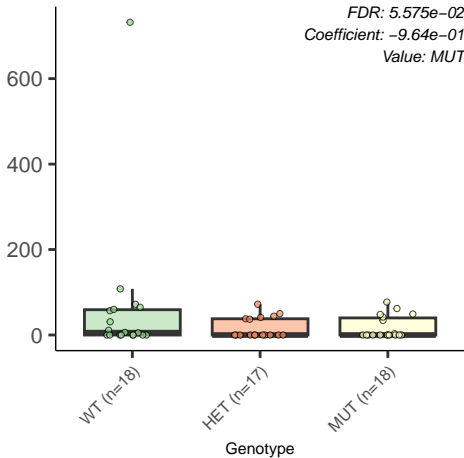
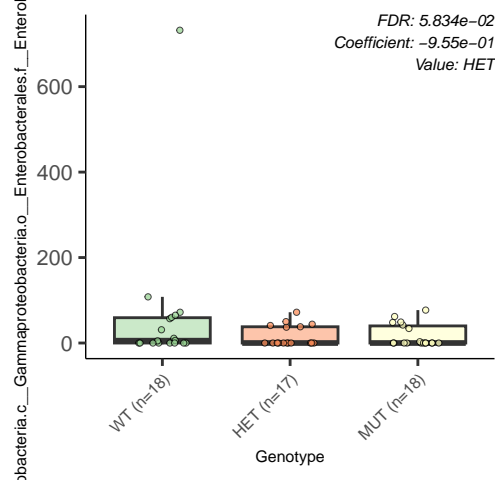


obacteria.c__Gammaproteobacteria.o__Enterobacterales.f__Enter

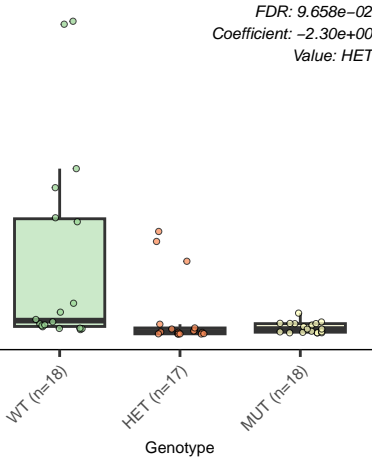
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Coefficient: -9.64e-01
Value: MUT





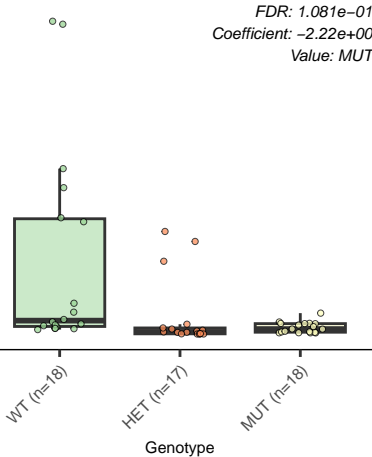
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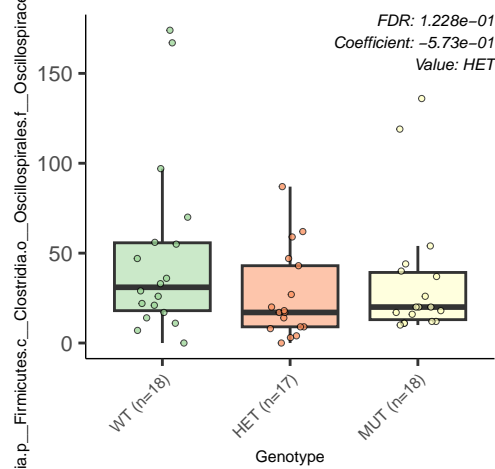
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Coefficient: -2.30×10^0
Value: HET

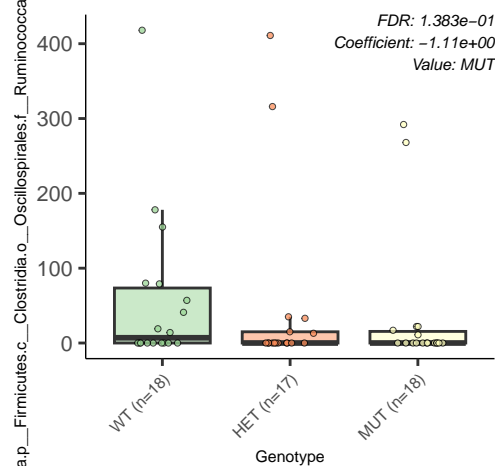


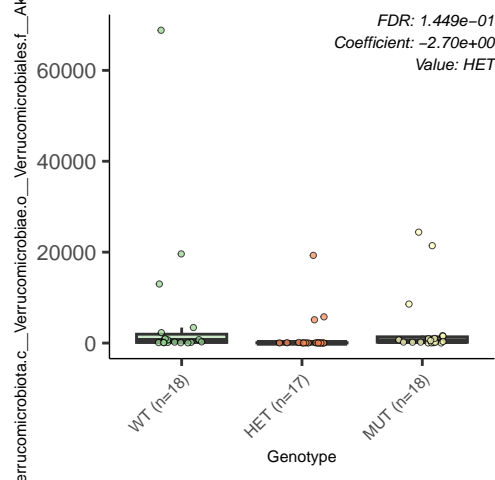
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FDR: 1.081e-01
Coefficient: -2.22e+00
Value: MUT



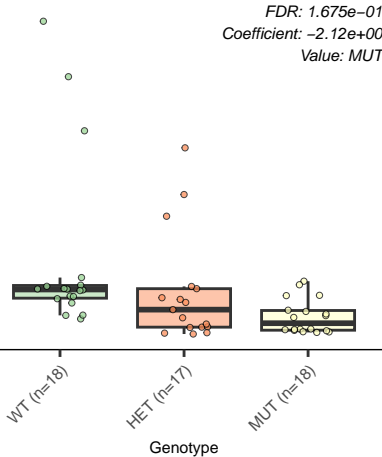




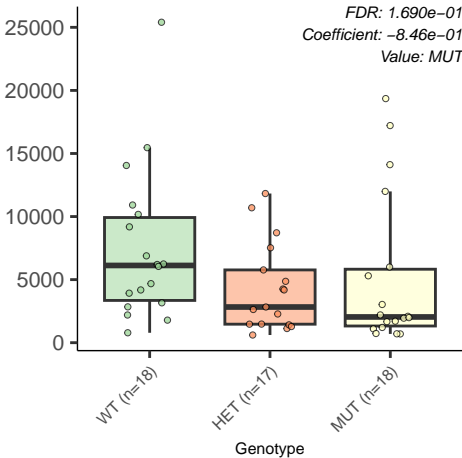


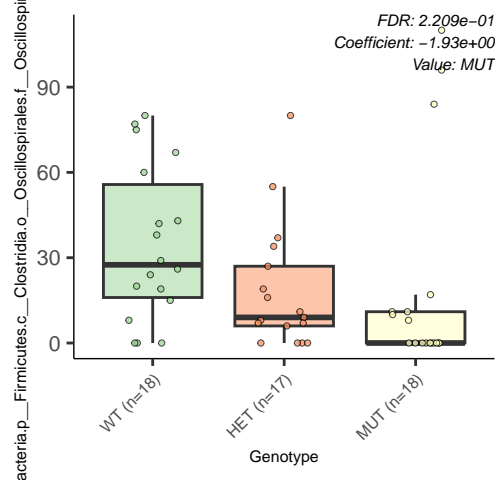
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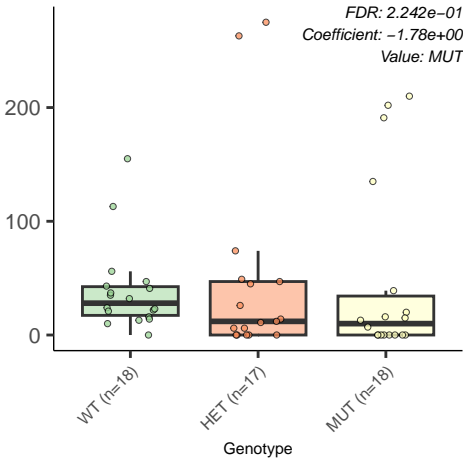
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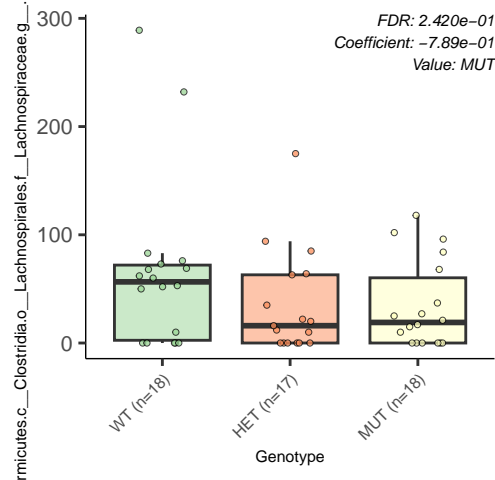


Bacteria.p_Firmicutes.c_Clostridia.o_Lachnospirales.f_Lach



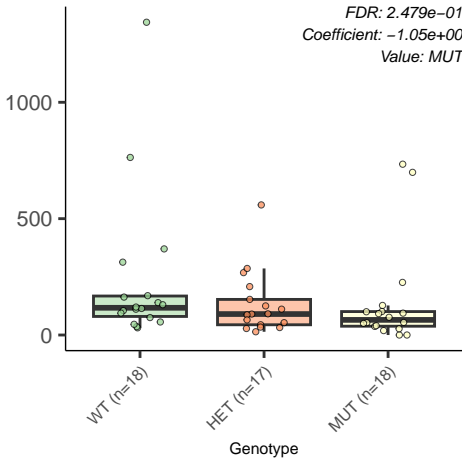






Bacteria.p__Firmicutes.c__Clostridia.o__Lachnospirales.f__Lachnospiraceae

FDR: 2.479e-01
Coefficient: -1.05e+00
Value: MUT



Firmicutes.c__Clostridia.o__Lachnospirales.f__Lachnospiraceae.g

