501.S14
F598825.S01
F598825.S03
F598825.S04
F598825.S06
F598825.S07
F598825.S22
F598825.S23
F598825.S25
F598825.S26
F598825.S28
F598825.S29
F598825.S30
0.00
0.25
0.50
0.75
1.00
Abundance
Class
Gammaproteobacteria
Bacilli
Bacteroidia
Betaproteobacteria
Actinomycetia
Clostridia
Fusobacteriia
Flavobacteriia
Saccharibacteria_(TM7)_[C−1]
Epsilonproteobacteria
Negativicutes
Spirochaetia
Erysipelotrichi
Deltaproteobacteria
Mollicutes
Alphaproteobacteria
Absconditabacteria_(SR1)_[C−1]
Synergistia
Bacteroidetes_[C−1]
Coriobacteriia
Magnoliopsida
Actinobacteria
Deferribacteres
Anaerolineae
Gracilibacteria_(GN02)_[C−2]
Other