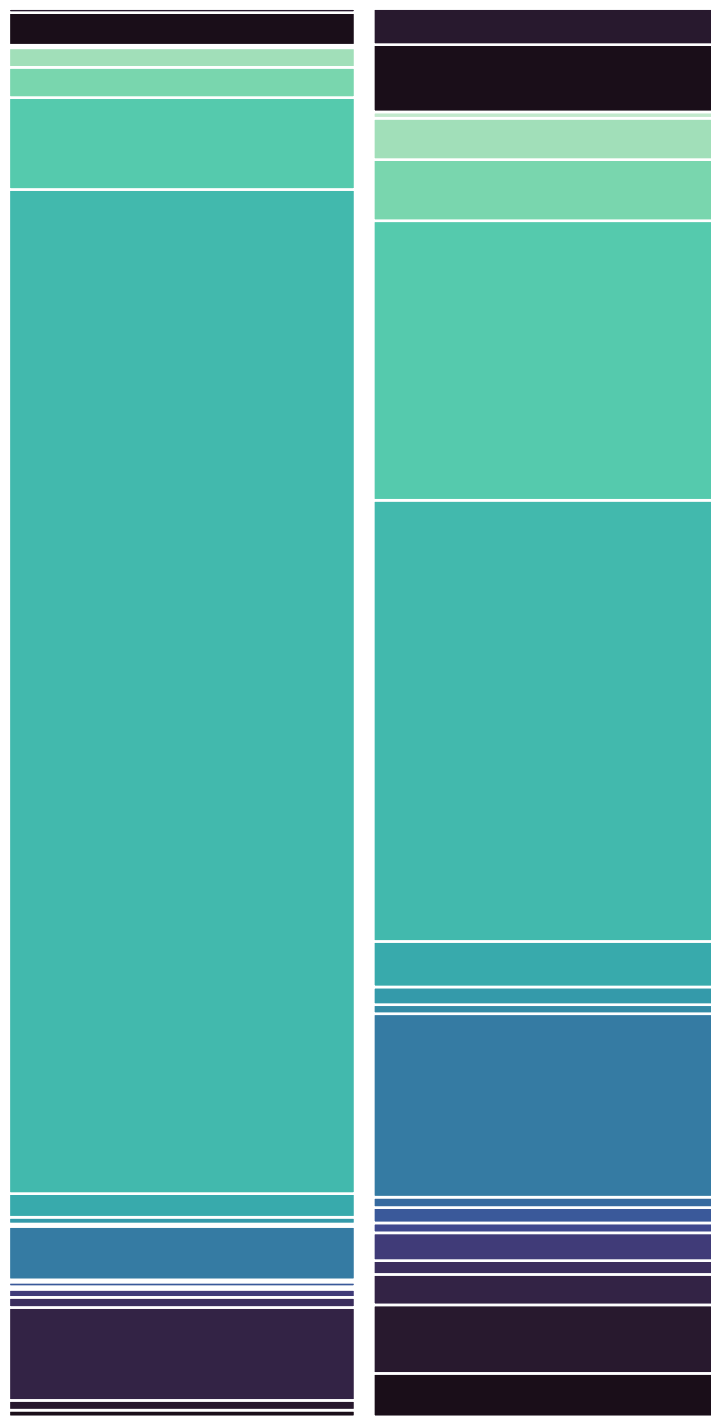


Relative Abundance (%)



CN

CMY

Group

- g_Alistipes
- g_Bacteroides
- g_Blautia
- g_Candidatus Saccharimonas
- g_Christensenellaceae R-7 group
- g_Desulfovibrio
- g_Helicobacter
- g_Incertae Sedis
- g_Ligilactobacillus
- g_Papillibacter
- g_Parabacteroides
- g_Parasutterella
- g_Prevotella_9
- g_Prevotellaceae NK3B31 group
- g_Rikenellaceae RC9 gut group
- g_Ruminococcus
- g_UCG-002
- g_[Eubacterium] oxidoreducens group
- g_gut metagenome