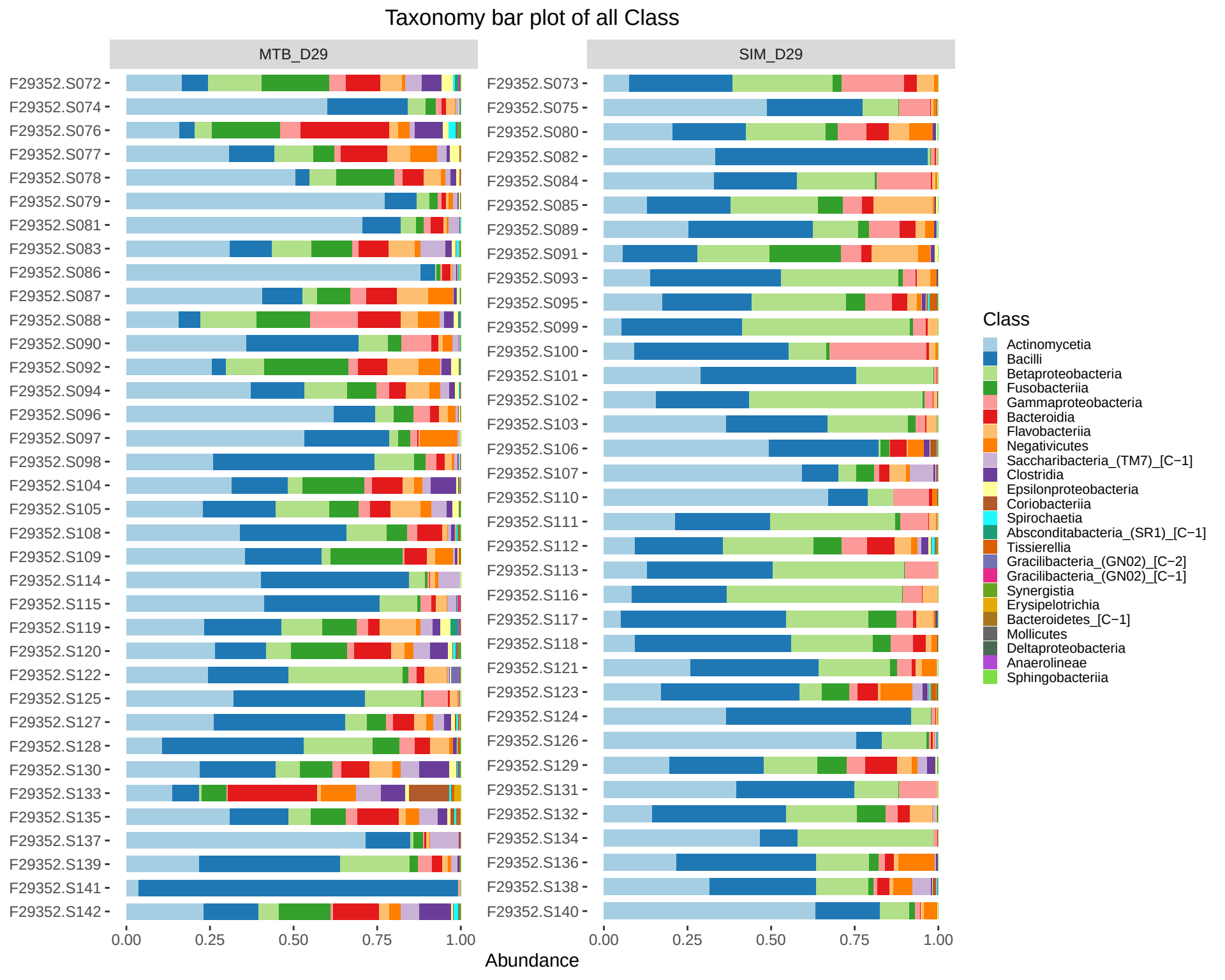
Taxonomy bar plot of all Class
MTB_D29
SIM_D29
F29352.S072
F29352.S074
F29352.S076
F29352.S077
F29352.S078
F29352.S079
F29352.S081
F29352.S083
F29352.S086
F29352.S087
F29352.S088
F29352.S090
F29352.S092
F29352.S094
F29352.S096
F29352.S097
F29352.S098
F29352.S104
F29352.S105
F29352.S108
F29352.S109
F29352.S114
F29352.S115
F29352.S119
F29352.S120
F29352.S122
F29352.S125
F29352.S127
F29352.S128
F29352.S130
F29352.S133
F29352.S135
F29352.S137
F29352.S139
F29352.S141
F29352.S142
F29352.S073
F29352.S075
F29352.S080
F29352.S082
F29352.S084
F29352.S085
F29352.S089
F29352.S091
F29352.S093
F29352.S095
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F29352.S116
F29352.S117
F29352.S118
F29352.S121
F29352.S123
F29352.S124
F29352.S126
F29352.S129
F29352.S131
F29352.S132
F29352.S134
F29352.S136
F29352.S138
F29352.S140
0.00
0.25
0.50
0.75
1.00
Abundance
Class
Actinomycetia
Bacilli
Betaproteobacteria
Fusobacteriia
Gammaproteobacteria
Bacteroidia
Flavobacteriia
Negativicutes
Saccharibacteria_(TM7)_[C−1]
Clostridia
Epsilonproteobacteria
Coriobacteriia
Spirochaetia
Absconditabacteria_(SR1)_[C−1]
Tissierellia
Gracilibacteria_(GN02)_[C−2]
Gracilibacteria_(GN02)_[C−1]
Synergistia
Erysipelotrichia
Bacteroidetes_[C−1]
Mollicutes
Deltaproteobacteria
Anaerolineae
Sphingobacteriia