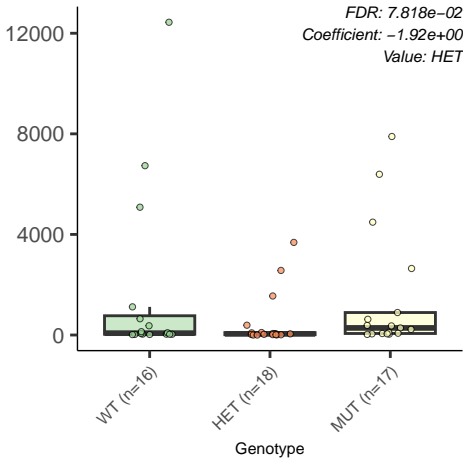
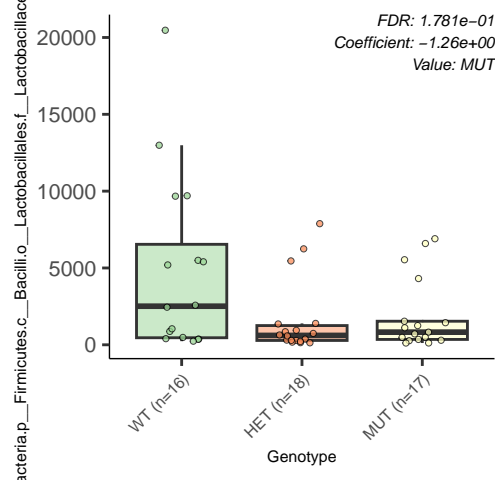


Verrucomicrobiota.c\_\_Verrucomicrobiae.o\_\_Verrucomicrobiales.f\_\_Ak





ria.p\_\_Firmicutes.c\_\_Bacilli.o\_\_Staphylococcales.f\_\_Staphylococca

FDR: 1.781e-01  
Coefficient: 1.42e+00  
Value: MUT

2000  
1500  
1000  
500  
0

WT (n=16)

HET (n=18)

MUT (n=17)

Genotype

