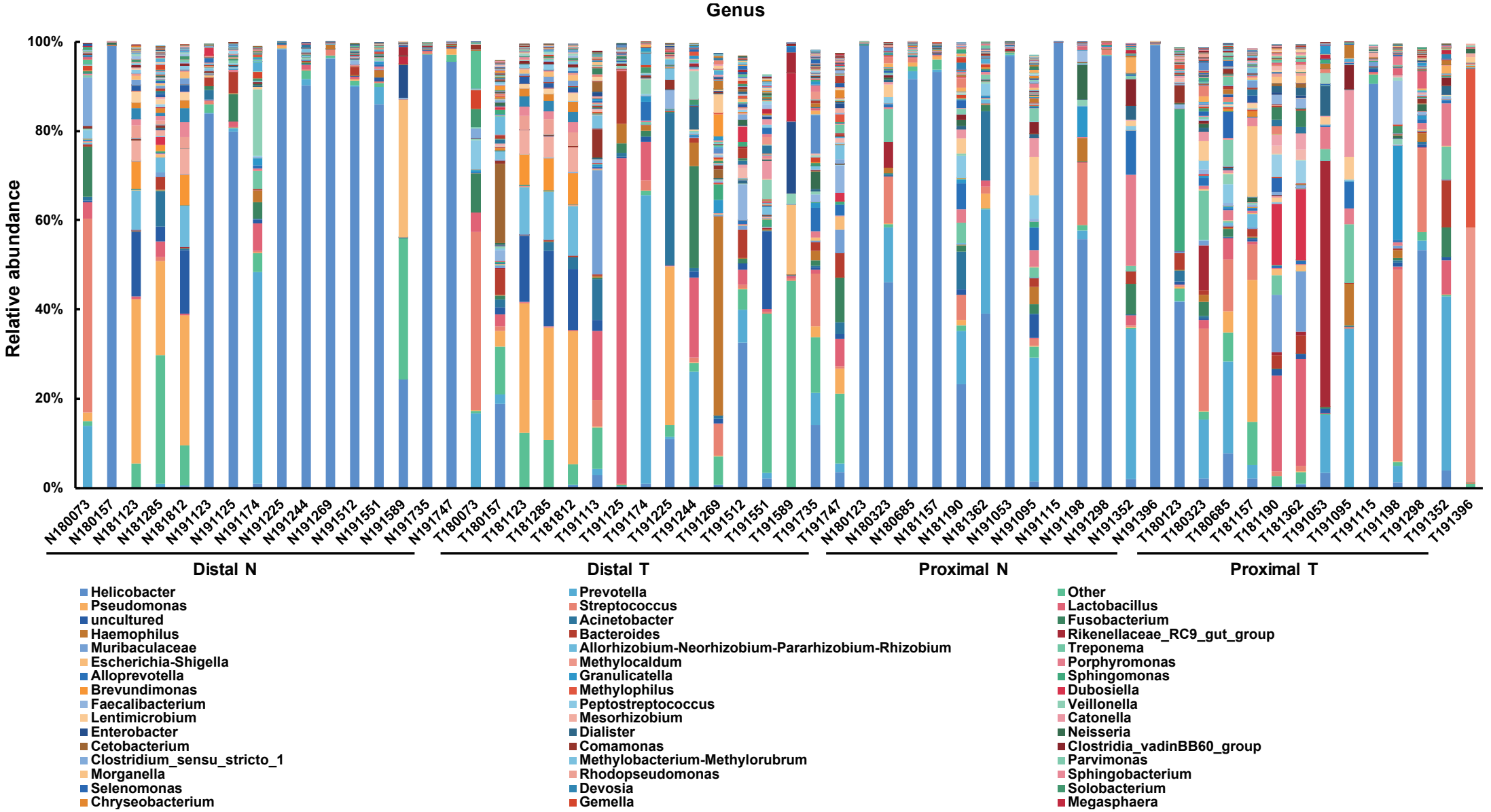
Genus
Relative abundance
100%
80%
60%
40%
20%
0%
N180073
N180157
N181123
N181285
N181812
N191123
N191125
N191174
N191225
N191244
N191269
N191512
N191551
N191589
N191735
N191747
T180073
T180157
T181123
T181285
T181812
T191113
T191125
T191174
T191225
T191244
T191269
T191512
T191551
T191589
T191735
T191747
N180123
N180323
N180685
N181157
N181190
N181362
N191053
N191095
N191115
N191198
N191298
N191352
N191396
T180123
T180323
T180685
T181157
T181190
T181362
T191053
T191095
T191115
T191198
T191298
T191352
T191396
Distal N
Distal T
Proximal N
Proximal T
Helicobacter
Pseudomonas
uncultured
Haemophilus
Muribaculaceae
Escherichia-Shigella
Alloprevotella
Brevundimonas
Faecalibacterium
Lentimicrobium
Enterobacter
Cetobacterium
Clostridium_sensu_stricto_1
Morganella
Selenomonas
Chryseobacterium
Prevotella
Streptococcus
Acinetobacter
Bacteroides
Allorhizobium-Neorhizobium-Pararhizobium-Rhizobium
Methylocaldum
Granulicatella
Methylophilus
Peptostreptococcus
Mesorhizobium
Dialister
Comamonas
Methylobacterium-Methylorubrum
Rhodopseudomonas
Devosia
Gemella
Other
Lactobacillus
Fusobacterium
Rikenellaceae_RC9_gut_group
Treponema
Porphyromonas
Sphingomonas
Dubosiella
Veillonella
Catonella
Neisseria
Clostridia_vadinBB60_group
Parvimonas
Sphingobacterium
Solobacterium
Megasphaera