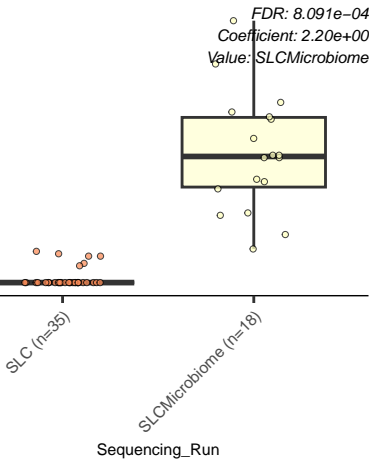
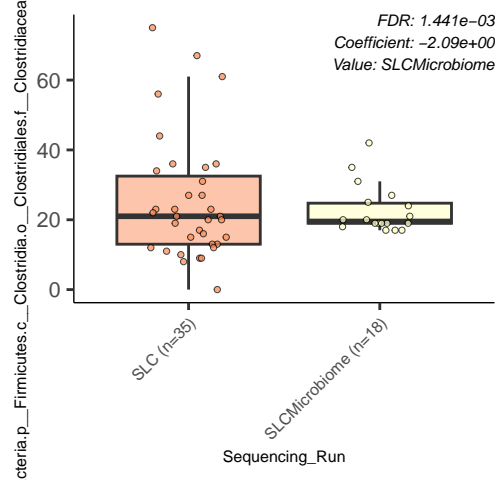
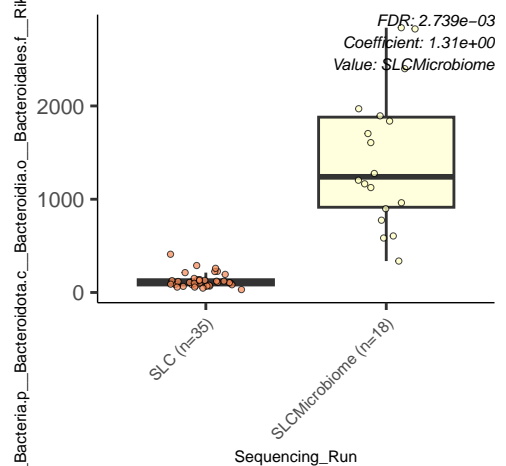


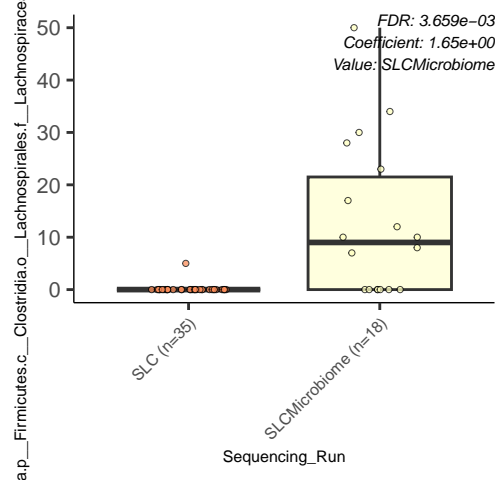
icutes.c\_\_Clostridia.o\_\_Peptostreptococcales.Tissierellales.f\_\_Anaer

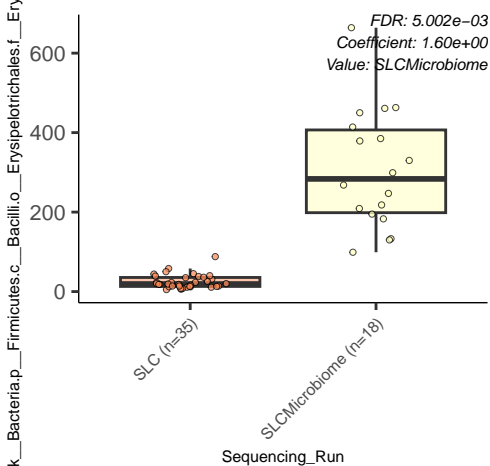


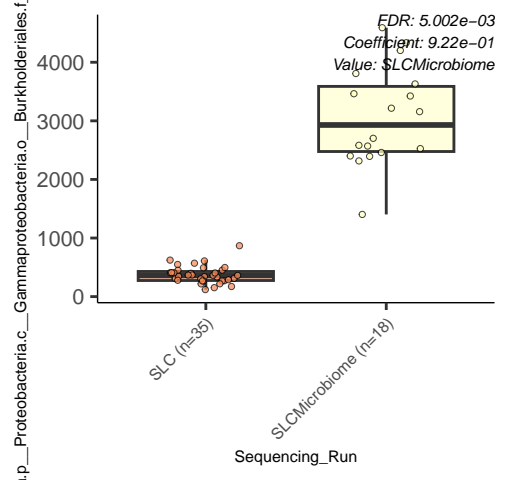


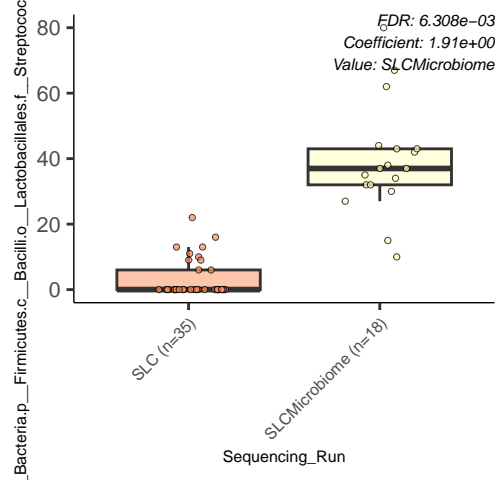












Bacteria.p\_\_Firmicutes.c\_\_Bacilli.o\_\_Erysipelotrichales.f\_\_Erysipel

20

10

0

SLC (n=35)

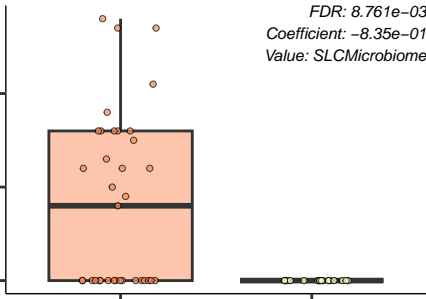
SLCMicrobiome (n=18)

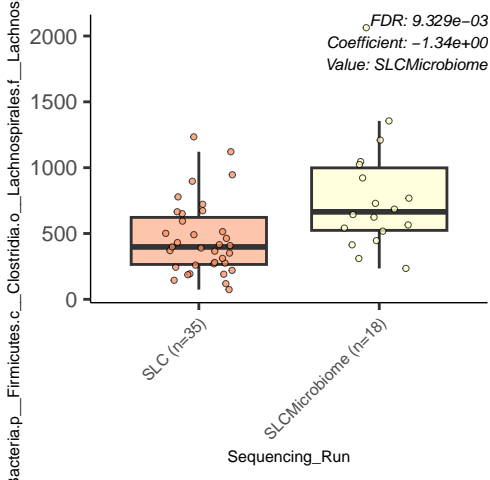
Sequencing\_Run

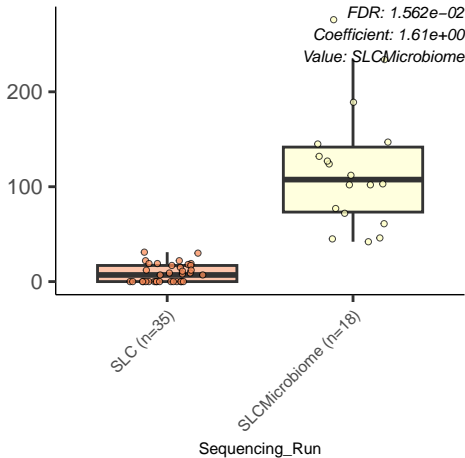
FDR: 8.761e-03

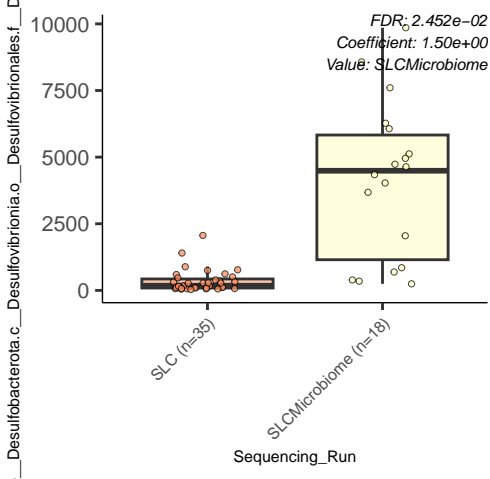
Coefficient: -8.35e-01

Value: SLCMicrobiome

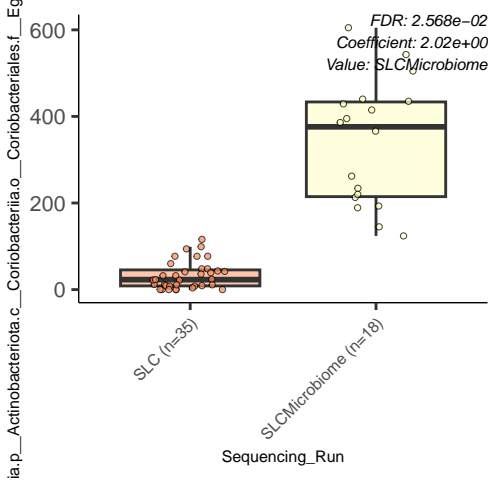


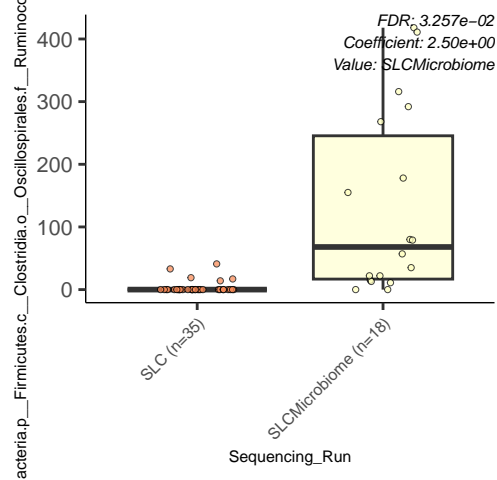


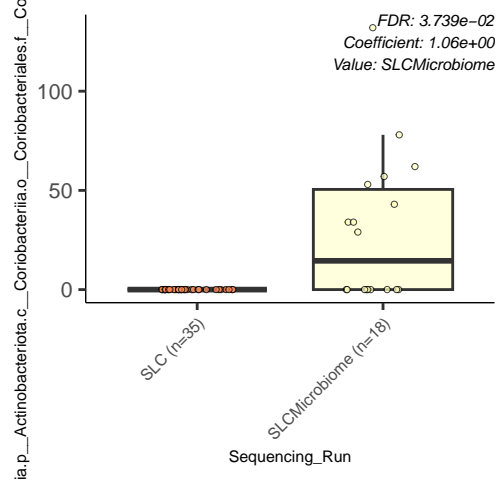


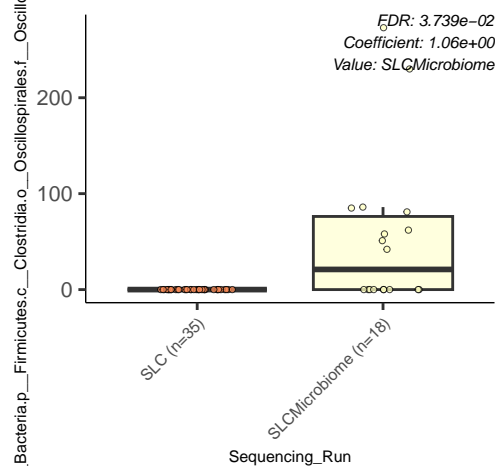


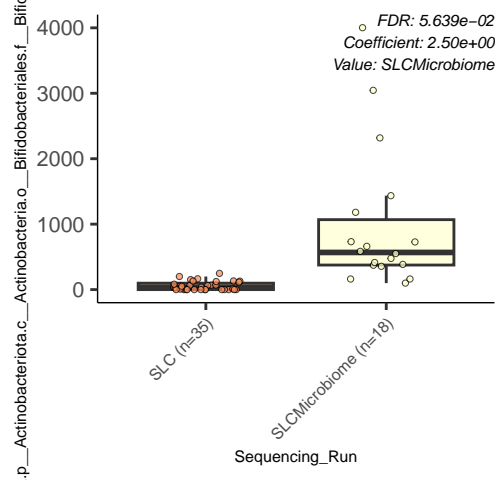


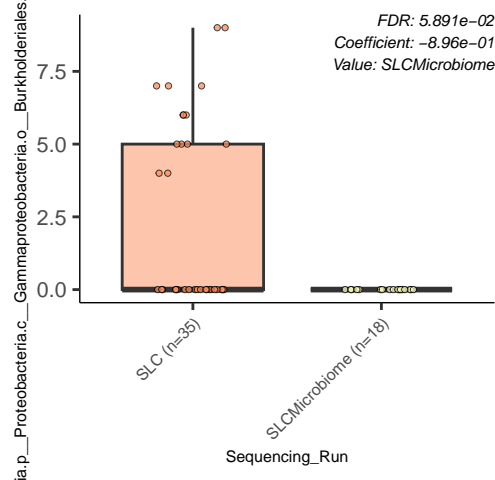


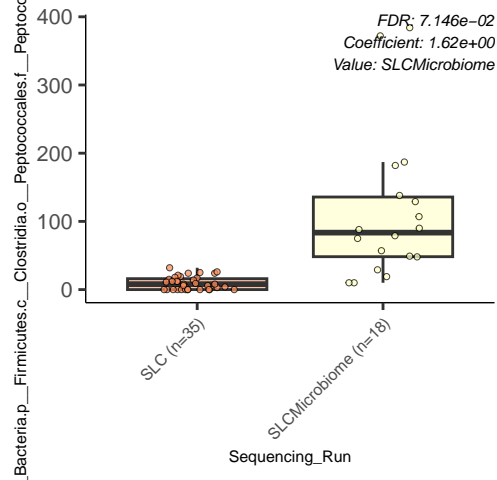




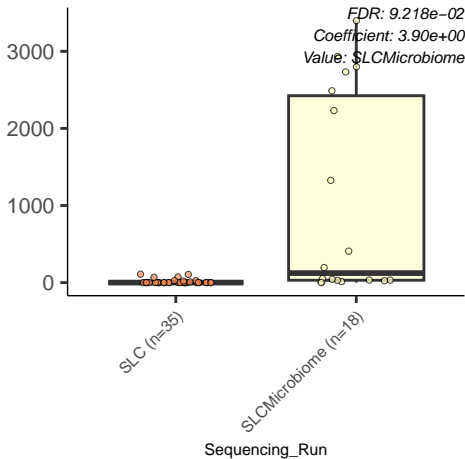






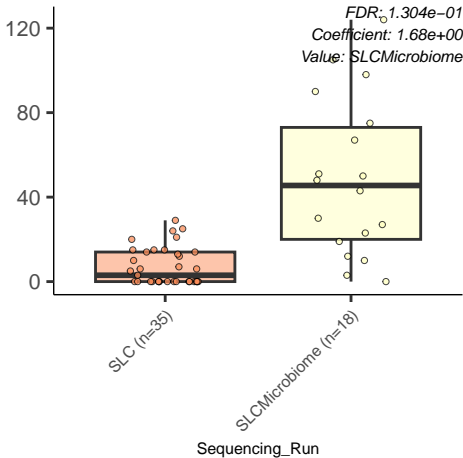


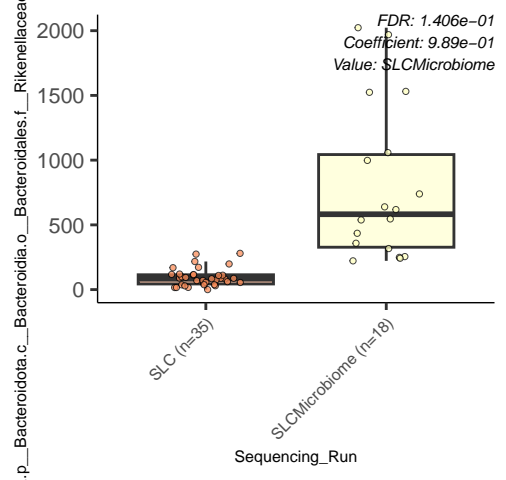
o\_Firmicutes.c\_Clostridia.o\_Lachnospirales.f\_Lachnospiraceae





cutes.c\_\_Clostridia.o\_\_Peptostreptococcales.Tissierellales.f\_\_Anaer





Firmicutes.c\_\_Clostridia.o\_\_Christensenellales.f\_\_Christensenellac

