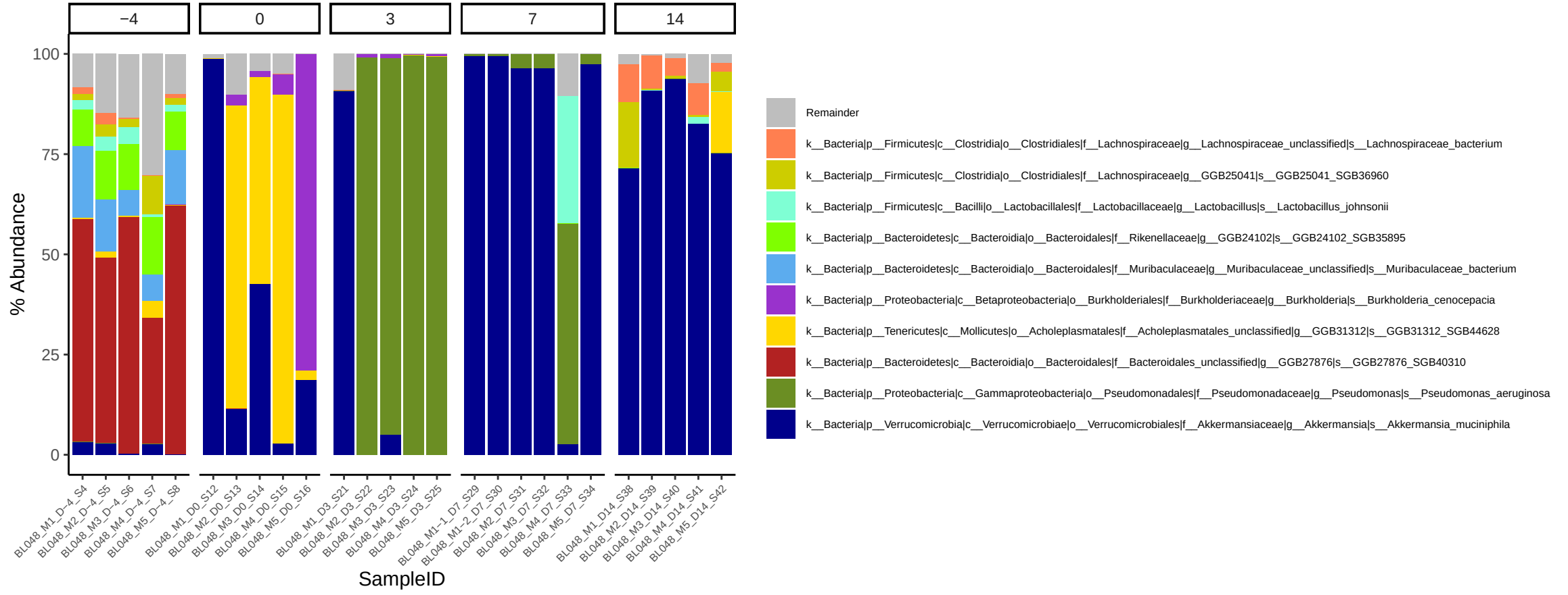
−4
0
3
7
14
100
75
50
25
0
% Abundance
BL048_M1_D−4_S4
BL048_M2_D−4_S5
BL048_M3_D−4_S6
BL048_M4_D−4_S7
BL048_M5_D−4_S8
BL048_M1_D0_S12
BL048_M2_D0_S13
BL048_M3_D0_S14
BL048_M4_D0_S15
BL048_M5_D0_S16
BL048_M1_D3_S21
BL048_M2_D3_S22
BL048_M3_D3_S23
BL048_M4_D3_S24
BL048_M5_D3_S25
BL048_M1−1_D7_S29
BL048_M1−2_D7_S30
BL048_M2_D7_S31
BL048_M3_D7_S32
BL048_M4_D7_S33
BL048_M5_D7_S34
BL048_M1_D14_S38
BL048_M2_D14_S39
BL048_M3_D14_S40
BL048_M4_D14_S41
BL048_M5_D14_S42
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__Clostridiales|f__Lachnospiraceae|g__GGB25041|s__GGB25041_SGB36960
k__Bacteria|p__Firmicutes|c__Bacilli|o__Lactobacillales|f__Lactobacillaceae|g__Lactobacillus|s__Lactobacillus_johnsonii
k__Bacteria|p__Bacteroidetes|c__Bacteroidia|o__Bacteroidales|f__Rikenellaceae|g__GGB24102|s__GGB24102_SGB35895
k__Bacteria|p__Bacteroidetes|c__Bacteroidia|o__Bacteroidales|f__Muribaculaceae|g__Muribaculaceae_unclassified|s__Muribaculaceae_bacterium
k__Bacteria|p__Proteobacteria|c__Betaproteobacteria|o__Burkholderiales|f__Burkholderiaceae|g__Burkholderia|s__Burkholderia_cenocepacia
k__Bacteria|p__Tenericutes|c__Mollicutes|o__Acholeplasmatales|f__Acholeplasmatales_unclassified|g__GGB31312|s__GGB31312_SGB44628
k__Bacteria|p__Bacteroidetes|c__Bacteroidia|o__Bacteroidales|f__Bacteroidales_unclassified|g__GGB27876|s__GGB27876_SGB40310
k__Bacteria|p__Proteobacteria|c__Gammaproteobacteria|o__Pseudomonadales|f__Pseudomonadaceae|g__Pseudomonas|s__Pseudomonas_aeruginosa
k__Bacteria|p__Verrucomicrobia|c__Verrucomicrobiae|o__Verrucomicrobiales|f__Akkermansiaceae|g__Akkermansia|s__Akkermansia_muciniphila