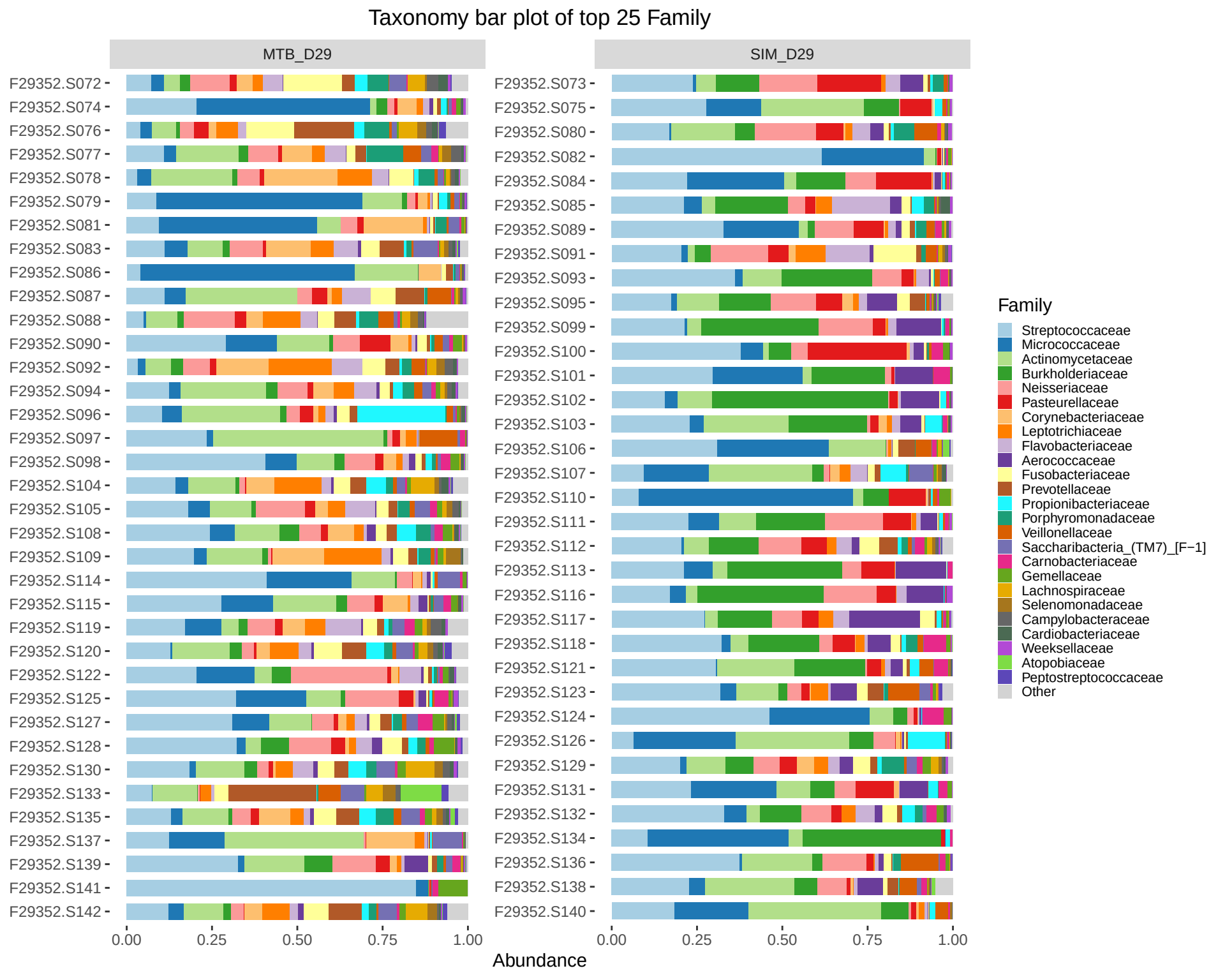
Taxonomy bar plot of top 25 Family
MTB_D29
SIM_D29
F29352.S072
F29352.S074
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F29352.S078
F29352.S079
F29352.S081
F29352.S083
F29352.S086
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F29352.S088
F29352.S090
F29352.S092
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F29352.S096
F29352.S097
F29352.S098
F29352.S104
F29352.S105
F29352.S108
F29352.S109
F29352.S114
F29352.S115
F29352.S119
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F29352.S122
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F29352.S127
F29352.S128
F29352.S130
F29352.S133
F29352.S135
F29352.S137
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F29352.S141
F29352.S142
F29352.S073
F29352.S075
F29352.S080
F29352.S082
F29352.S084
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F29352.S095
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F29352.S110
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F29352.S118
F29352.S121
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F29352.S134
F29352.S136
F29352.S138
F29352.S140
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0.25
0.50
0.75
1.00
Abundance
Family
Streptococcaceae
Micrococcaceae
Actinomycetaceae
Burkholderiaceae
Neisseriaceae
Pasteurellaceae
Corynebacteriaceae
Leptotrichiaceae
Flavobacteriaceae
Aerococcaceae
Fusobacteriaceae
Prevotellaceae
Propionibacteriaceae
Porphyromonadaceae
Veillonellaceae
Saccharibacteria_(TM7)_[F−1]
Carnobacteriaceae
Gemellaceae
Lachnospiraceae
Selenomonadaceae
Campylobacteraceae
Cardiobacteriaceae
Weeksellaceae
Atopobiaceae
Peptostreptococcaceae
Other