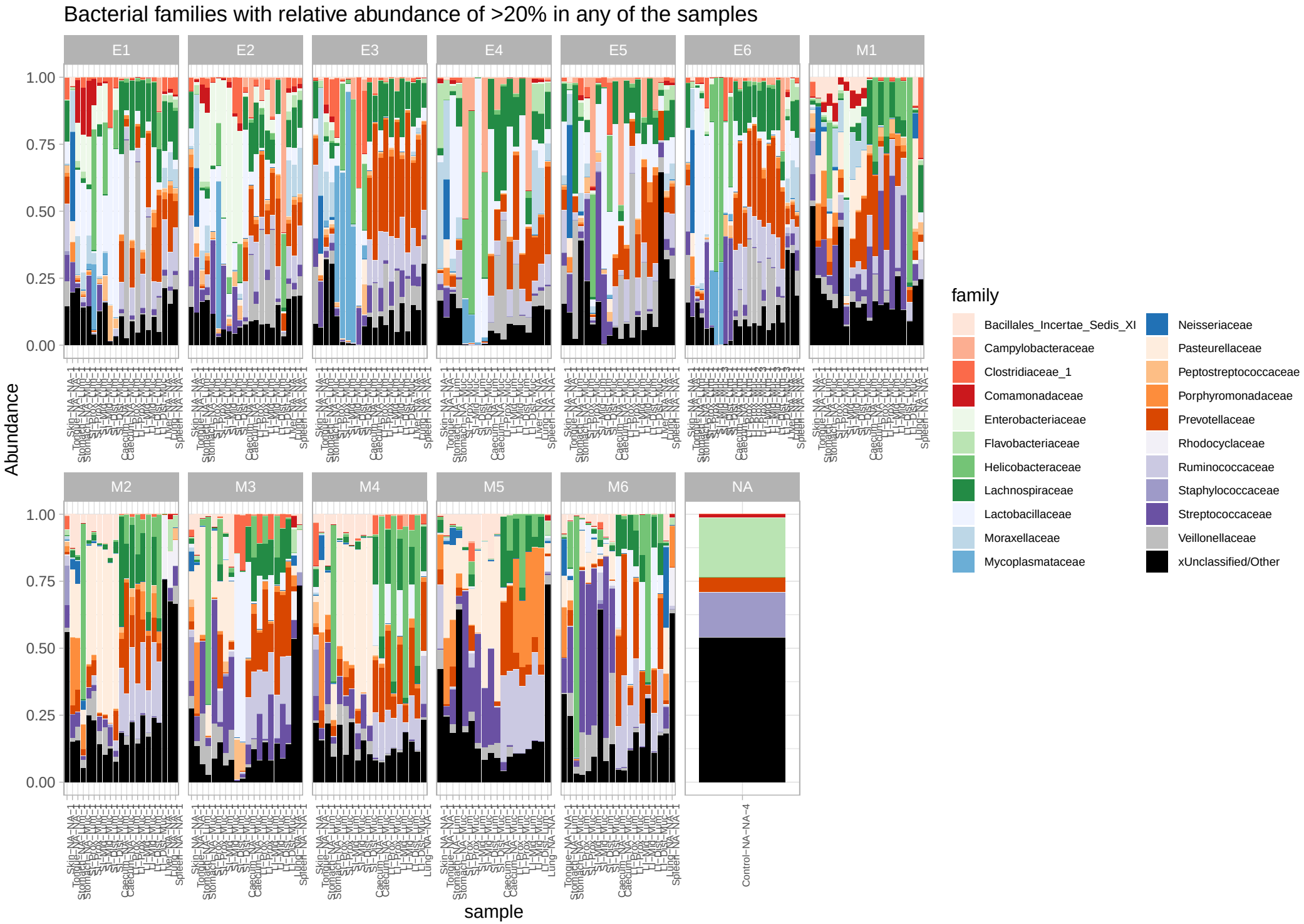
Bacterial families with relative abundance of >20% in any of the samples
E1
E2
E3
E4
E5
E6
M1
M2
M3
M4
M5
M6
NA
Abundance
1.00
0.75
0.50
0.25
0.00
sample
Control−NA−4
family
Bacillales_Incertae_Sedis_XI
Campylobacteraceae
Clostridiaceae_1
Comamonadaceae
Enterobacteriaceae
Flavobacteriaceae
Helicobacteraceae
Lachnospiraceae
Lactobacillaceae
Moraxellaceae
Mycoplasmataceae
Neisseriaceae
Pasteurellaceae
Peptostreptococcaceae
Porphyromonadaceae
Prevotellaceae
Rhodocyclaceae
Ruminococcaceae
Staphylococcaceae
Streptococcaceae
Veillonellaceae
xUnclassified/Other