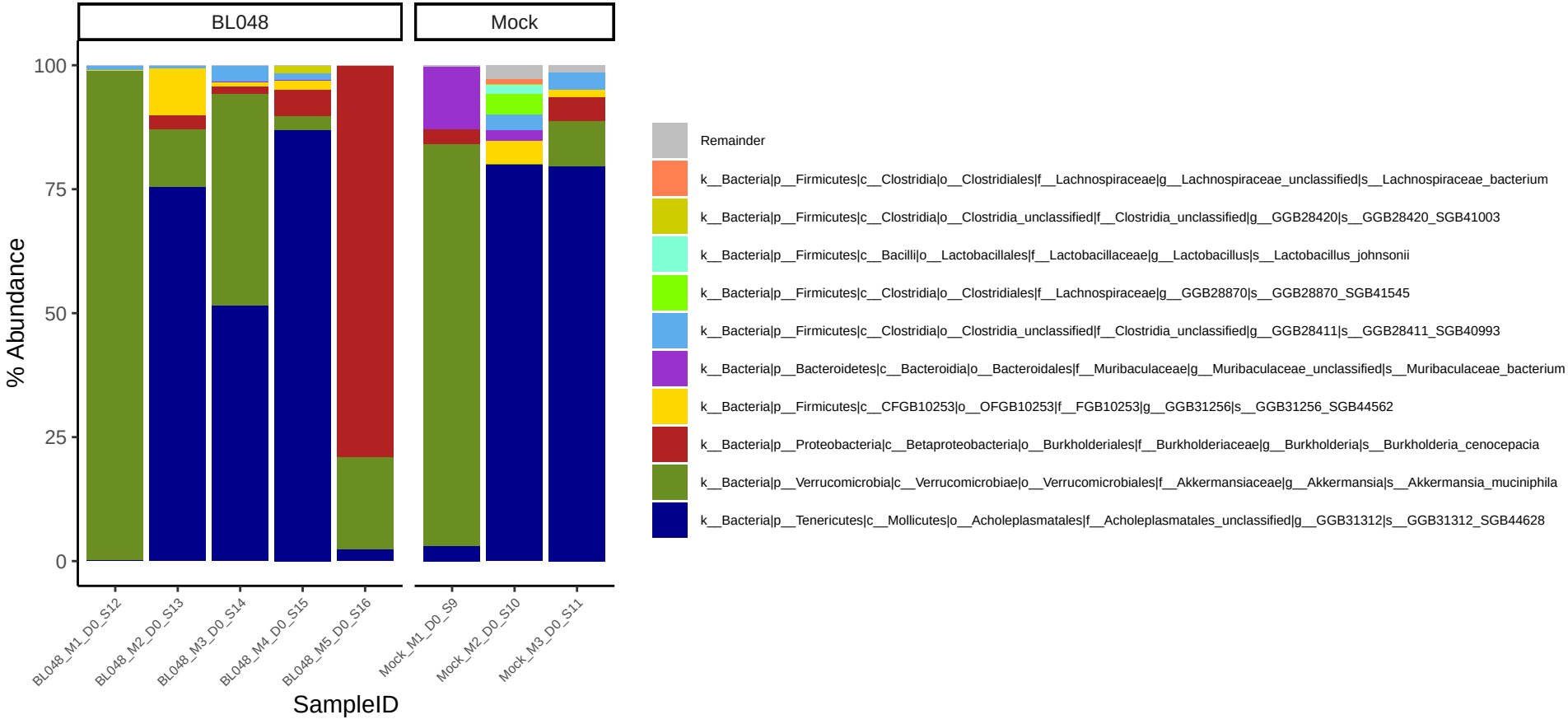
BL048
Mock
% Abundance
100
75
50
25
0
BL048_M1_D0_S12
BL048_M2_D0_S13
BL048_M3_D0_S14
BL048_M4_D0_S15
BL048_M5_D0_S16
Mock_M1_D0_S9
Mock_M2_D0_S10
Mock_M3_D0_S11
SampleID
Remainder
k__Bacteria|p__Firmicutes|c__Clostridia|o__Clostridiales|f__Lachnospiraceae|g__Lachnospiraceae_unclassified|s__Lachnospiraceae_bacterium
k__Bacteria|p__Firmicutes|c__Clostridia|o__Clostridia_unclassified|f__Clostridia_unclassified|g__GGB28420|s__GGB28420_SGB41003
k__Bacteria|p__Firmicutes|c__Bacilli|o__Lactobacillales|f__Lactobacillaceae|g__Lactobacillus|s__Lactobacillus_johnsonii
k__Bacteria|p__Firmicutes|c__Clostridia|o__Clostridiales|f__Lachnospiraceae|g__GGB28870|s__GGB28870_SGB41545
k__Bacteria|p__Firmicutes|c__Clostridia|o__Clostridia_unclassified|f__Clostridia_unclassified|g__GGB28411|s__GGB28411_SGB40993
k__Bacteria|p__Bacteroidetes|c__Bacteroidia|o__Bacteroidales|f__Muribaculaceae|g__Muribaculaceae_unclassified|s__Muribaculaceae_bacterium
k__Bacteria|p__Firmicutes|c__CFGB10253|o__OFGB10253|f__FGB10253|g__GGB31256|s__GGB31256_SGB44562
k__Bacteria|p__Proteobacteria|c__Betaproteobacteria|o__Burkholderiales|f__Burkholderiaceae|g__Burkholderia|s__Burkholderia_cenocepacia
k__Bacteria|p__Verrucomicrobia|c__Verrucomicrobiae|o__Verrucomicrobiales|f__Akkermansiaceae|g__Akkermansia|s__Akkermansia_muciniphila
k__Bacteria|p__Tenericutes|c__Mollicutes|o__Acholeplasmatales|f__Acholeplasmatales_unclassified|g__GGB31312|s__GGB31312_SGB44628