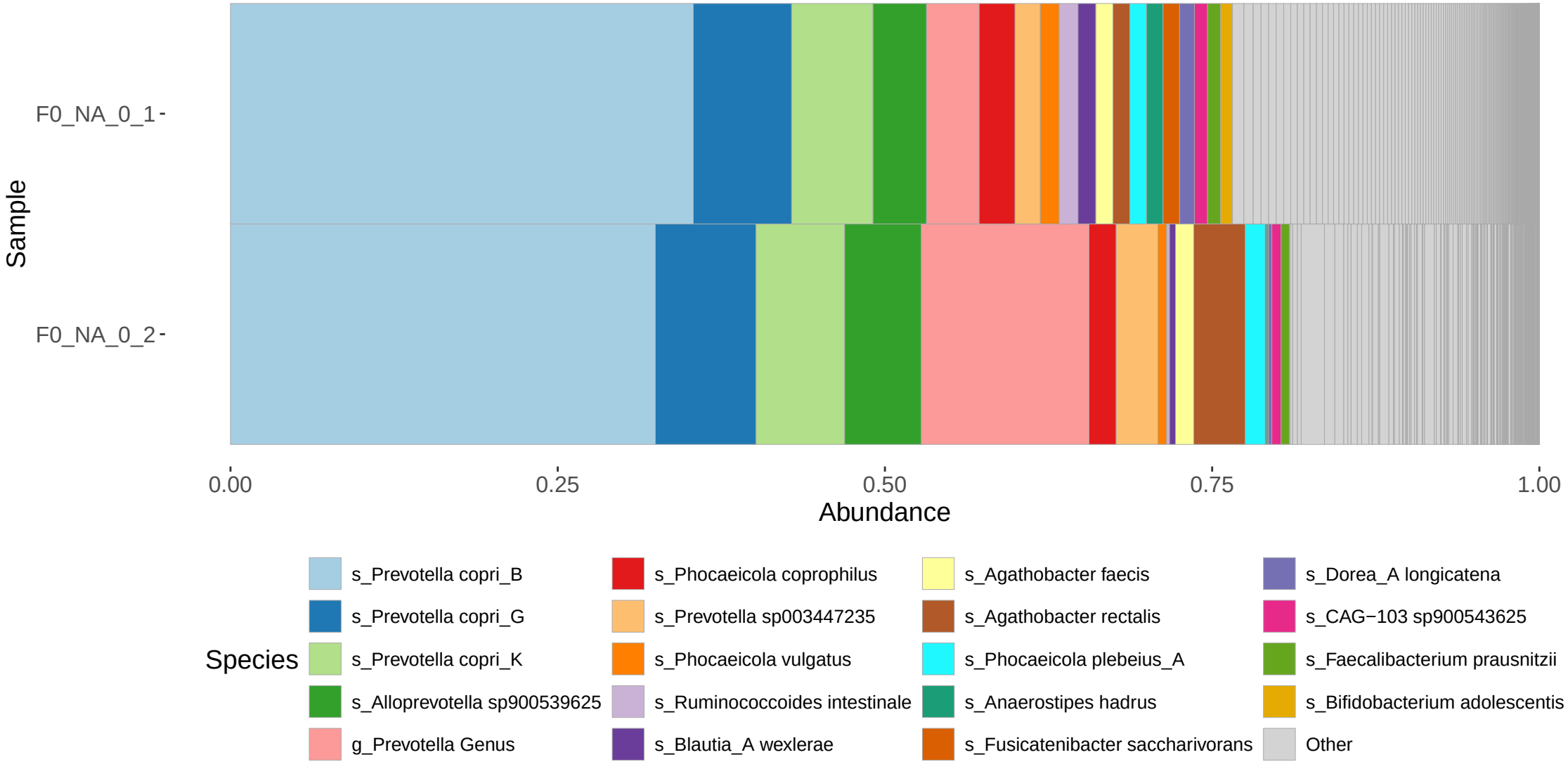
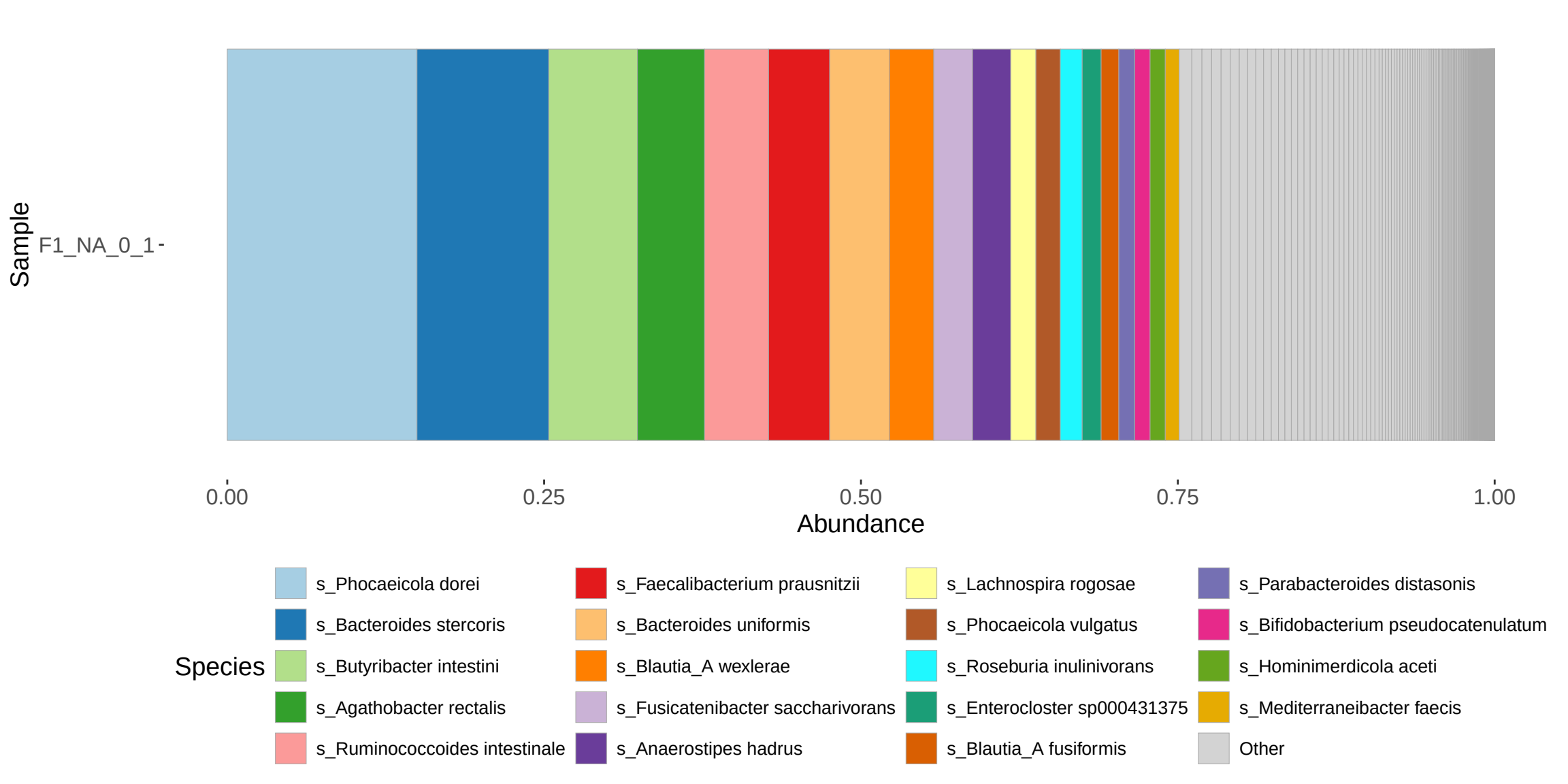
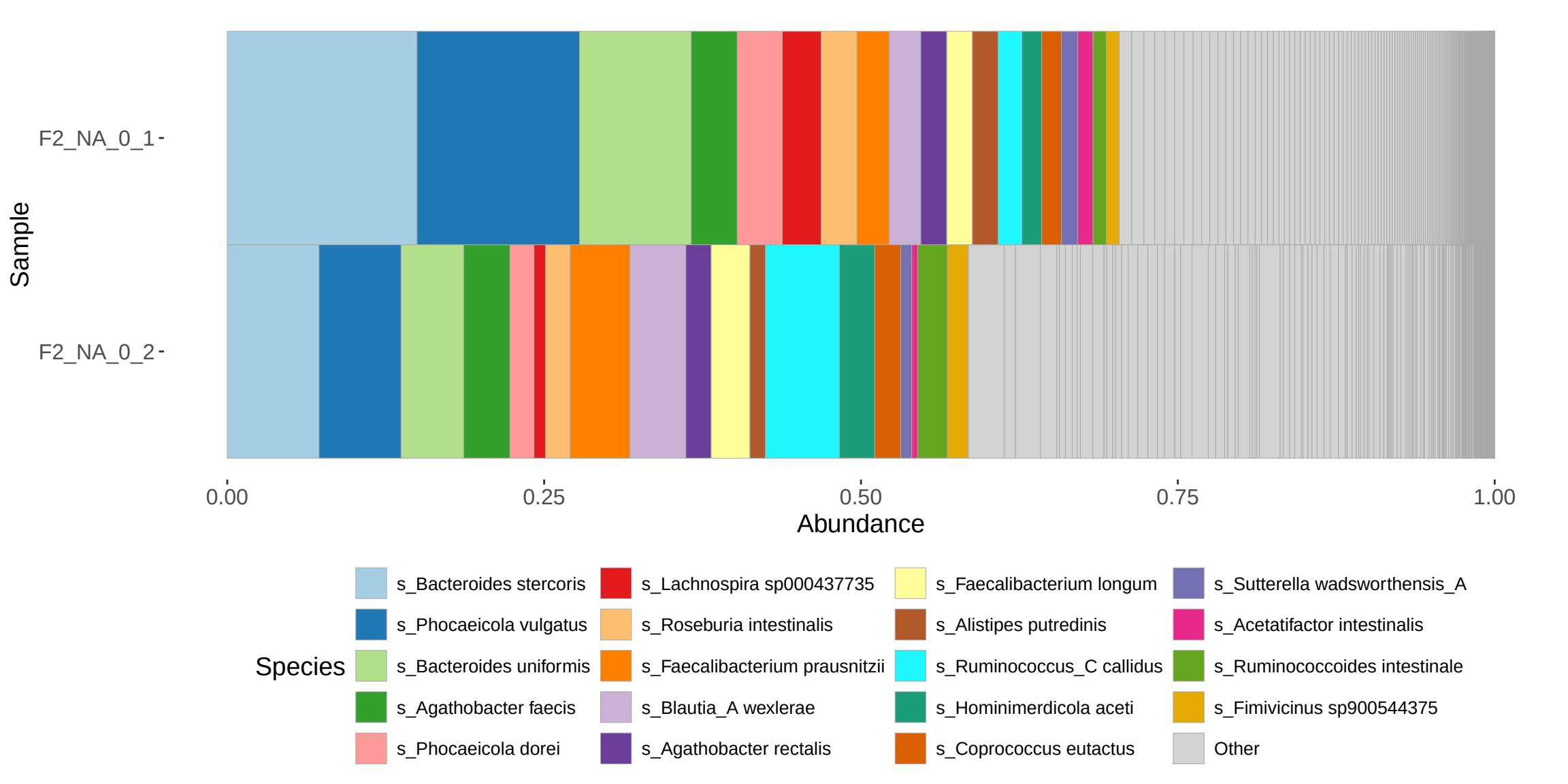
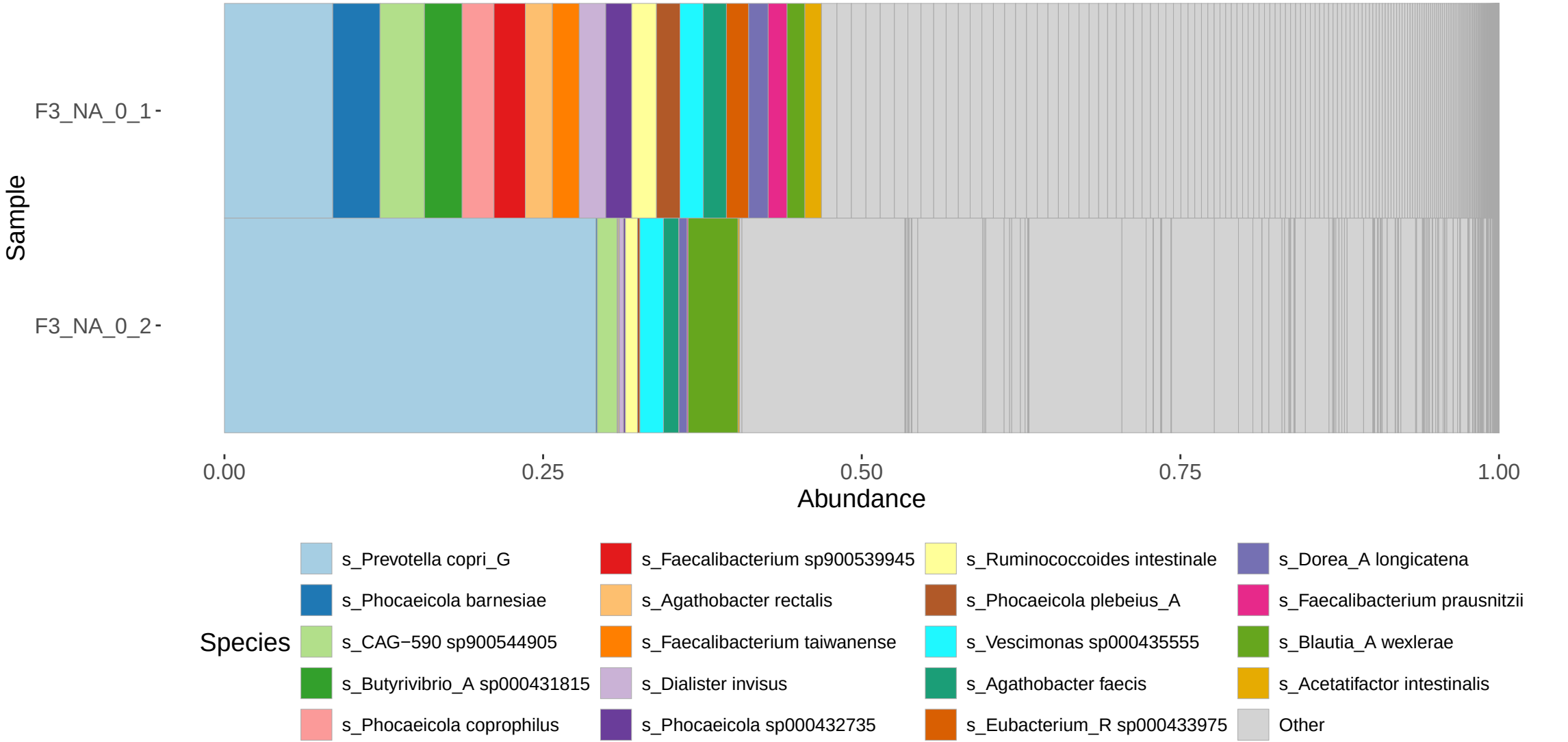
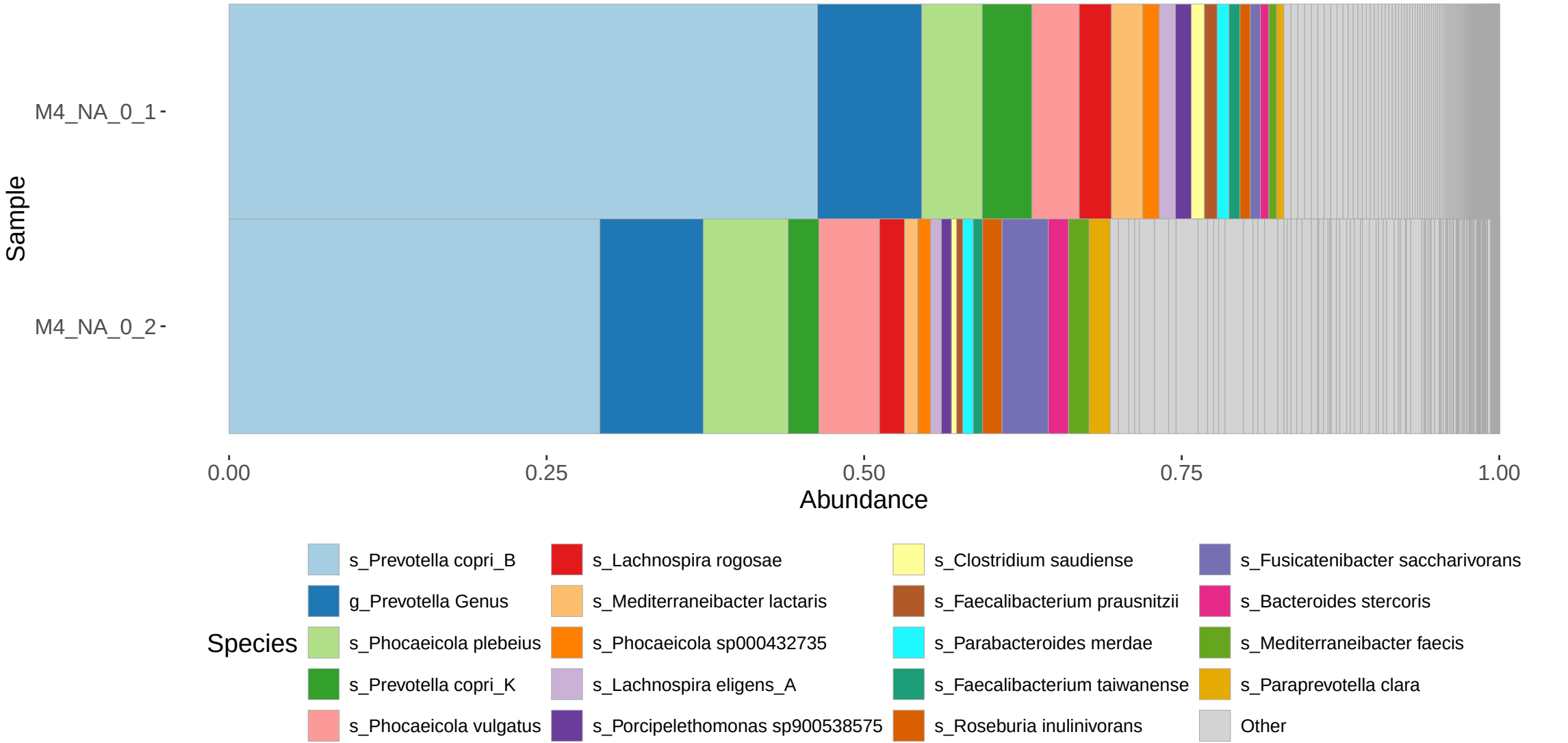
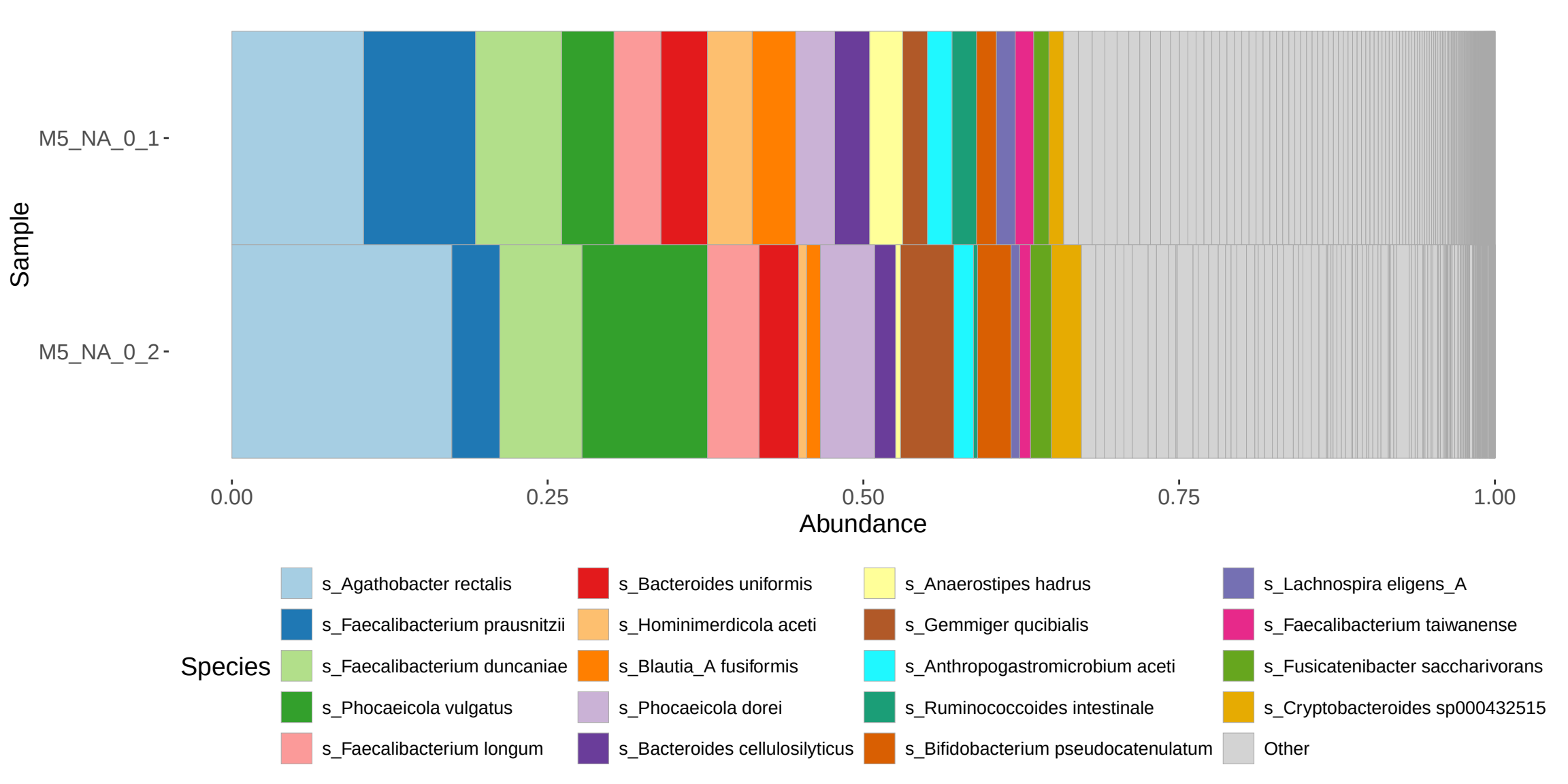

Sample
F0_NA_0_1
F0_NA_0_2
0.00
0.25
0.50
0.75
1.00
Abundance
Species
s_Prevotella copri_B
s_Prevotella copri_G
s_Prevotella copri_K
s_Alloprevotella sp900539625
g_Prevotella Genus
s_Phocaeicola coprophilus
s_Prevotella sp003447235
s_Phocaeicola vulgatus
s_Ruminococcoides intestinale
s_Blautia_A wexlerae
s_Agathobacter faecis
s_Agathobacter rectalis
s_Phocaeicola plebeius_A
s_Anaerostipes hadrus
s_Fusicatenibacter saccharivorans
s_Dorea_A longicatena
s_CAG−103 sp900543625
s_Faecalibacterium prausnitzii
s_Bifidobacterium adolescentis
Other

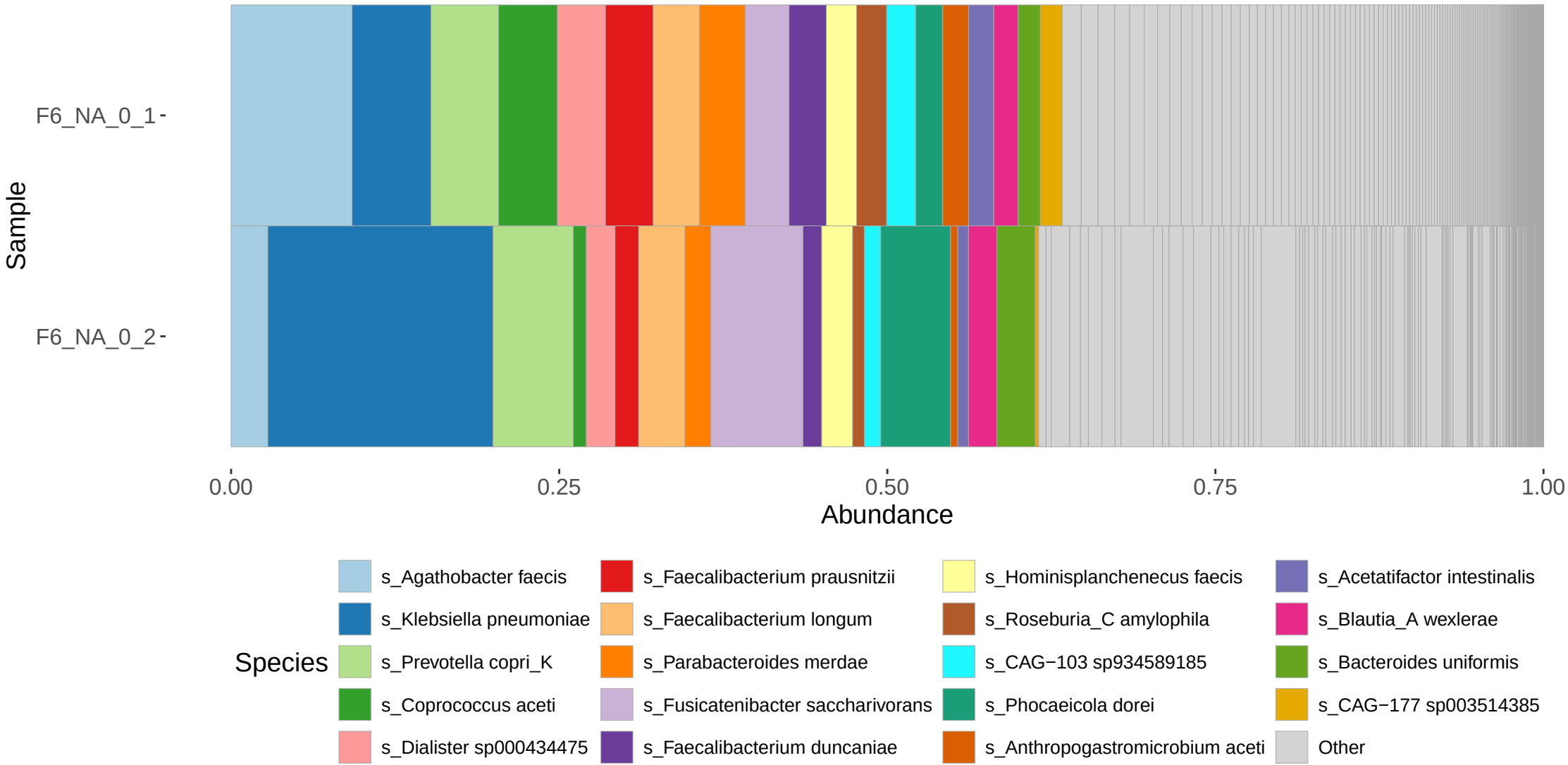

Sample
F1_NA_0_1
0.00
0.25
0.50
0.75
1.00
Abundance
Species
s_Phocaeicola dorei
s_Bacteroides stercoris
s_Butyribacter intestini
s_Agathobacter rectalis
s_Ruminococcoides intestinale
s_Faecalibacterium prausnitzii
s_Bacteroides uniformis
s_Blautia_A wexlerae
s_Fusicatenibacter saccharivorans
s_Anaerostipes hadrus
s_Lachnospira rogosae
s_Phocaeicola vulgatus
s_Roseburia inulinivorans
s_Enterocloster sp000431375
s_Blautia_A fusiformis
s_Parabacteroides distasonis
s_Bifidobacterium pseudocatenulatum
s_Hominimerdicola aceti
s_Mediterraneibacter faecis
Other

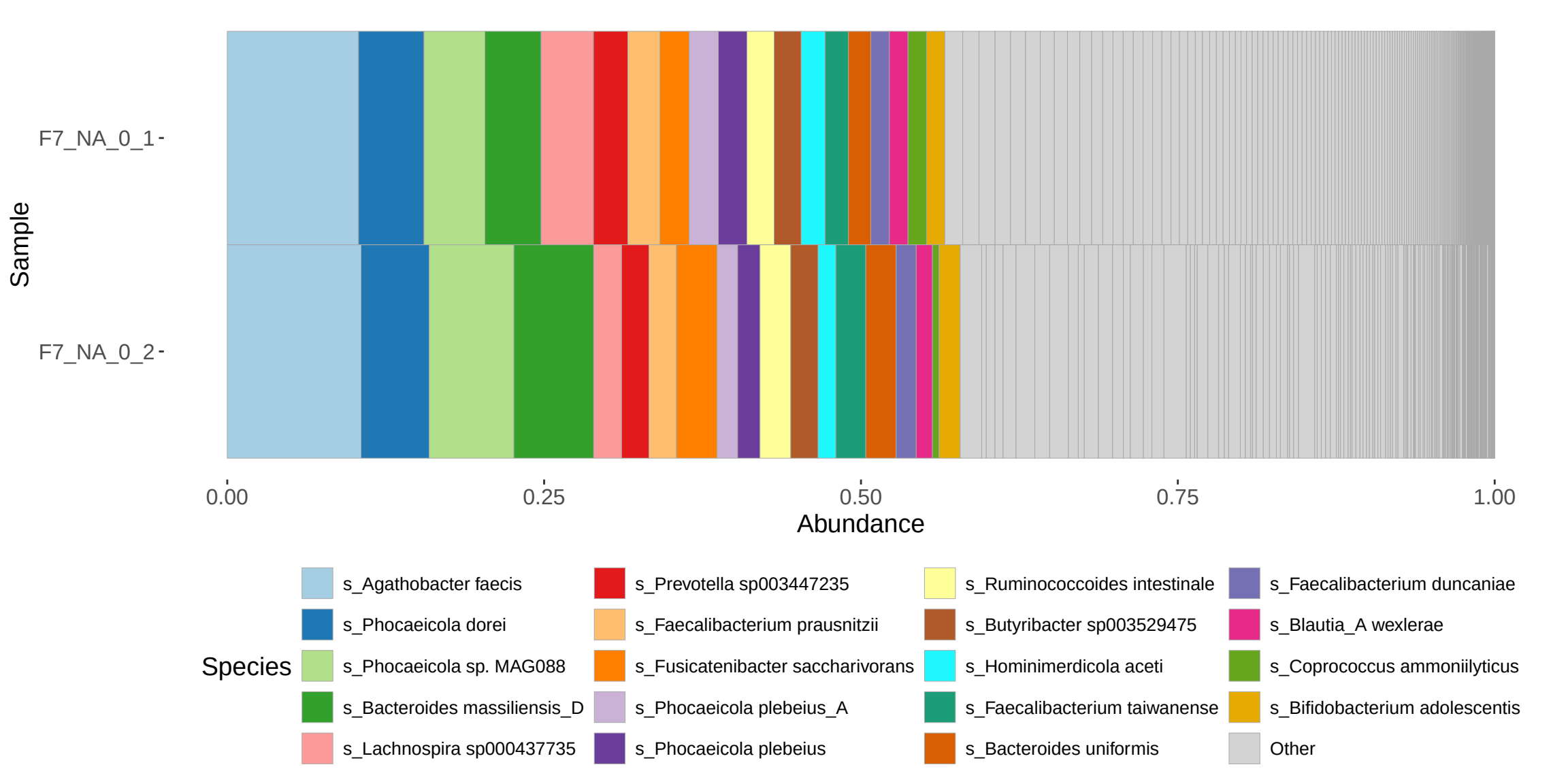

F2_NA_0_1
F2_NA_0_2
Sample
0.00
0.25
0.50
0.75
1.00
Abundance
Species
s_Bacteroides stercoris
s_Phocaeicola vulgatus
s_Bacteroides uniformis
s_Agathobacter faecis
s_Phocaeicola dorei
s_Lachnospira sp000437735
s_Roseburia intestinalis
s_Faecalibacterium prausnitzii
s_Blautia_A wexlerae
s_Agathobacter rectalis
s_Faecalibacterium longum
s_Alistipes putredinis
s_Ruminococcus_C callidus
s_Hominimerdicola aceti
s_Coprococcus eutactus
s_Sutterella wadsworthensis_A
s_Acetatifactor intestinalis
s_Ruminococcoides intestinale
s_Fimivicinus sp900544375
Other

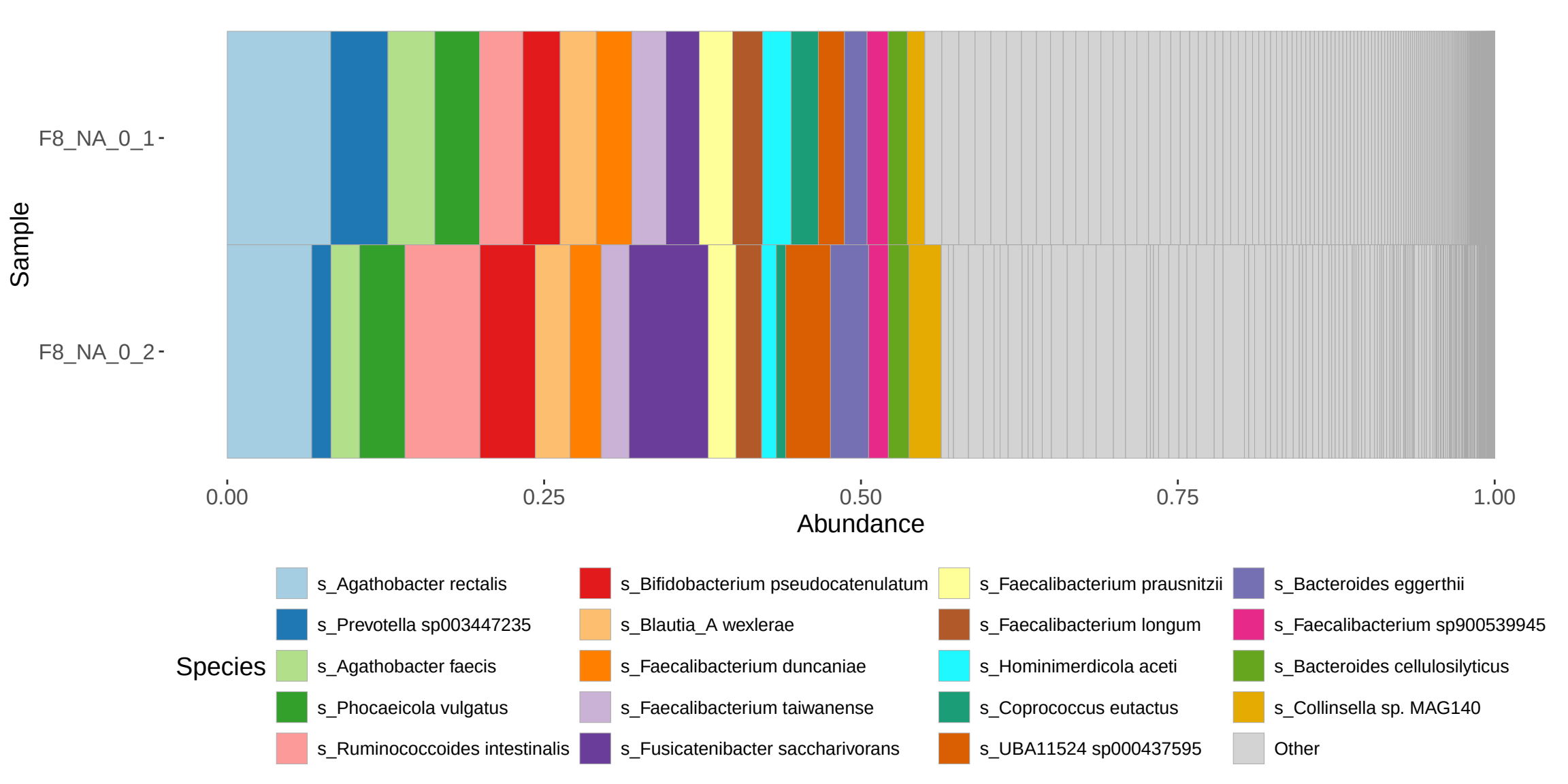

Sample
F3_NA_0_1
F3_NA_0_2
0.00
0.25
0.50
0.75
1.00
Abundance
Species
s_Prevotella copri_G
s_Phocaeicola barnesiae
s_CAG−590 sp900544905
s_Butyrivibrio_A sp000431815
s_Phocaeicola coprophilus
s_Faecalibacterium sp900539945
s_Agathobacter rectalis
s_Faecalibacterium taiwanense
s_Dialister invisus
s_Phocaeicola sp000432735
s_Ruminococcoides intestinale
s_Phocaeicola plebeius_A
s_Vescimonas sp000435555
s_Agathobacter faecis
s_Eubacterium_R sp000433975
s_Dorea_A longicatena
s_Faecalibacterium prausnitzii
s_Blautia_A wexlerae
s_Acetatifactor intestinalis
Other

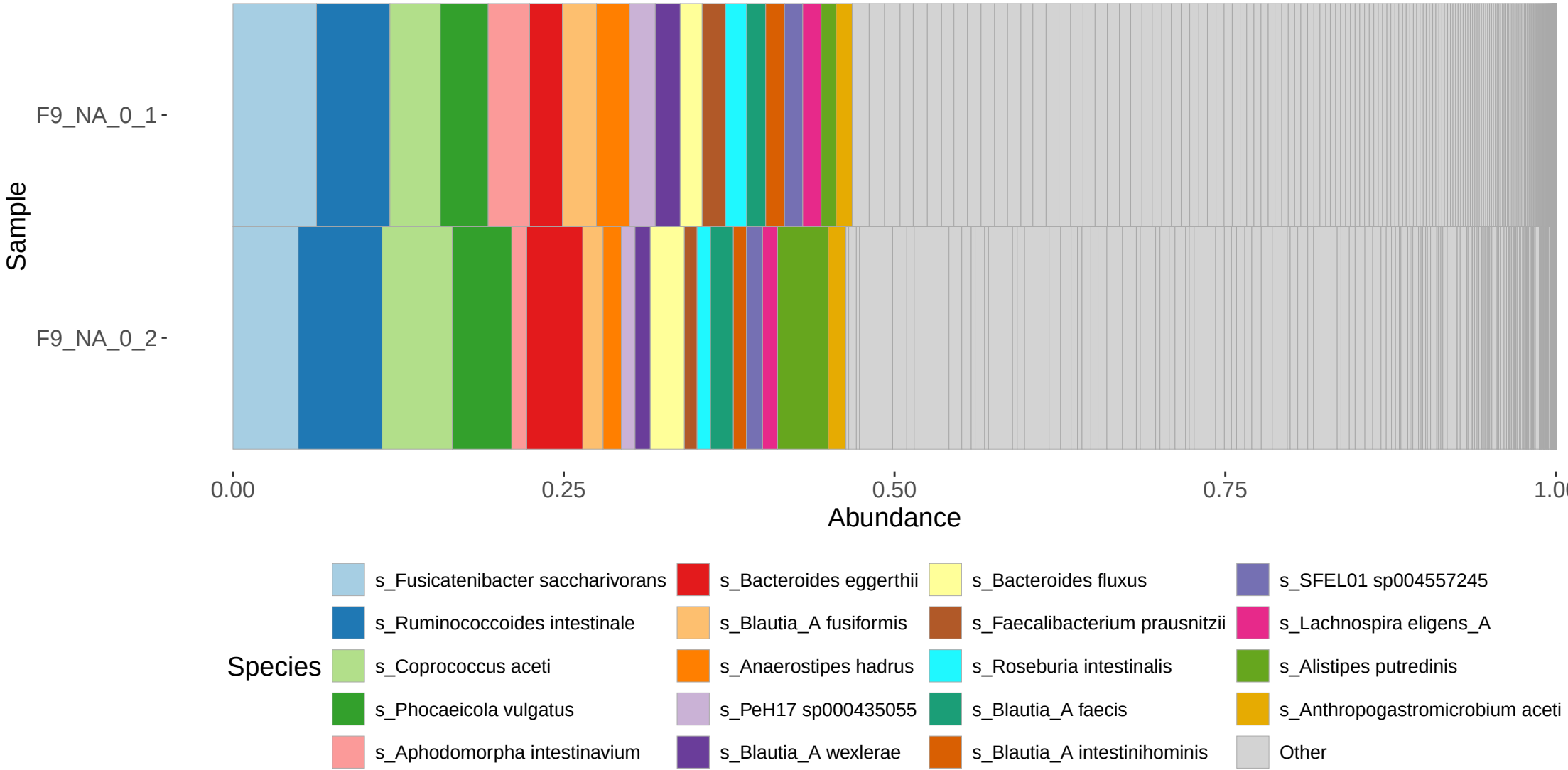

Sample
M4_NA_0_1
M4_NA_0_2
0.00
0.25
0.50
0.75
1.00
Abundance
Species
s_Prevotella copri_B
g_Prevotella Genus
s_Phocaeicola plebeius
s_Prevotella copri_K
s_Phocaeicola vulgatus
s_Lachnospira rogosae
s_Mediterraneibacter lactaris
s_Phocaeicola sp000432735
s_Lachnospira eligens_A
s_Porcipelethomonas sp900538575
s_Clostridium saudiense
s_Faecalibacterium prausnitzii
s_Parabacteroides merdae
s_Faecalibacterium taiwanense
s_Roseburia inulinivorans
s_Fusicatenibacter saccharivorans
s_Bacteroides stercoris
s_Mediterraneibacter faecis
s_Paraprevotella clara
Other


M5_NA_0_1
M5_NA_0_2
Sample
0.00
0.25
0.50
0.75
1.00
Abundance
Species
s_Agathobacter rectalis
s_Faecalibacterium prausnitzii
s_Faecalibacterium duncaniae
s_Phocaeicola vulgatus
s_Faecalibacterium longum
s_Bacteroides uniformis
s_Hominimerdicola aceti
s_Blautia_A fusiformis
s_Phocaeicola dorei
s_Bacteroides cellulosilyticus
s_Anaerostipes hadrus
s_Gemmiger qucibialis
s_Anthropogastromicrobium aceti
s_Ruminococcoides intestinale
s_Bifidobacterium pseudocatenulatum
s_Lachnospira eligens_A
s_Faecalibacterium taiwanense
s_Fusicatenibacter saccharivorans
s_Cryptobacteroides sp000432515
Other


Sample
F6_NA_0_1
F6_NA_0_2
0.00
0.25
0.50
0.75
1.00
Abundance
Species
s_Agathobacter faecis
s_Klebsiella pneumoniae
s_Prevotella copri_K
s_Coprococcus aceti
s_Dialister sp000434475
s_Faecalibacterium prausnitzii
s_Faecalibacterium longum
s_Parabacteroides merdae
s_Fusicatenibacter saccharivorans
s_Faecalibacterium duncaniae
s_Hominisplanchenecus faecis
s_Roseburia_C amylophila
s_CAG−103 sp934589185
s_Phocaeicola dorei
s_Anthropogastromicrobium aceti
s_Acetatifactor intestinalis
s_Blautia_A wexlerae
s_Bacteroides uniformis
s_CAG−177 sp003514385
Other


F7_NA_0_1
F7_NA_0_2
Sample
0.00
0.25
0.50
0.75
1.00
Abundance
Species
s_Agathobacter faecis
s_Phocaeicola dorei
s_Phocaeicola sp. MAG088
s_Bacteroides massiliensis_D
s_Lachnospira sp000437735
s_Prevotella sp003447235
s_Faecalibacterium prausnitzii
s_Fusicatenibacter saccharivorans
s_Phocaeicola plebeius_A
s_Phocaeicola plebeius
s_Ruminococcoides intestinale
s_Butyribacter sp003529475
s_Hominimerdicola aceti
s_Faecalibacterium taiwanense
s_Bacteroides uniformis
s_Faecalibacterium duncaniae
s_Blautia_A wexlerae
s_Coprococcus ammoniilyticus
s_Bifidobacterium adolescentis
Other


Sample
F8_NA_0_1
F8_NA_0_2
0.00
0.25
0.50
0.75
1.00
Abundance
Species
s_Agathobacter rectalis
s_Prevotella sp003447235
s_Agathobacter faecis
s_Phocaeicola vulgatus
s_Ruminococcoides intestinalis
s_Bifidobacterium pseudocatenulatum
s_Blautia_A wexlerae
s_Faecalibacterium duncaniae
s_Faecalibacterium taiwanense
s_Fusicatenibacter saccharivorans
s_Faecalibacterium prausnitzii
s_Faecalibacterium longum
s_Hominimerdicola aceti
s_Coprococcus eutactus
s_UBA11524 sp000437595
s_Bacteroides eggerthii
s_Faecalibacterium sp900539945
s_Bacteroides cellulosilyticus
s_Collinsella sp. MAG140
Other


Sample
F9_NA_0_1
F9_NA_0_2
0.00
0.25
0.50
0.75
1.00
Abundance
Species
s_Fusicatenibacter saccharivorans
s_Ruminococcoides intestinale
s_Coprococcus aceti
s_Phocaeicola vulgatus
s_Aphodomorpha intestinavium
s_Bacteroides eggerthii
s_Blautia_A fusiformis
s_Anaerostipes hadrus
s_PeH17 sp000435055
s_Blautia_A wexlerae
s_Bacteroides fluxus
s_Faecalibacterium prausnitzii
s_Roseburia intestinalis
s_Blautia_A faecis
s_Blautia_A intestinihominis
s_SFEL01 sp004557245
s_Lachnospira eligens_A
s_Alistipes putredinis
s_Anthropogastromicrobium aceti
Other