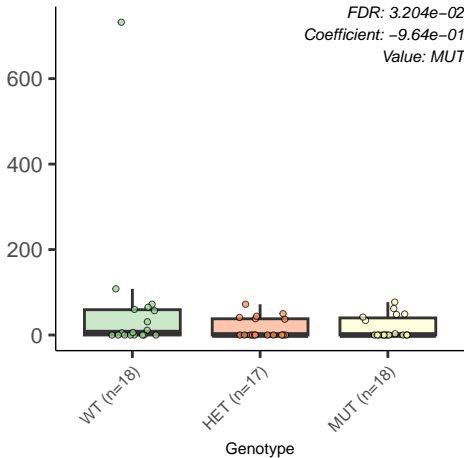


obacteria.c__Gammaproteobacteria.o__Enterobacterales.f__Enterobacteriaceae

FDR: 3.204e-02
Coefficient: -9.64e-01
Value: MUT



obacteria.c__Gammaproteobacteria.o__Enterobacterales.f__Enter

FDR: 3.427e-02
Coefficient: -9.55e-01
Value: HET

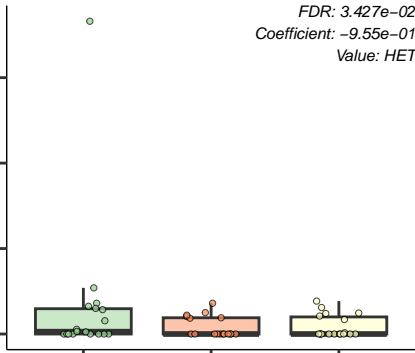
600
400
200
0

WT (n=18)

HET (n=17)

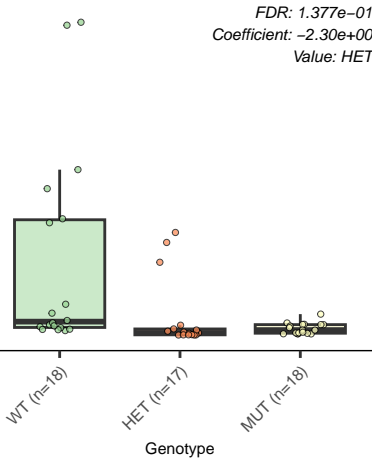
MUT (n=18)

Genotype

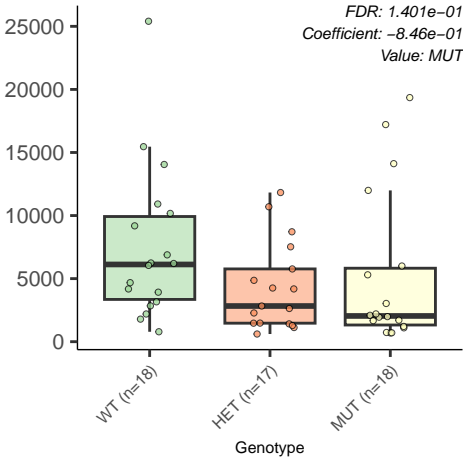


bacteria.p__Firmicutes.c__Bacilli.o__Lactobacillales.f__Lactobacillaceae

FDR: 1.377e-01
Coefficient: -2.30e+00
Value: HET



Bacteria.p__Firmicutes.c__Clostridia.o__Lachnospirales.f__Lachnospiraceae.g__Lachnospira



bacteria.p__Firmicutes.c__Bacilli.o__Lactobacillales.f__Lactobacillaceae

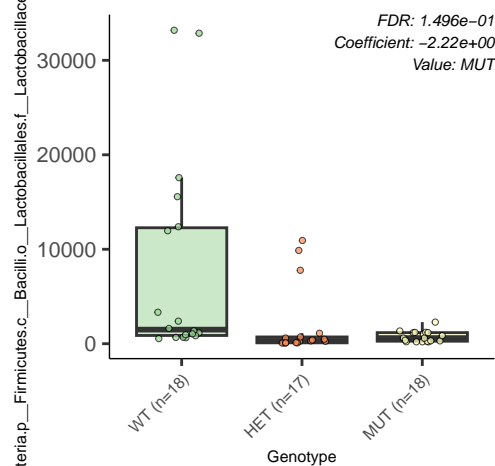
FDR: $1.496e-01$
Coefficient: $-2.22e+00$
Value: MUT

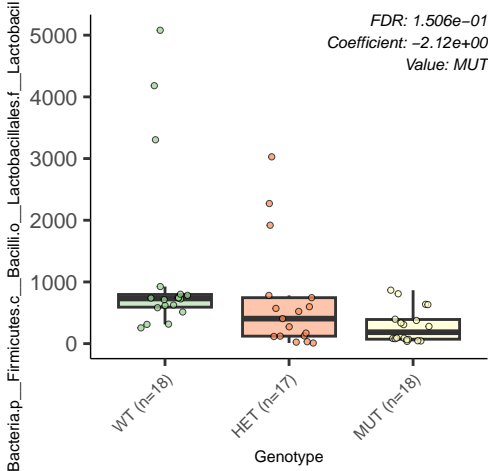
WT (n=18)

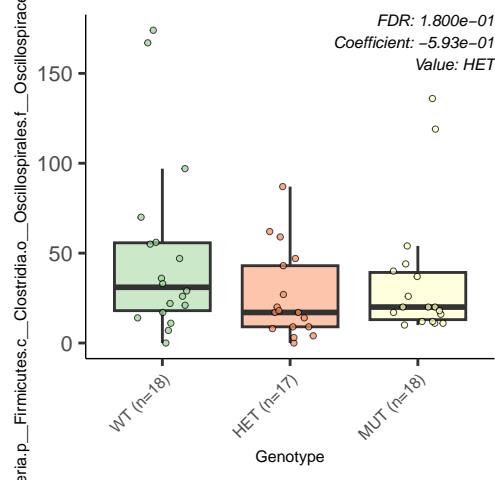
HET (n=17)

MUT (n=18)

Genotype







ria.p__Firmicutes.c__Clostridia.o__Lachnospirales.f__Lachnospira

20

10

0

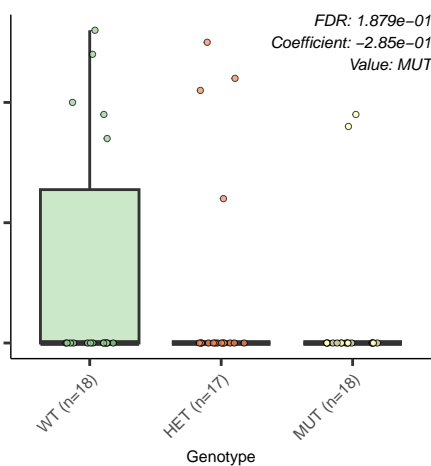
WT (n=18)

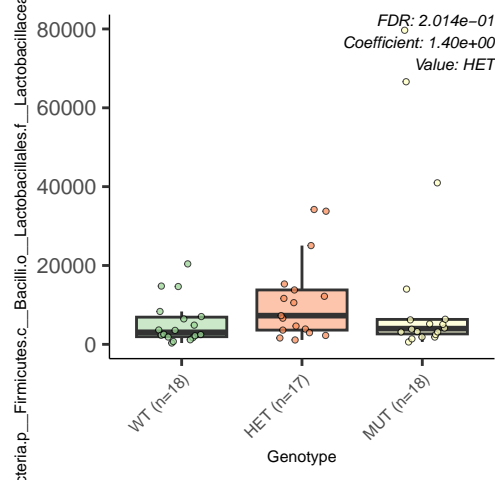
HET (n=17)

MUT (n=18)

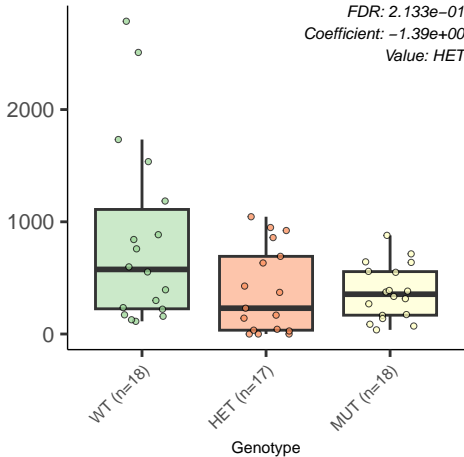
Genotype

FDR: 1.879e-01
Coefficient: -2.85e-01
Value: MUT





FDR: $2.133\text{e-}01$
Coefficient: $-1.39\text{e}+00$
Value: HET



bacteria.p__Firmicutes.c__Clostridia.o__Oscillospirales.f__Oscillospira

