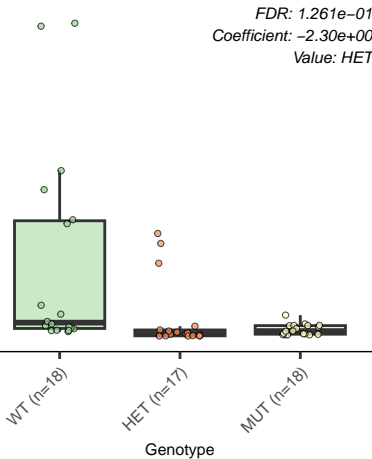


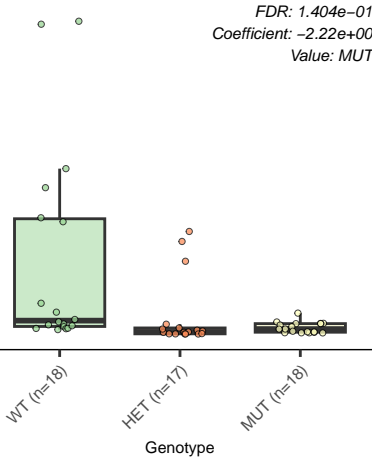
bacteria.p__Firmicutes.c__Bacilli.o__Lactobacillales.f__Lactobacillaceae

FDR: 1.261e-01
Coefficient: -2.30e+00
Value: HET

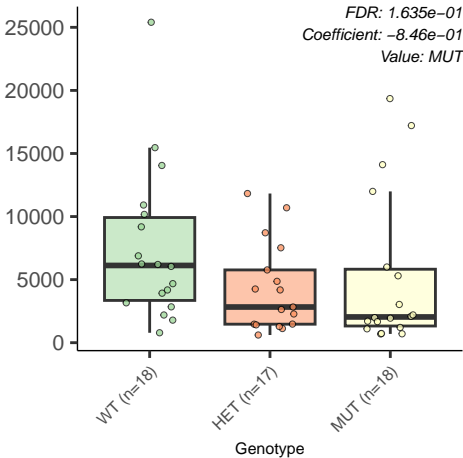


bacteria.p__Firmicutes.c__Bacilli.o__Lactobacillales.f__Lactobacillaceae

FDR: 1.404×10^{-1}
Coefficient: -2.22×10^0
Value: MUT

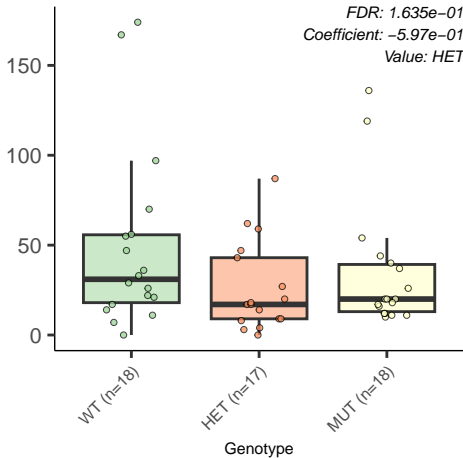


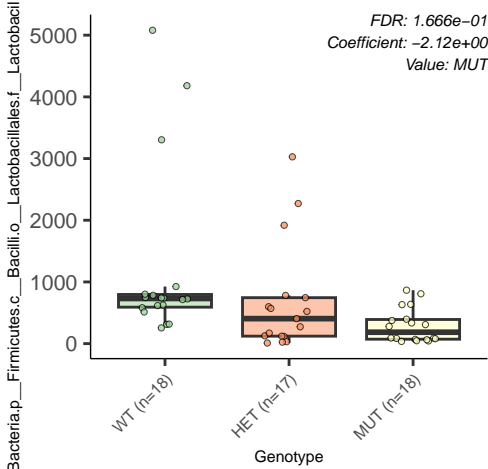
Bacteria.p__Firmicutes.c__Clostridia.o__Lachnospirales.f__Lachnospiraceae.g__Lachnospira

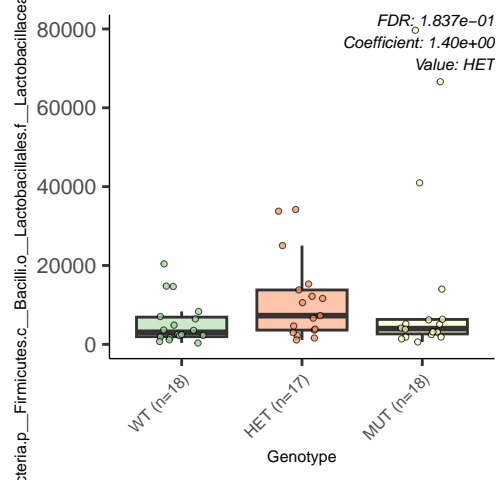


eria.p__Firmicutes.c__Clostridia.o__Oscillospirales.f__Oscillospirac

FDR: 1.635e-01
Coefficient: -5.97e-01
Value: HET

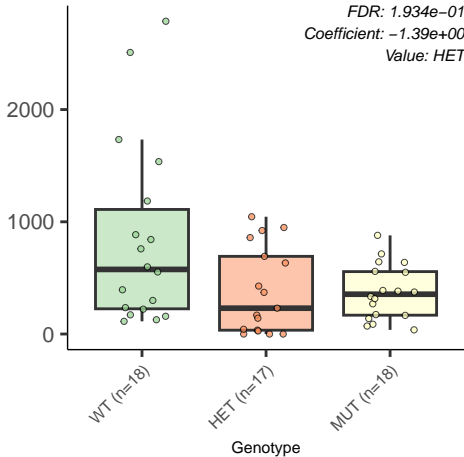


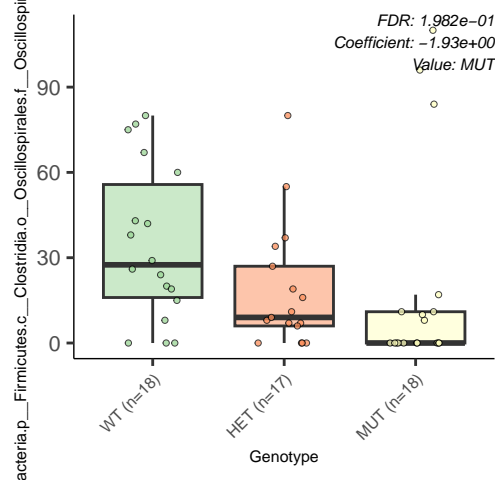


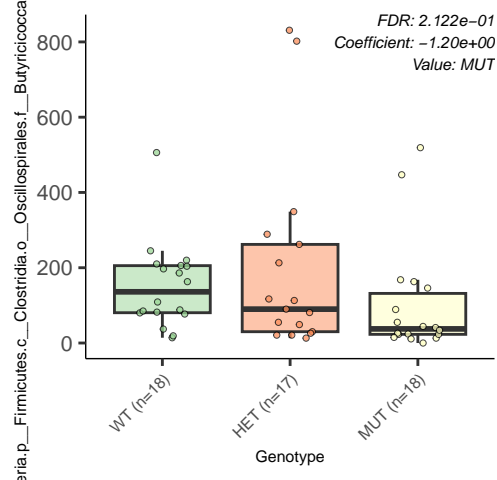


inobacteriota.c__Coriobacteriia.o__Coriobacteriales.f__Atopobiaceae

FDR: $1.934e-01$
Coefficient: $-1.39e+00$
Value: HET







terial.p__Firmicutes.c__Clostridia.o__Lachnospirales.f__Lachnospira

