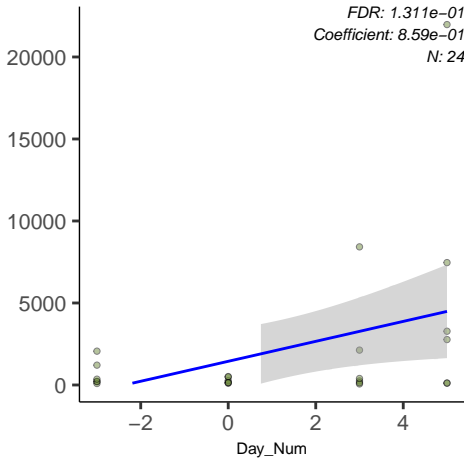
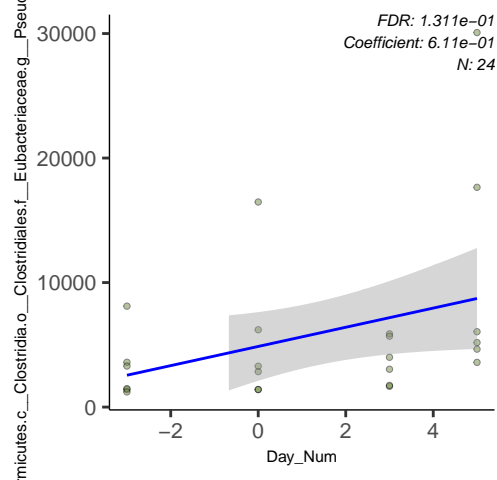
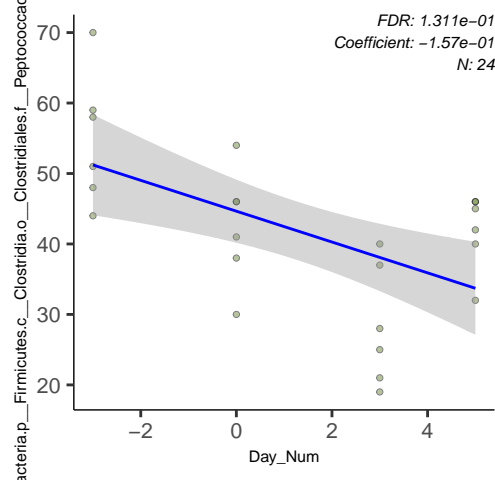


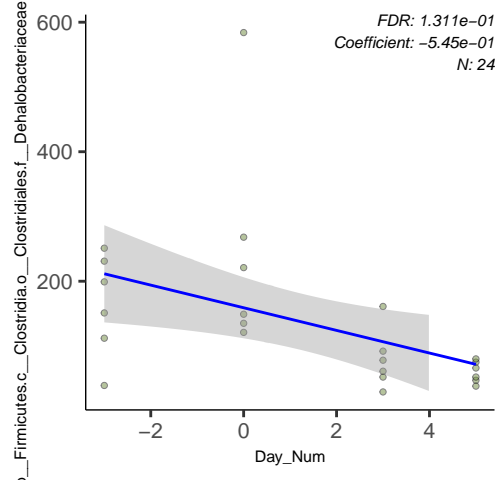
bacteria.p__Firmicutes.c__Clostridia.o__Clostridiales.f__Clostridiaceae

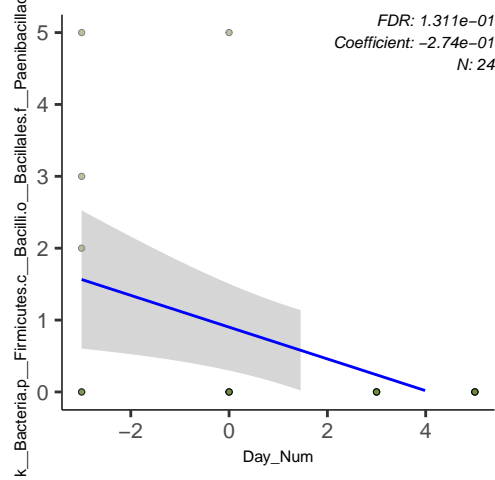
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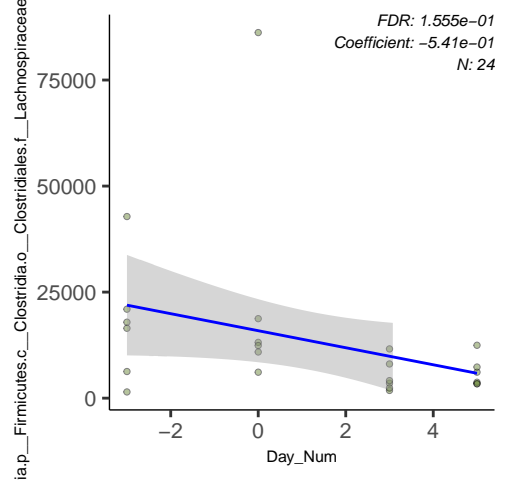


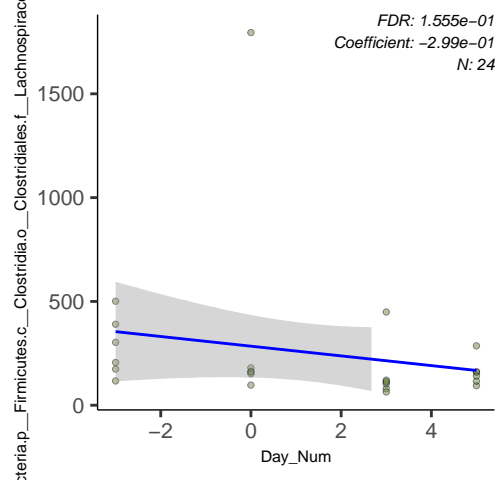


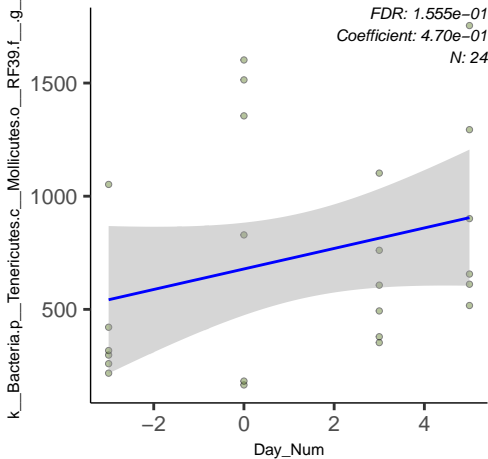


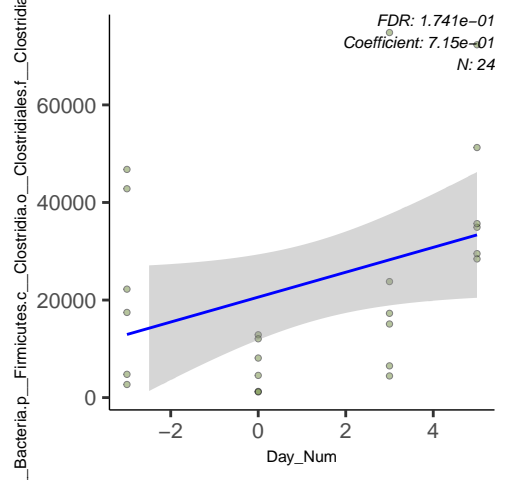


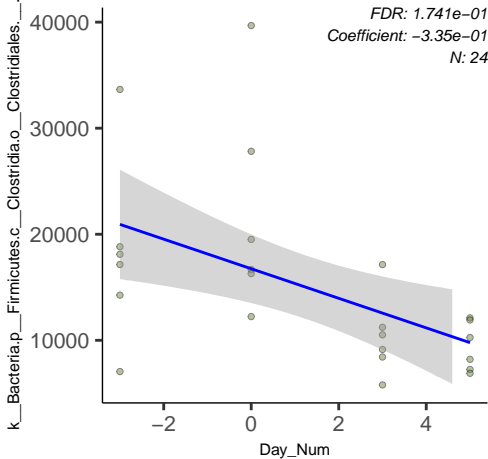


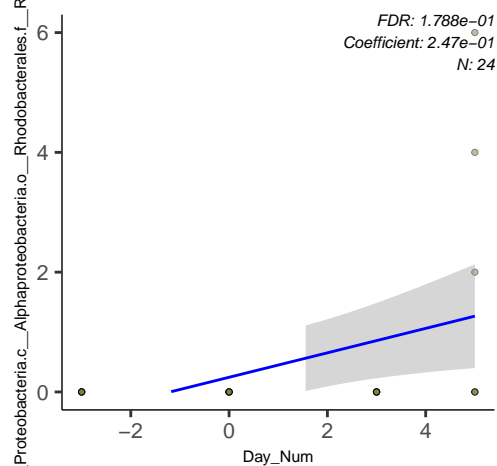






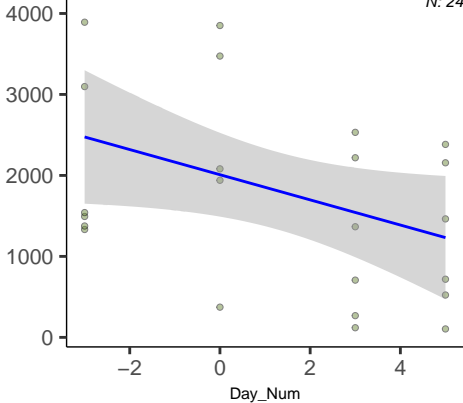


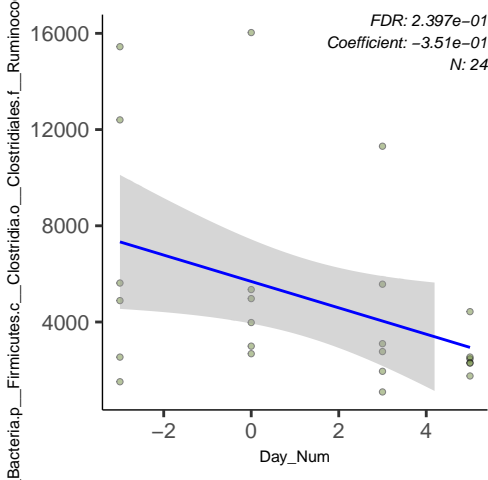


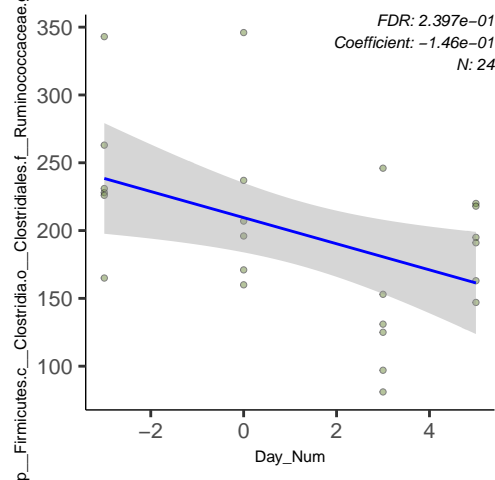


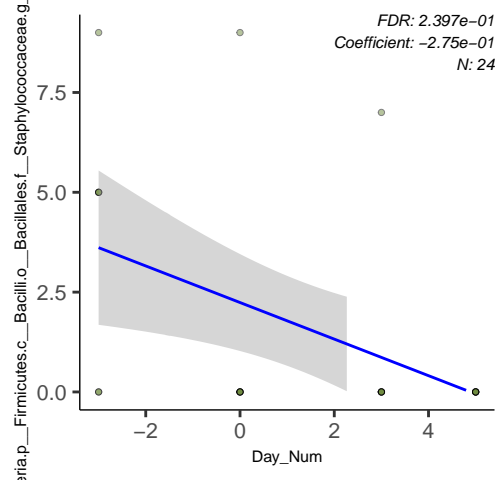
ria.p__Firmicutes.c__Clostridia.o__Clostridiales.f__Lachnospiraceae

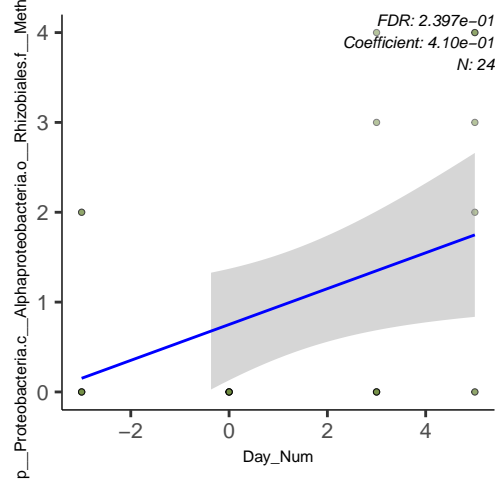
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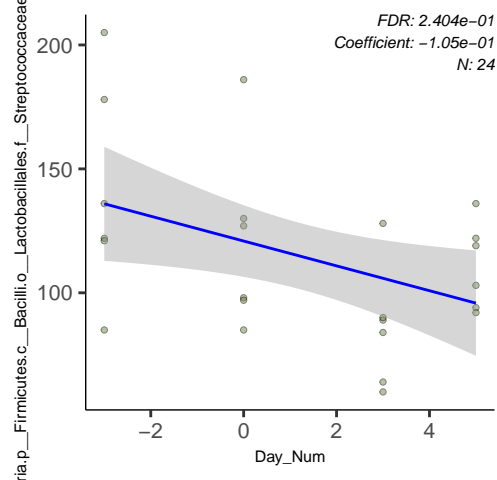


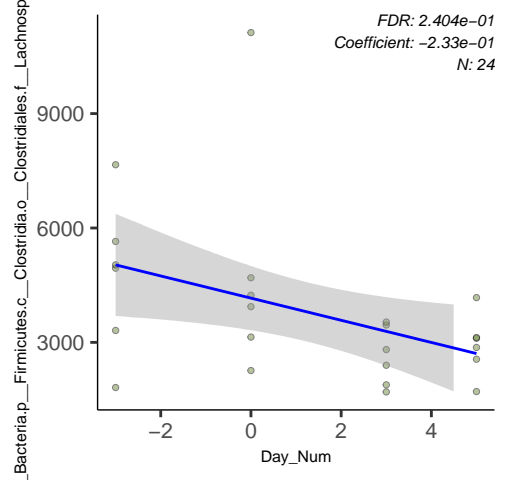












ia.p__Firmicutes.c__Clostridia.o__Clostridiales.f__Ruminococcaceae

FDR: 2.404e-01
Coefficient: -4.03e-01
N: 24

