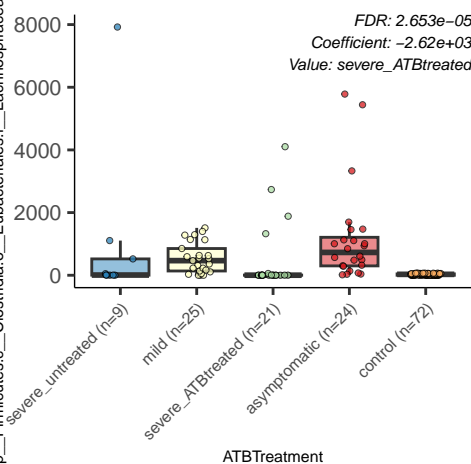
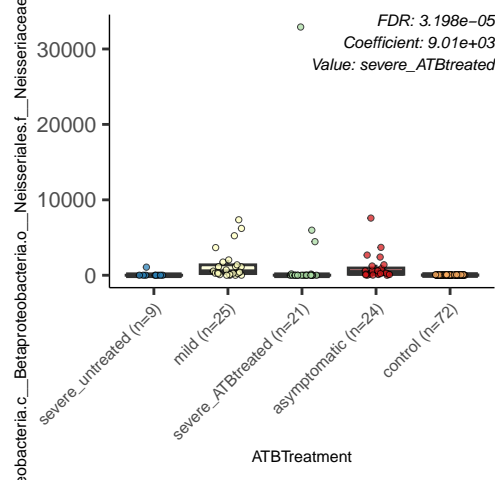
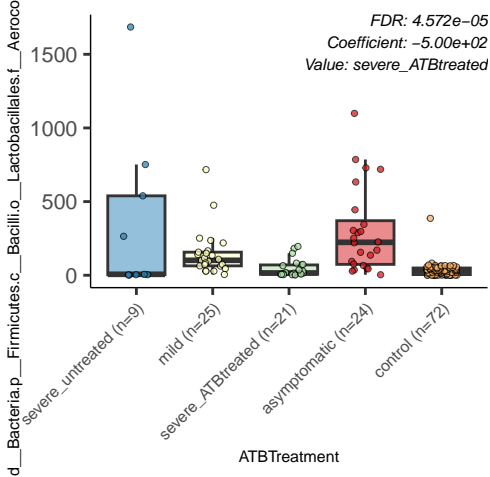
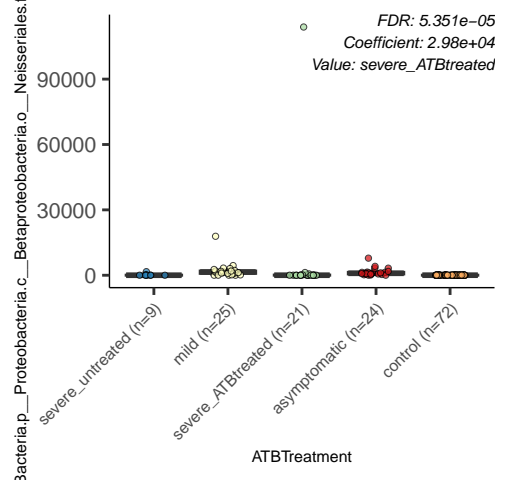


FDR: $2.653e-05$
Coefficient: $-2.62e+03$
Value: severe_ATBtreated









eria.p_Actinobacteria.c_Actinomycetia.o_Actinomycetales.f_Ac

$FDR: 5.414e-05$
 $Coefficient: -3.20e+03$
 $Value: severe_ATBtreated$

10000

5000

0

severe_untreated (n=9)

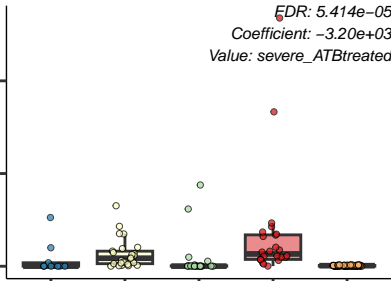
mild (n=25)

severe_ATBtreated (n=21)

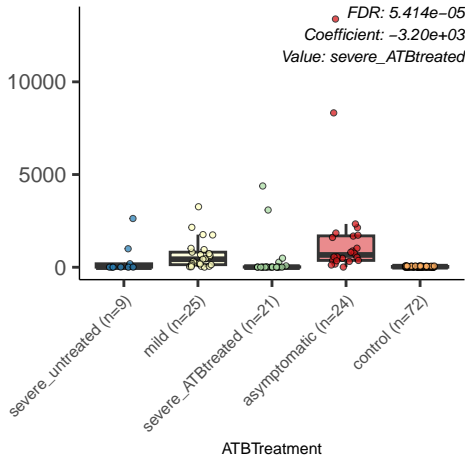
asymptomatic (n=24)

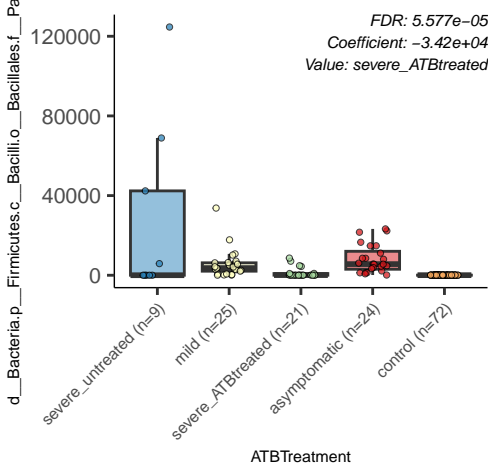
control (n=72)

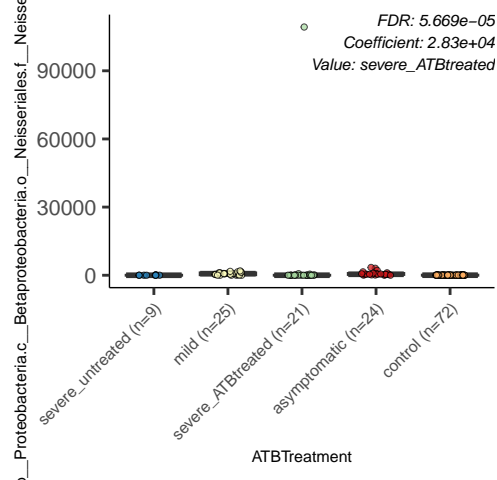
ATBTreatment

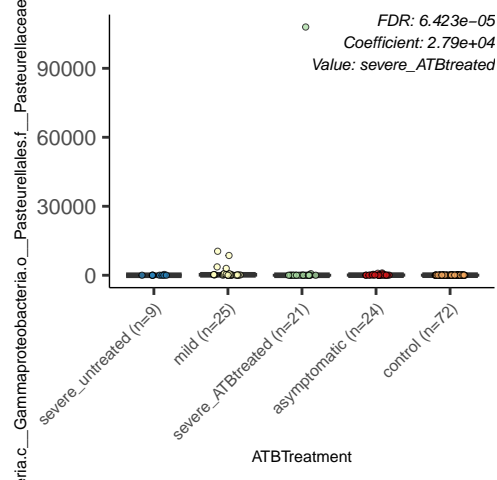


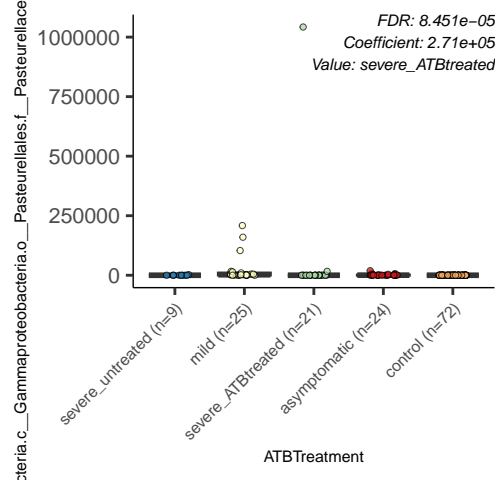
bacteria.c_Actinomycetia.o_Actinomycetales.f_Actinomycetaceae

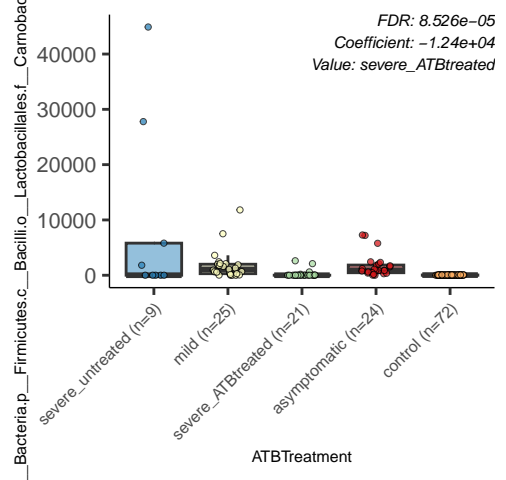


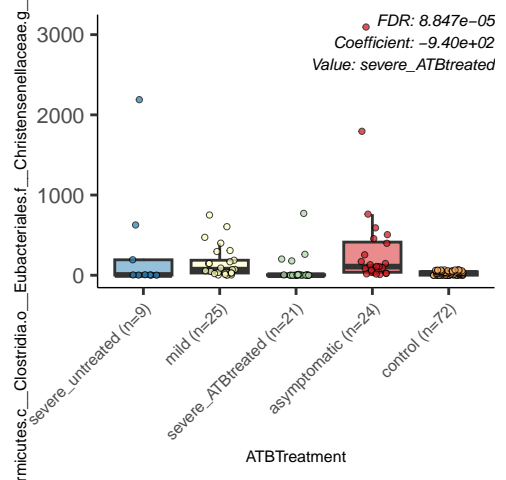


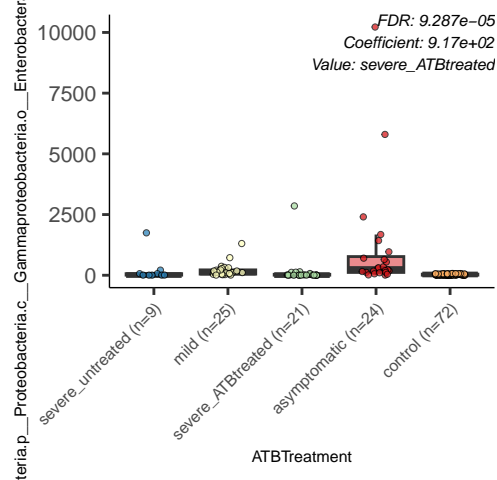


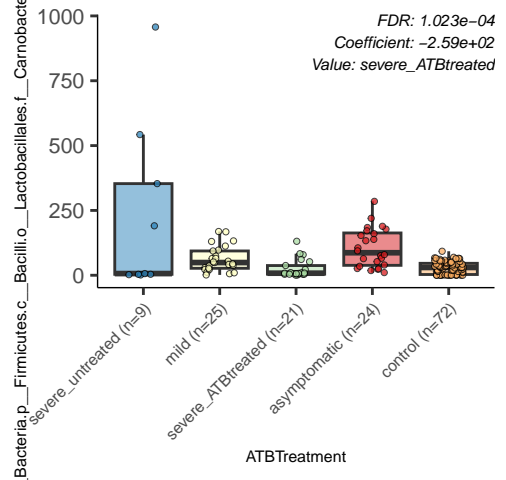


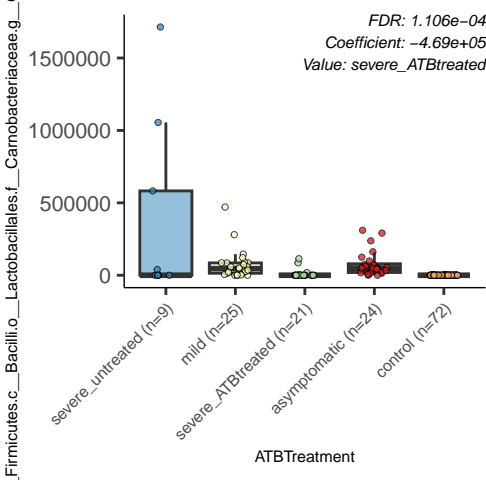


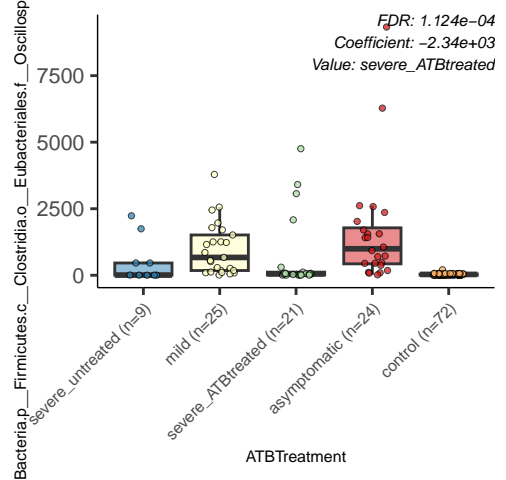




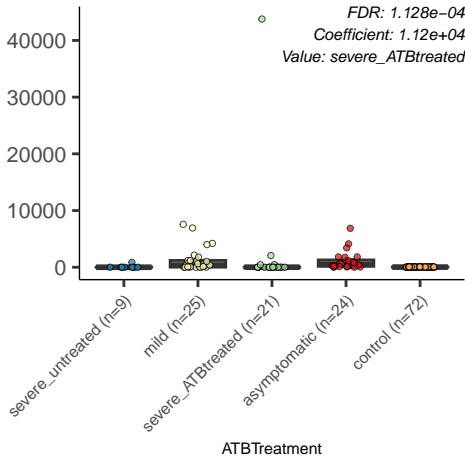






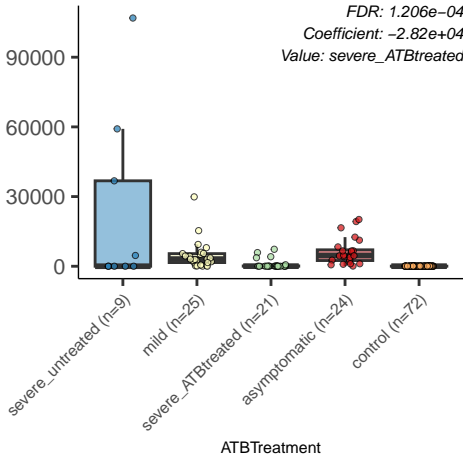


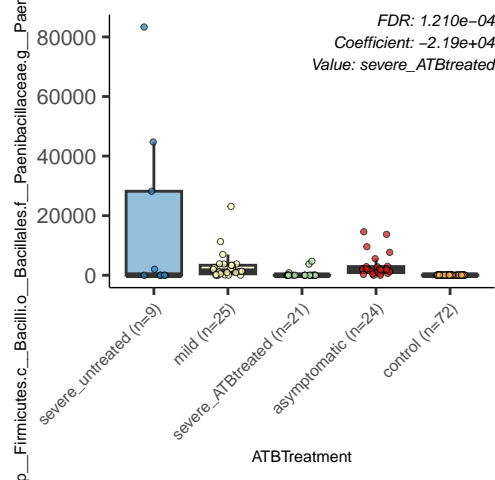
a.c__Gammaproteobacteria.o__Pasteurellales.f__Pasteurellaceae.g



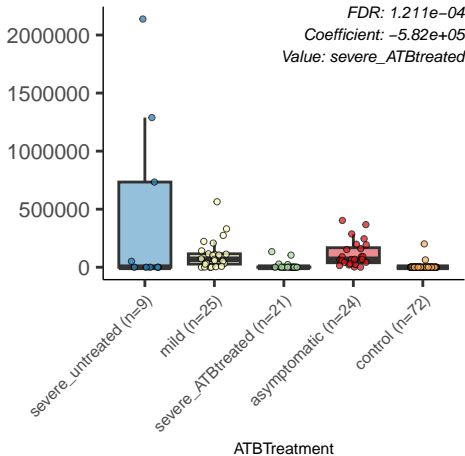
d_Bacteria.p_Firmicutes.c_Bacilli.o_Bacillales.f_Paenibacillales

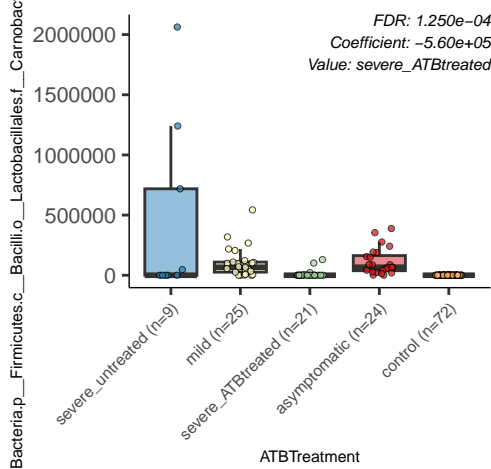
FDR: 1.206e-04
Coefficient: -2.82e+04
Value: severe_ATBtreated

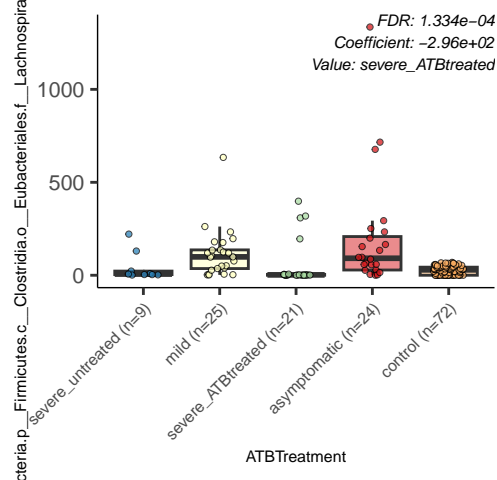


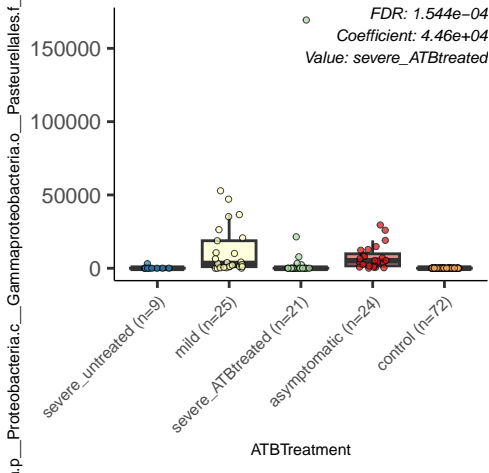


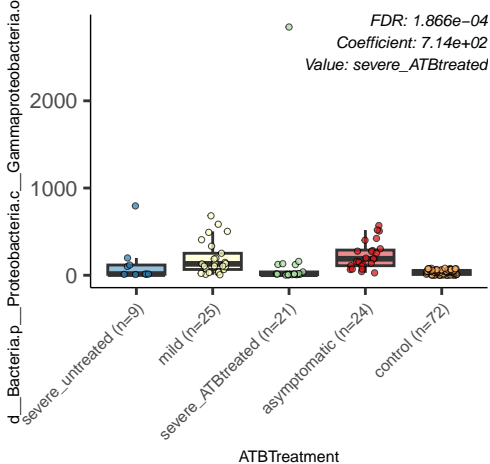
d_Bacteria.p_Firmicutes.c_Bacilli.o_Lactobacillales.f





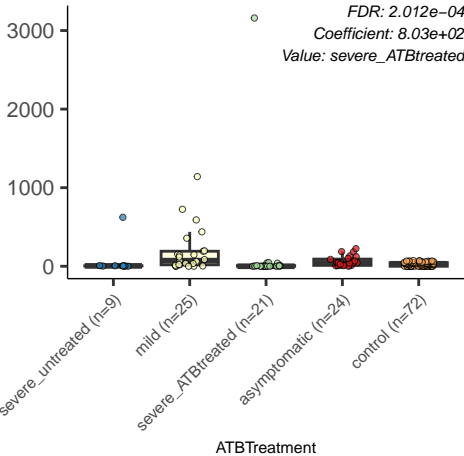


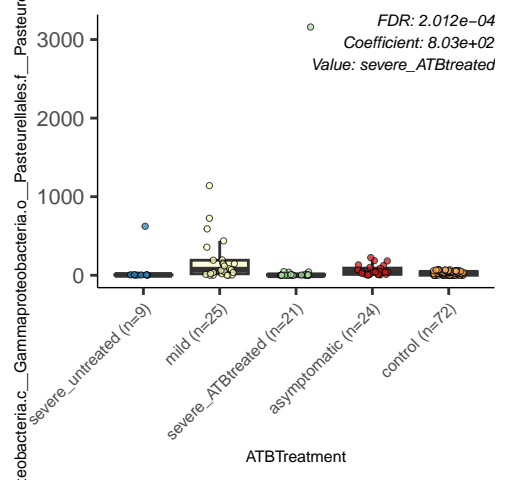


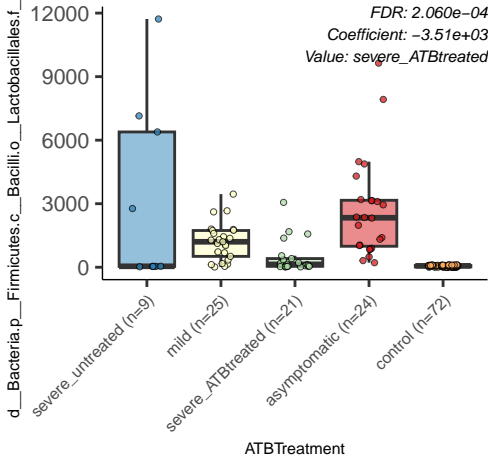


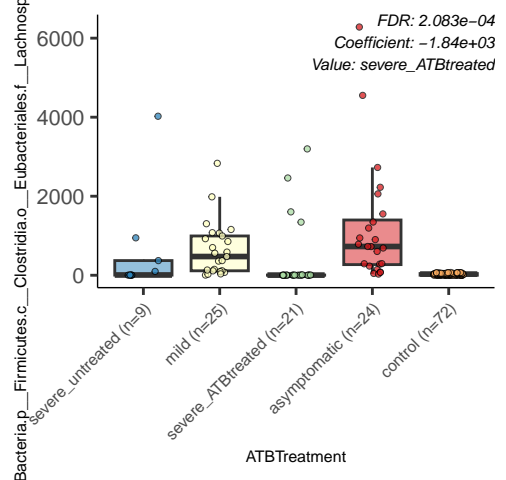
ria.p__Proteobacteria.c__Gammaproteobacteria.o__Pasteurellales.

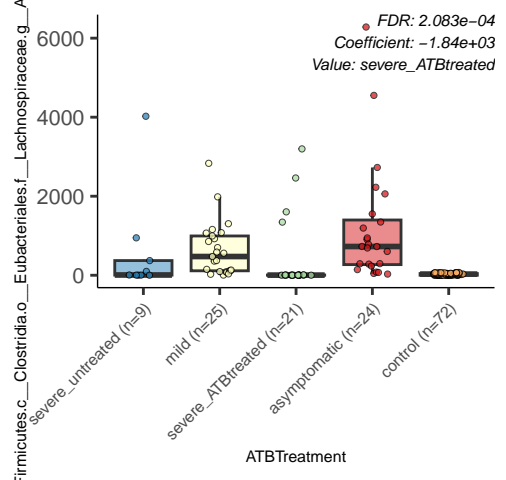
FDR: 2.012e-04
Coefficient: 8.03e+02
Value: severe_ATBtreated

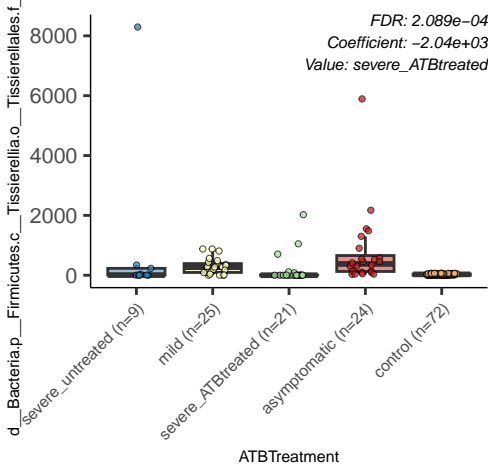


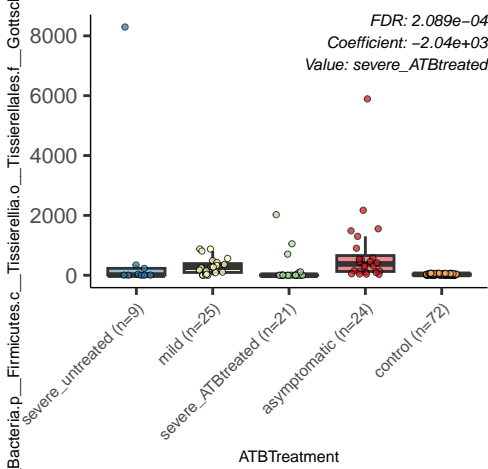


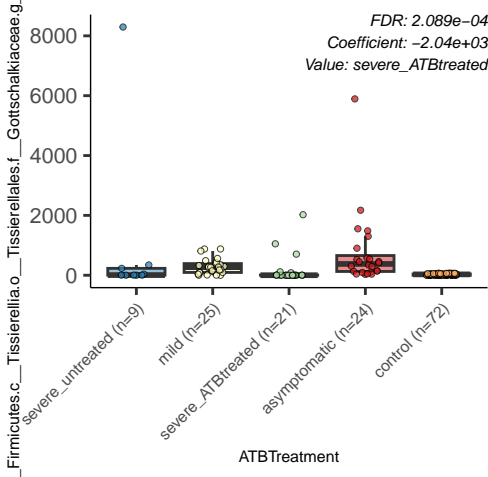


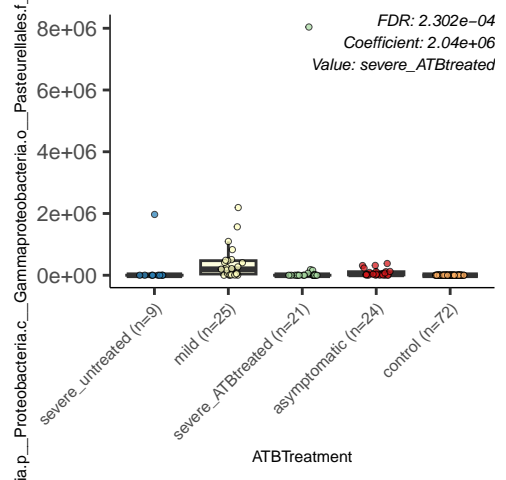


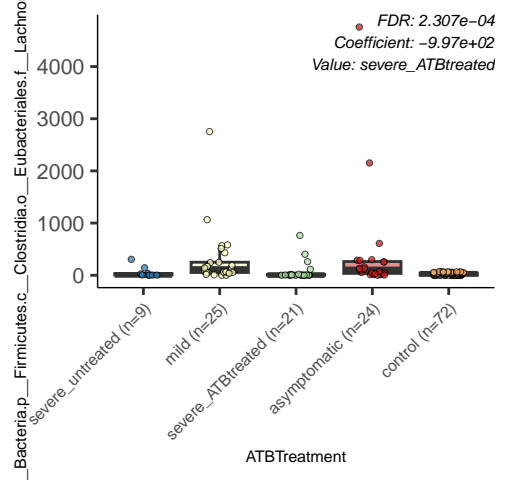


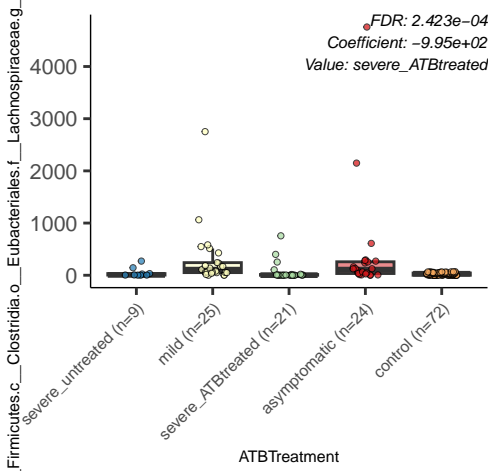




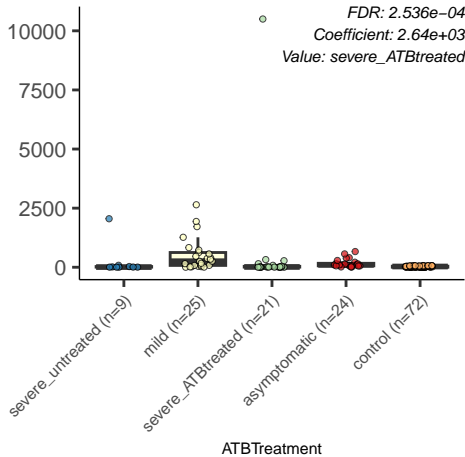


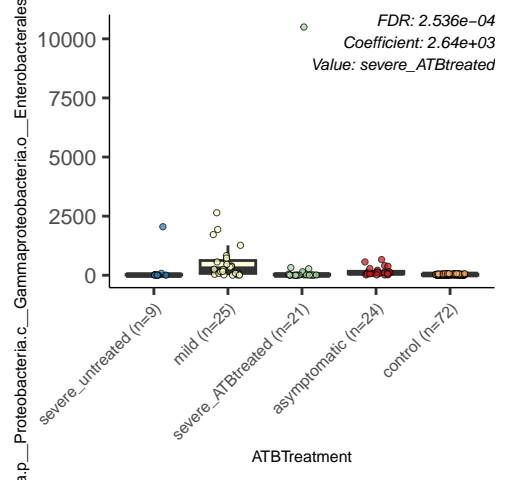


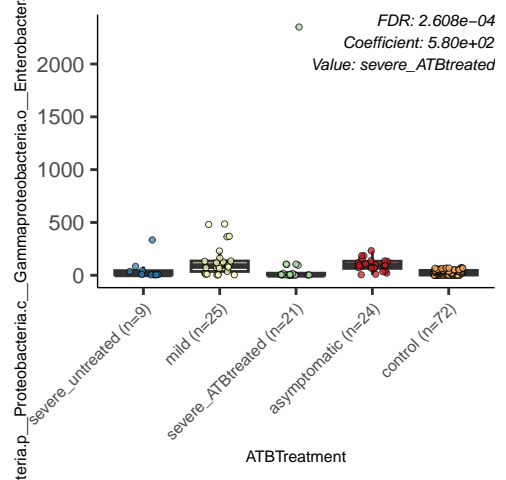


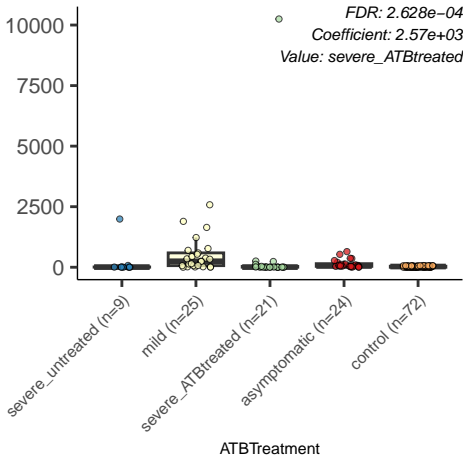


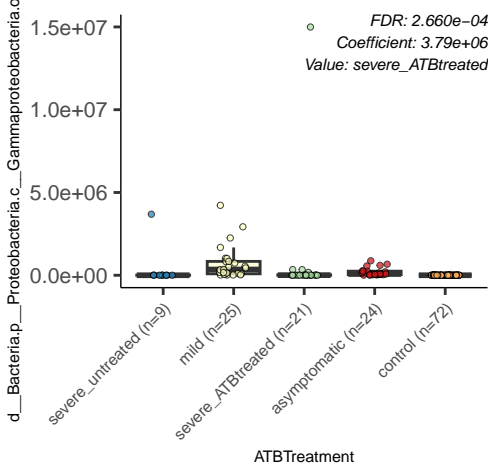
bacteria.p__Proteobacteria.c__Gammaproteobacteria.o__Enterobact

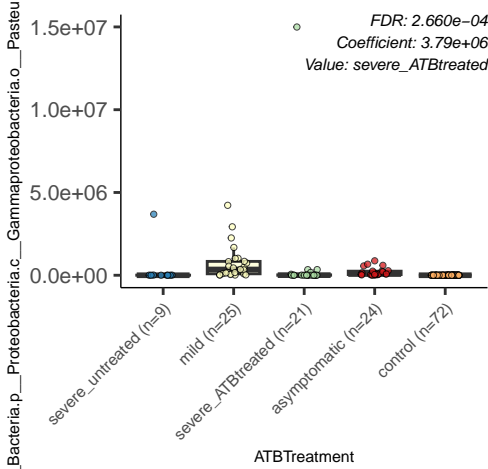


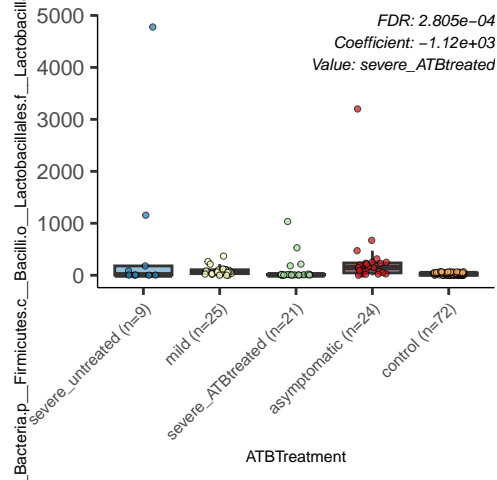


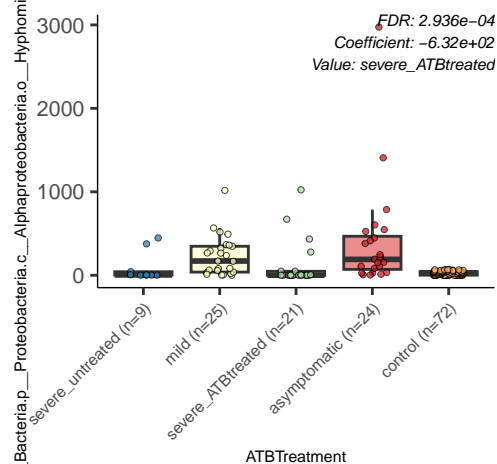


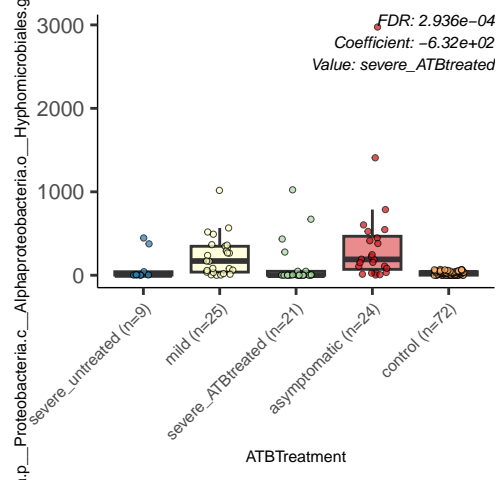


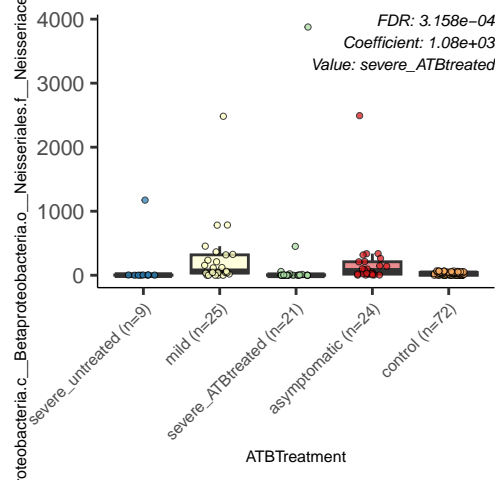


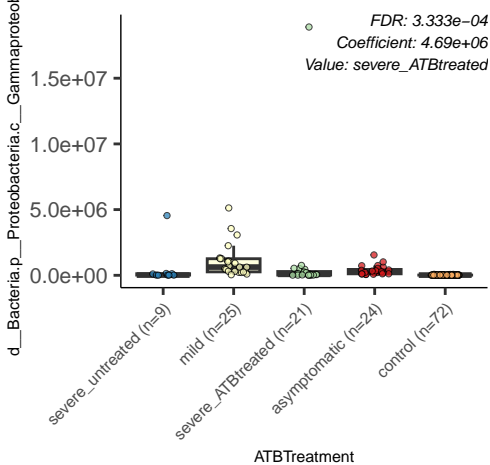


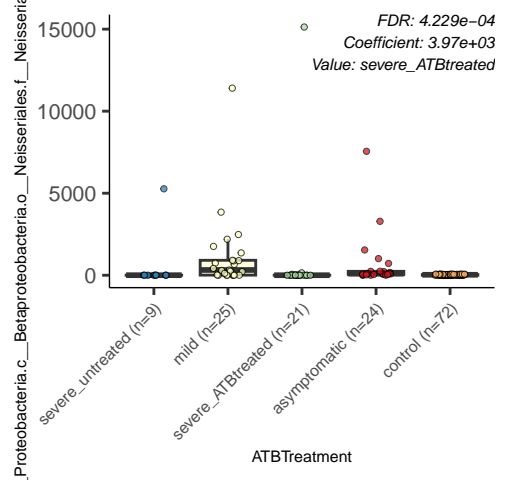


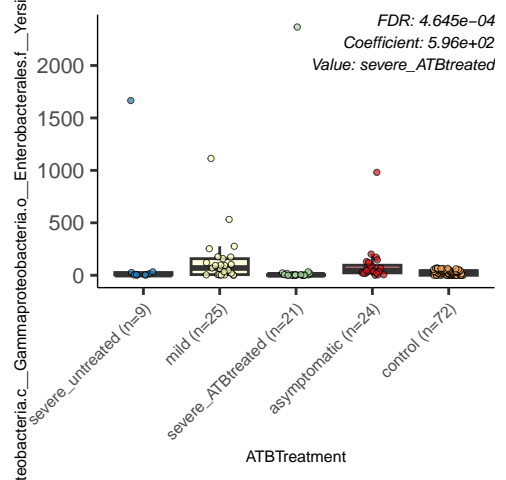


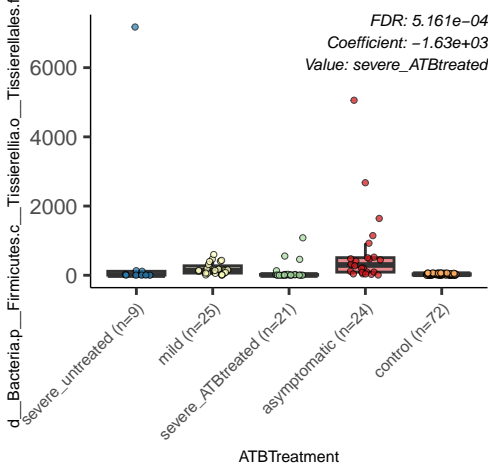


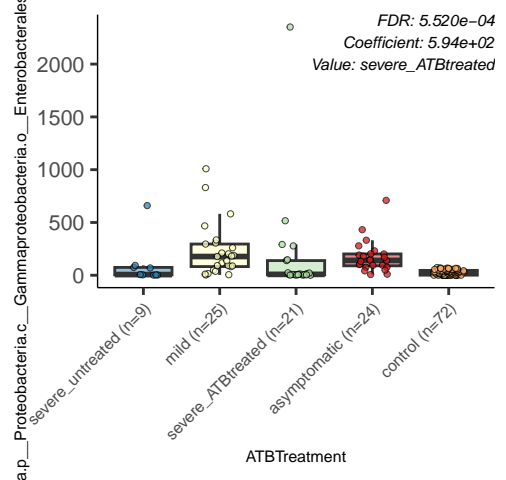


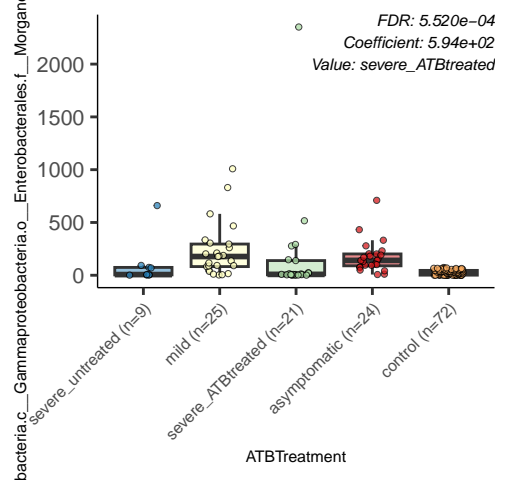




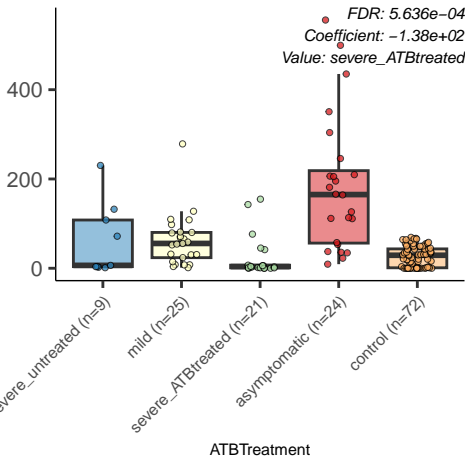


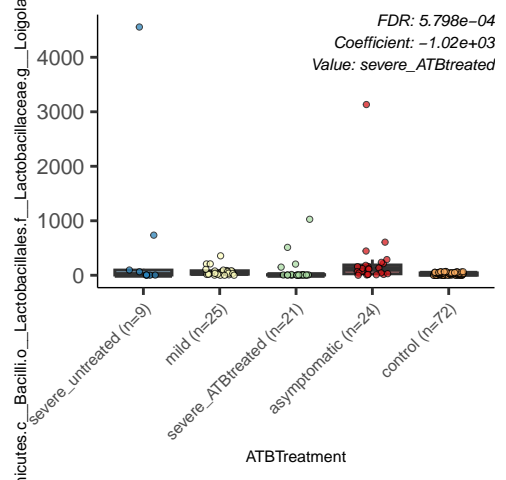


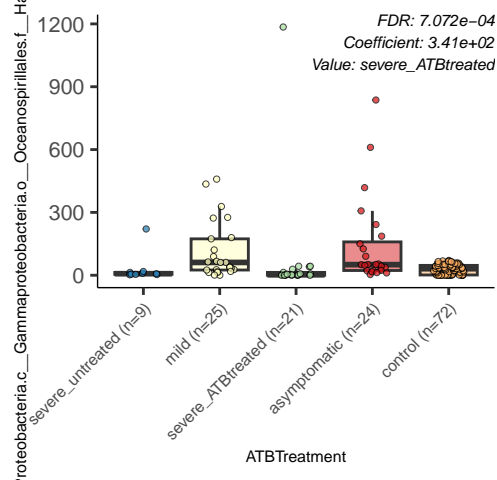




d_Bacteria.p_Firmicutes.c_Clostridia.o_Halanaerobiales.f

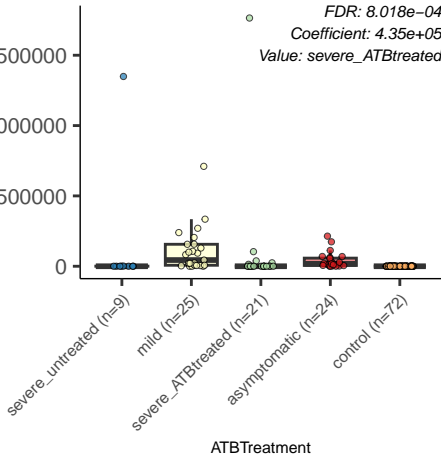


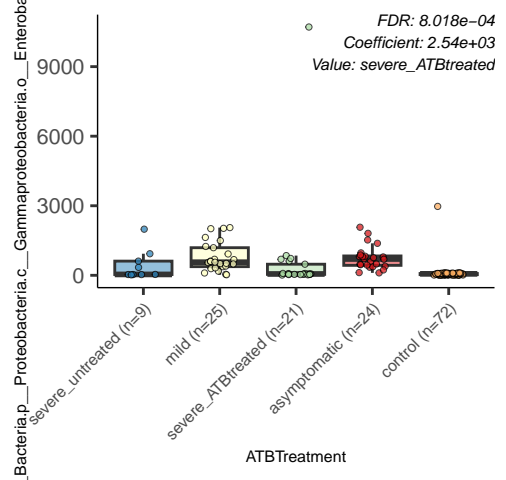




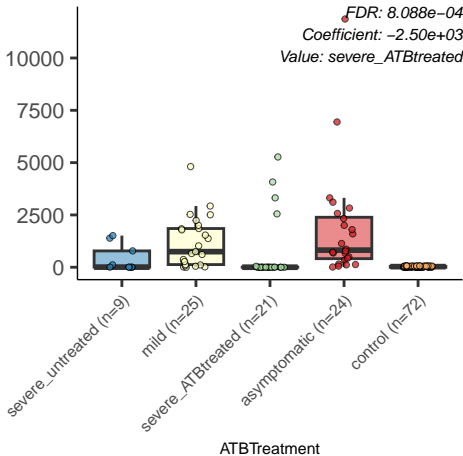
terial.c__Gammaproteobacteria.o__Pasteurellales.f__Pasteurellaceae

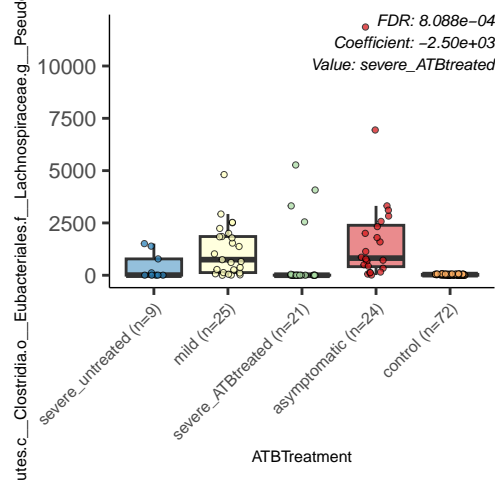
FDR: 8.018e-04
Coefficient: 4.35e+05
Value: severe_ATBtreated

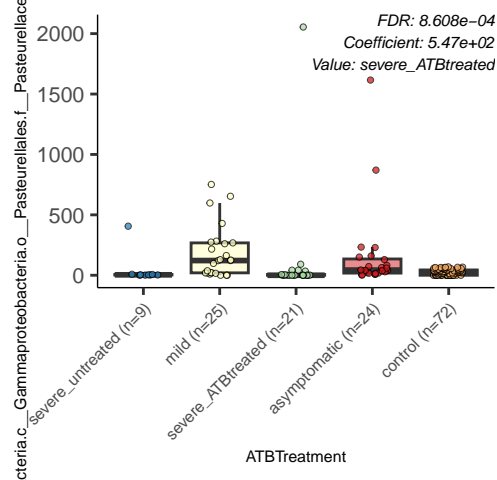


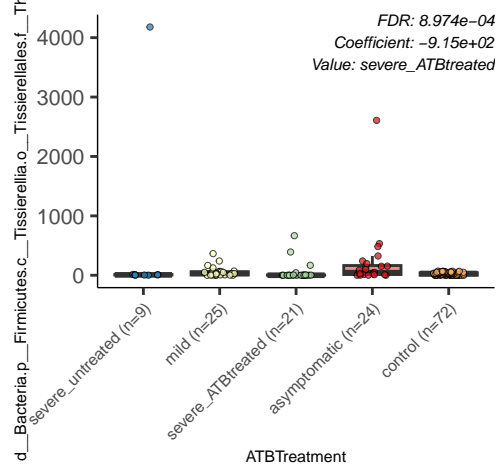


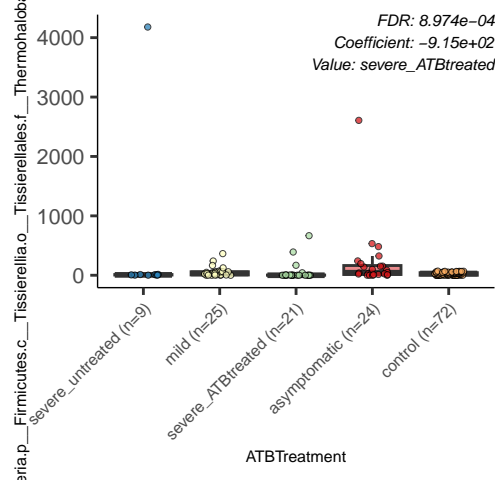
bacteria.p_Firmicutes.c_Clostridia.o_Eubacteriales.f_Lachnosp

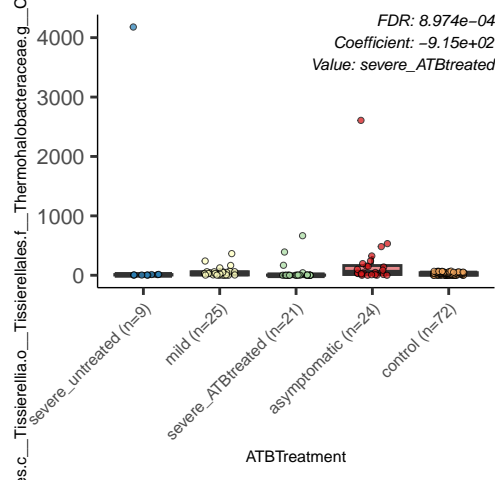


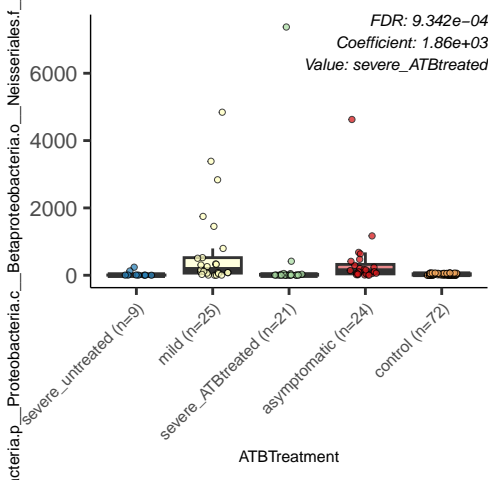


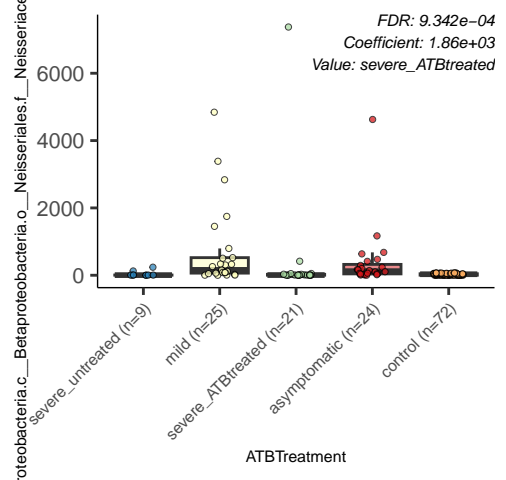


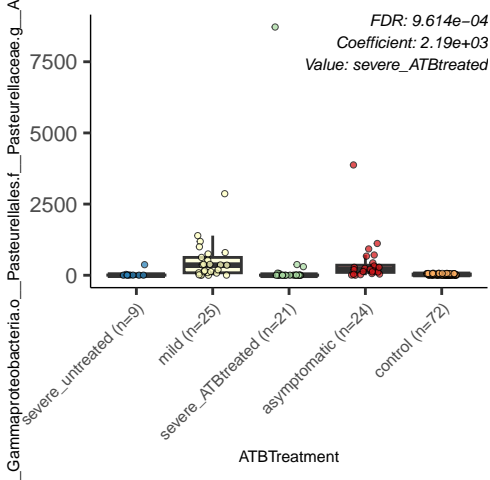


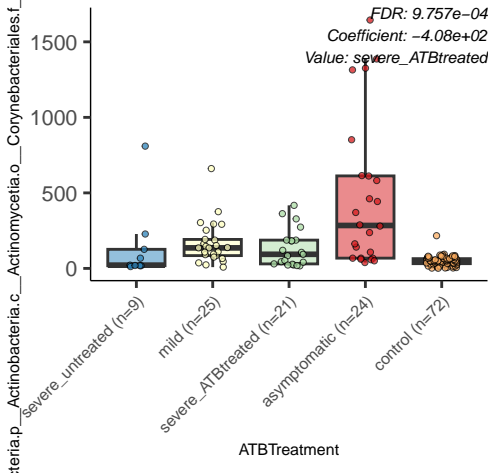


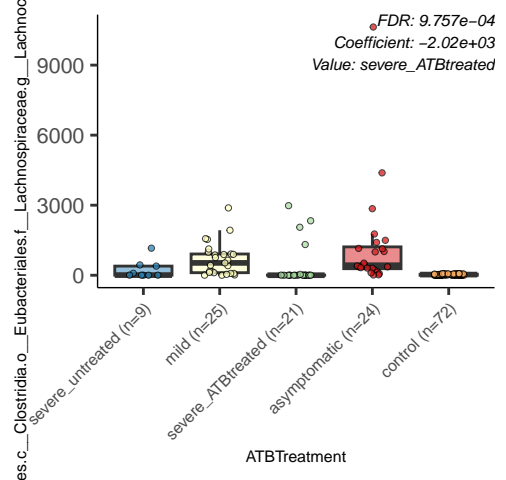






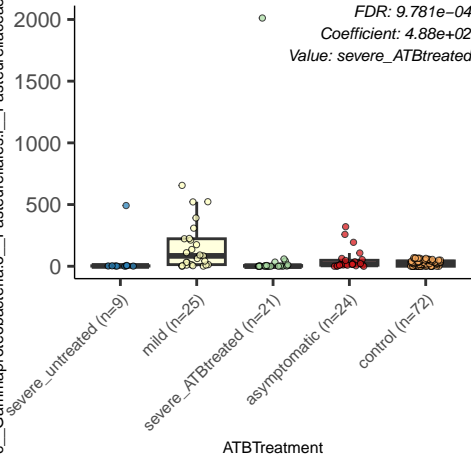


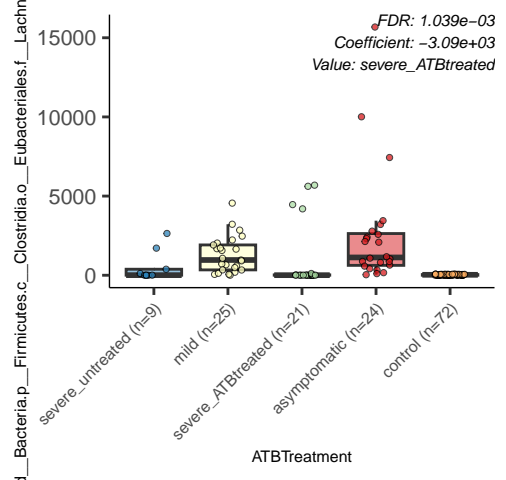


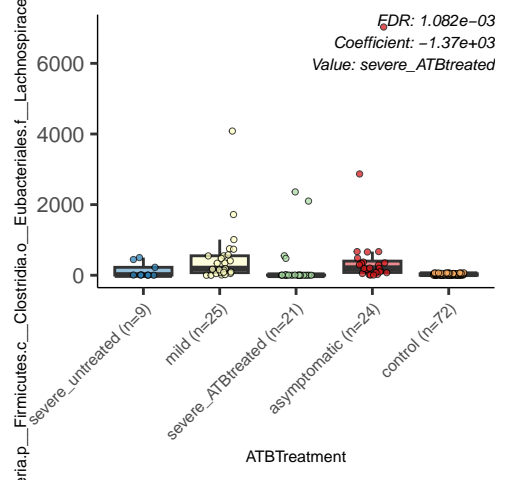


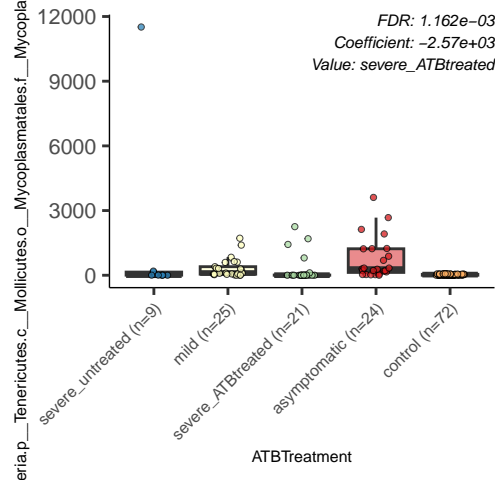
ia.c__Gammaproteobacteria.o__Pasteurellales.f__Pasteurellaceae.

FDR: $9.781e-04$
Coefficient: $4.88e+02$
Value: severe_ATBtreated

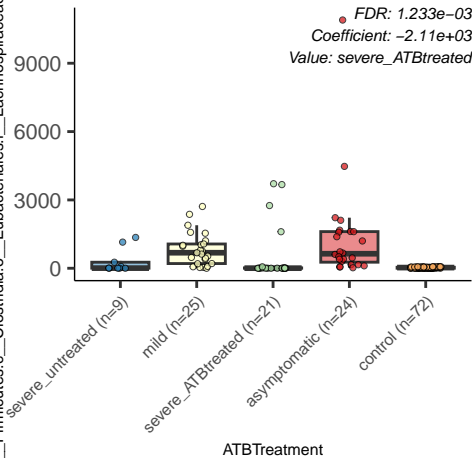


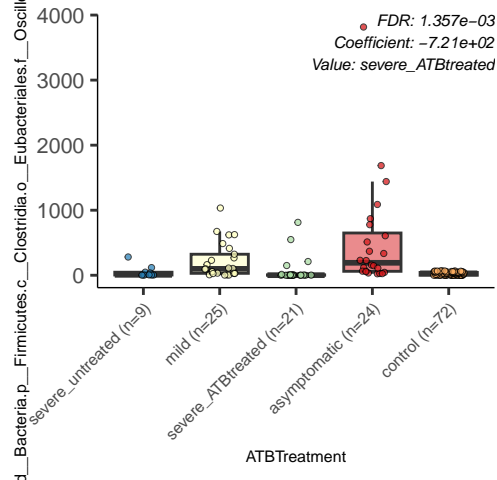


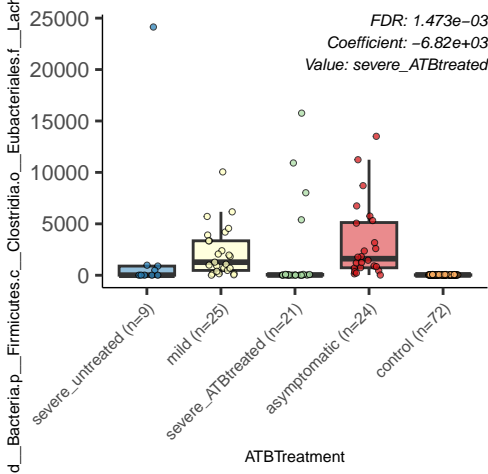


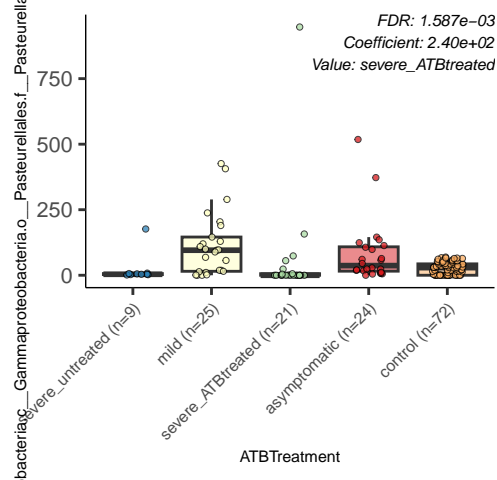


p_Firmicutes.c_Clostridia.o_Eubacteriales.f_Lachnospiraceae

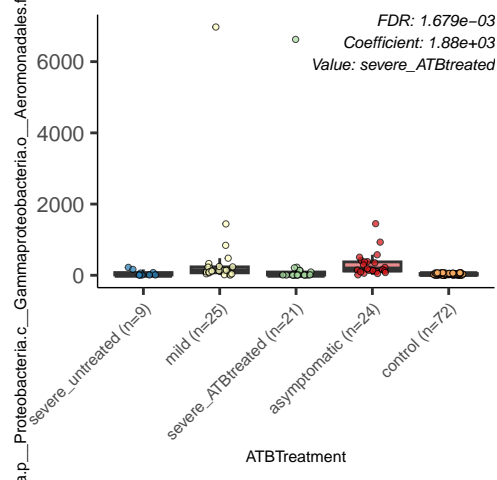


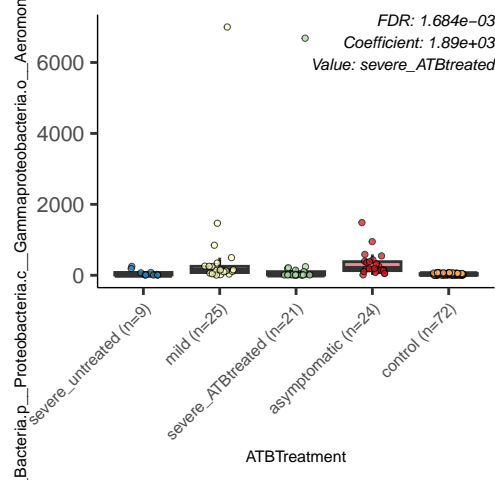


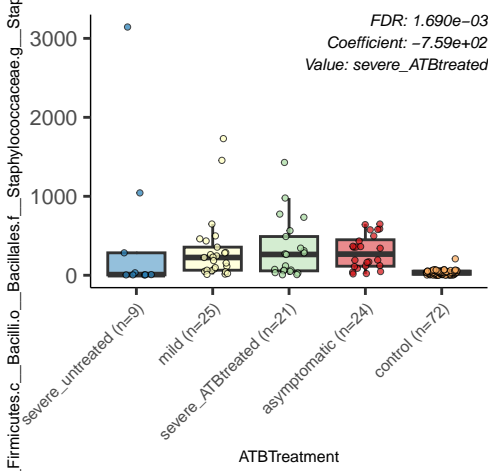


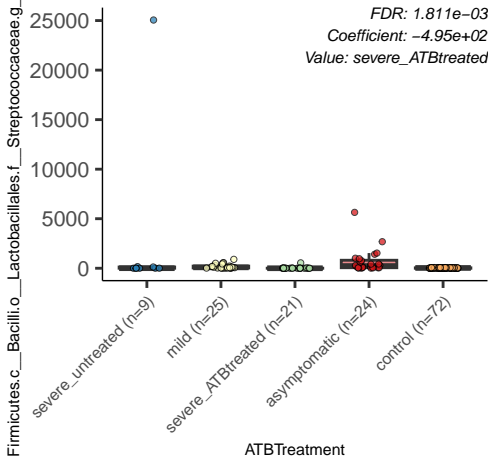


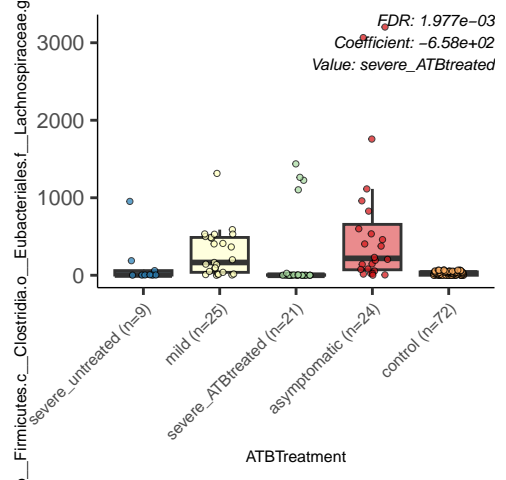
a.p__Proteobacteria.c__Gammaproteobacteria.o__Aeromonadales.

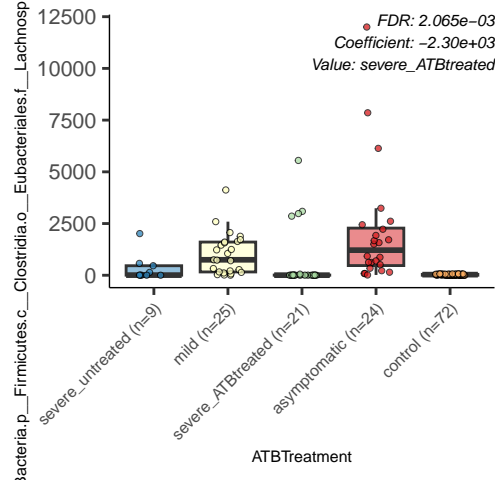


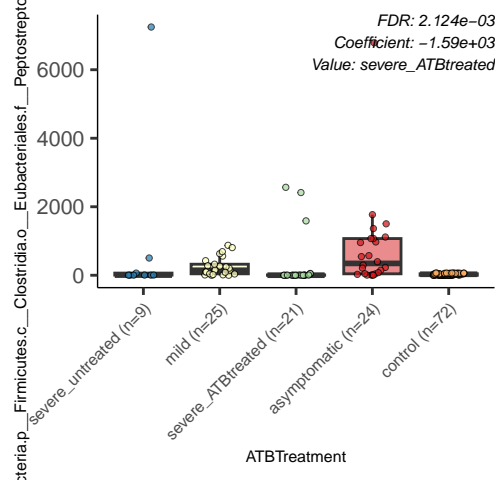


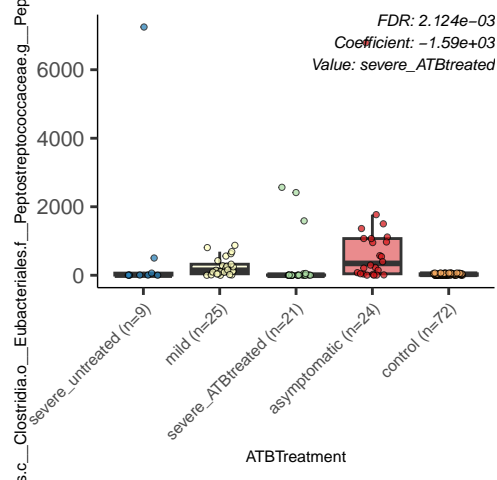


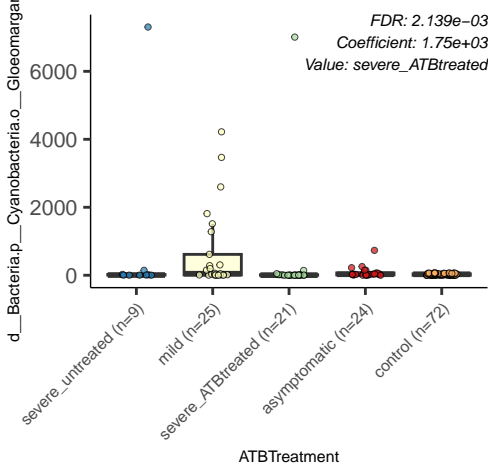


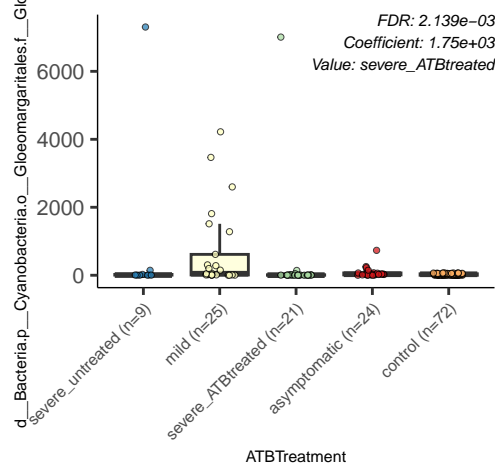


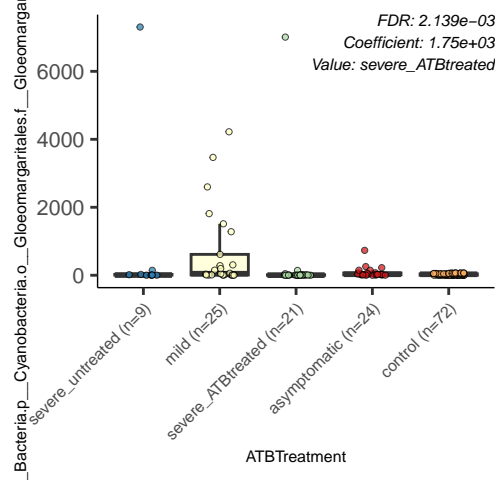


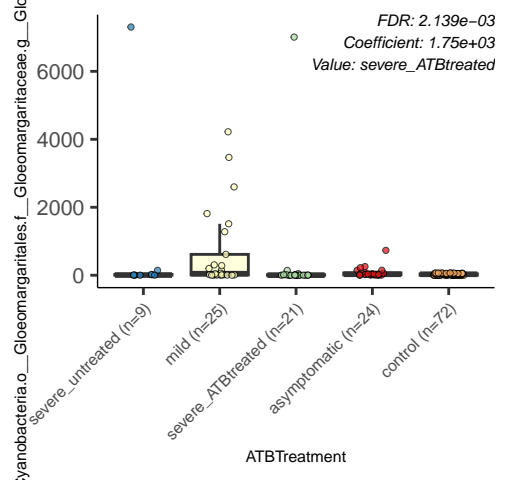


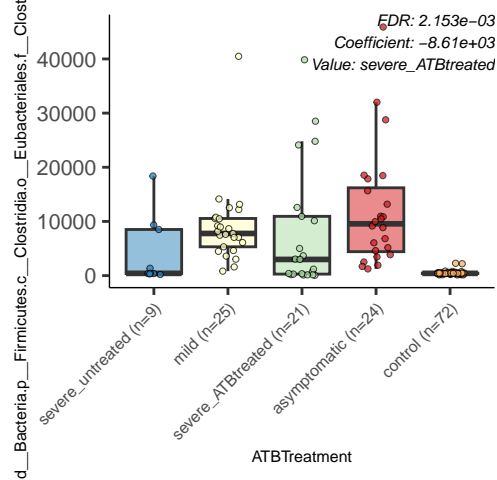




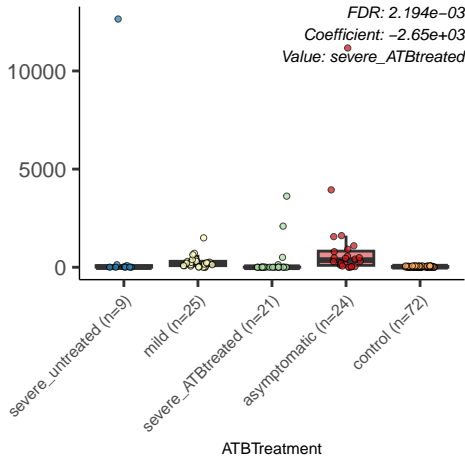




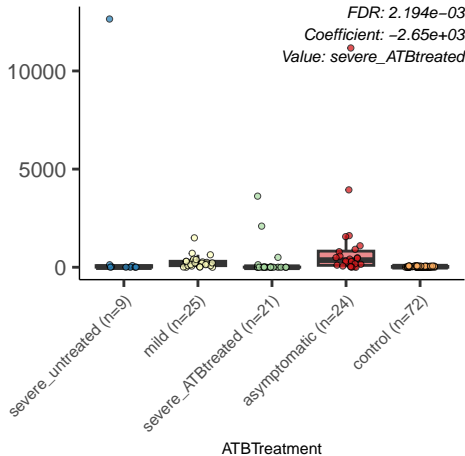


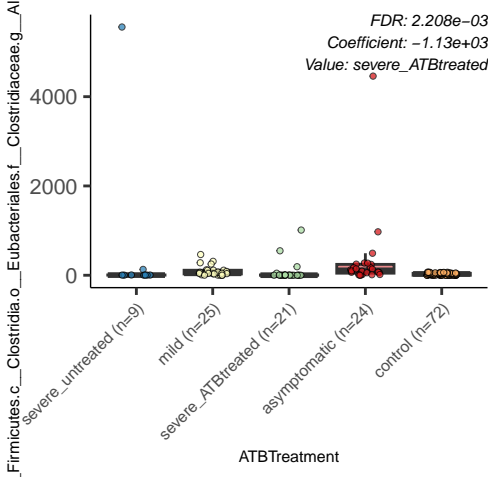


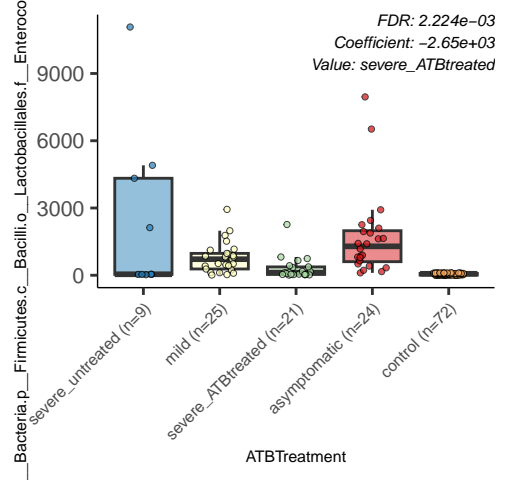
Bacteria.p_Firmicutes.c_Clostridia.o_Eubacteriales.f_Peptostreptococcaceae

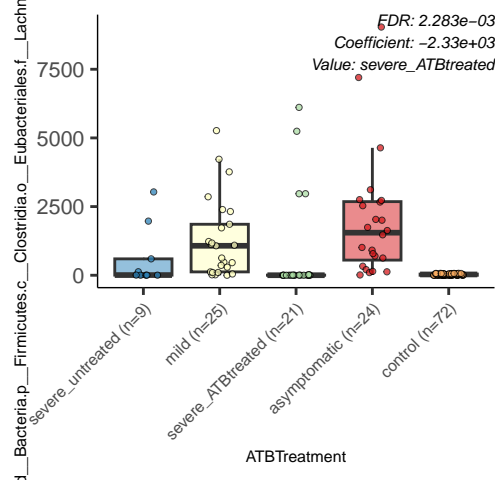


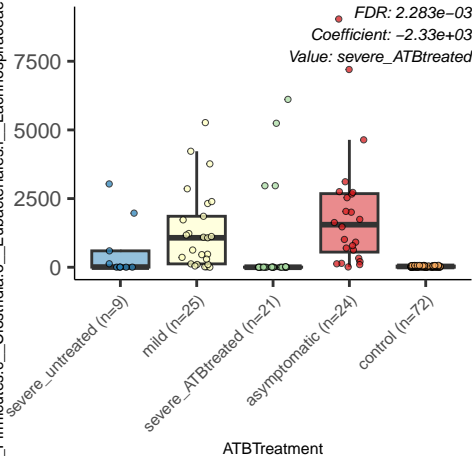
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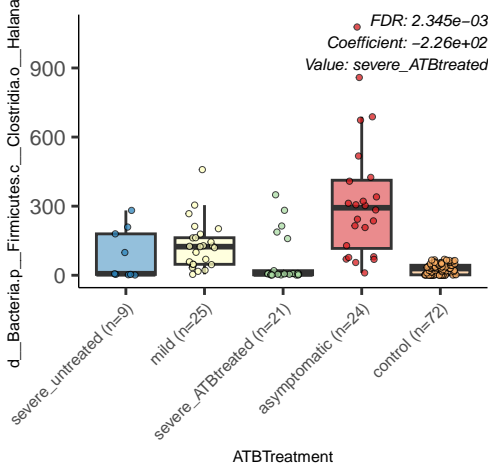


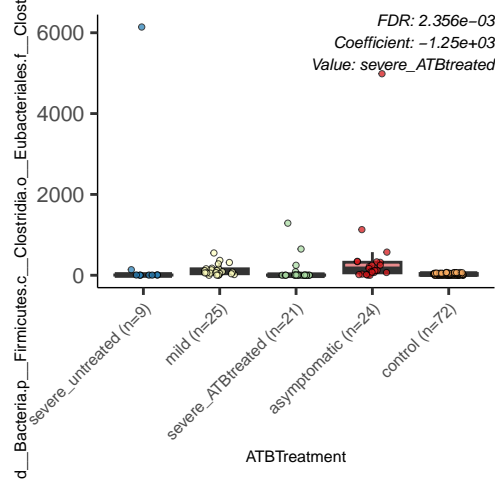


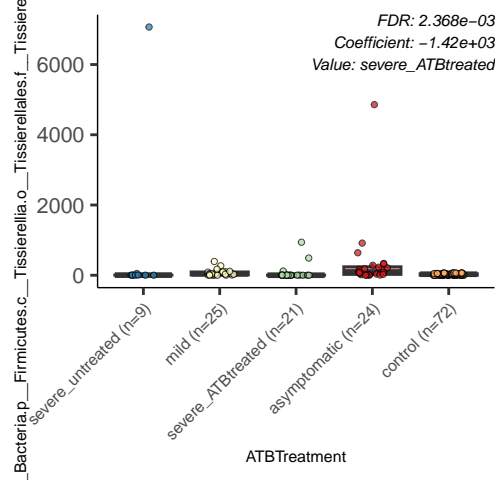


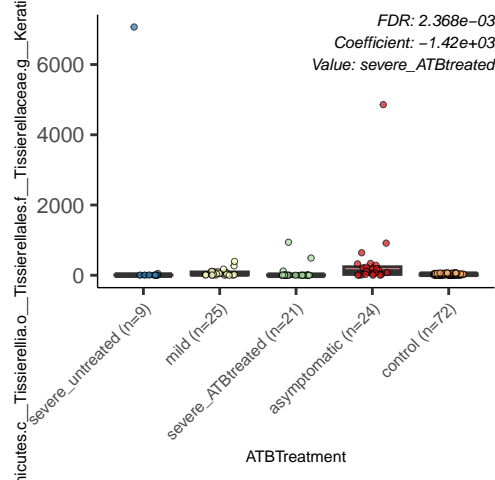


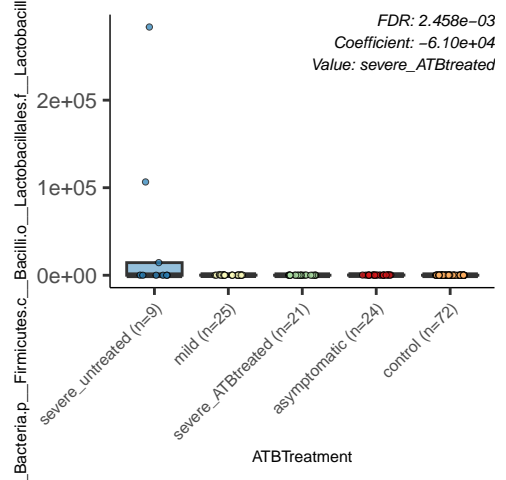


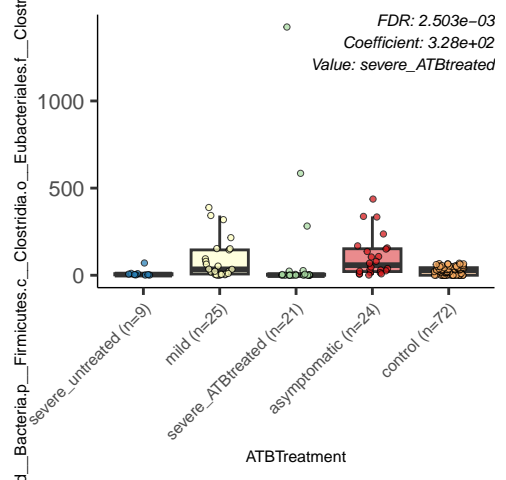




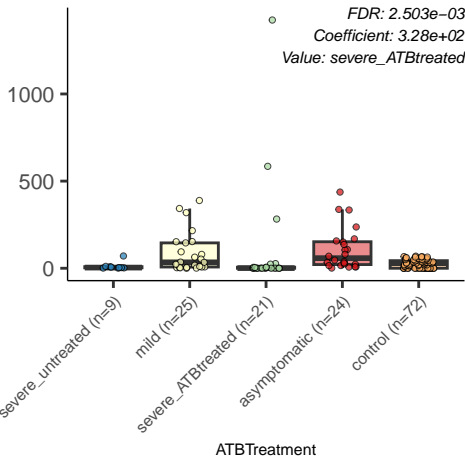


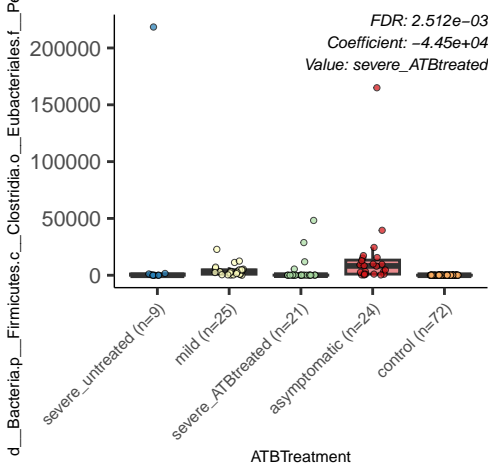


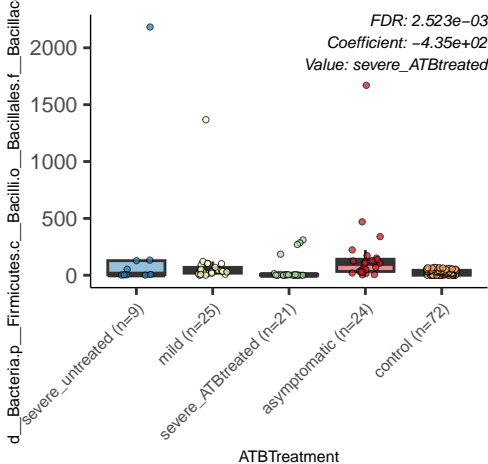


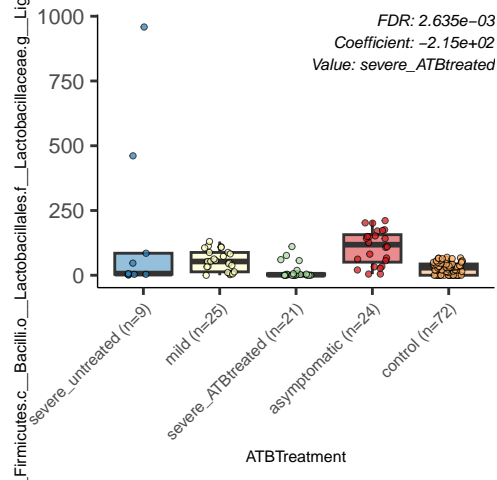


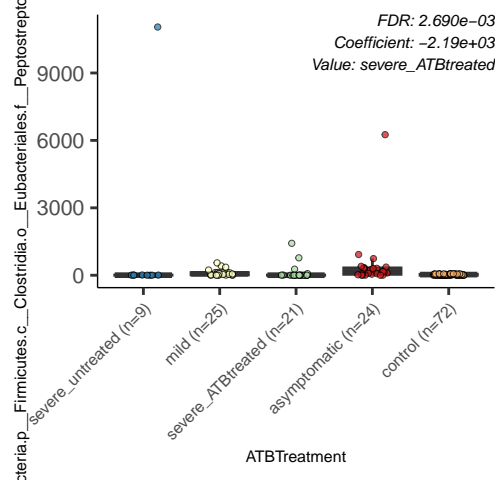
p_Firmicutes.c_Clostridia.o_Eubacteriales.f_Clostridiaceae.g



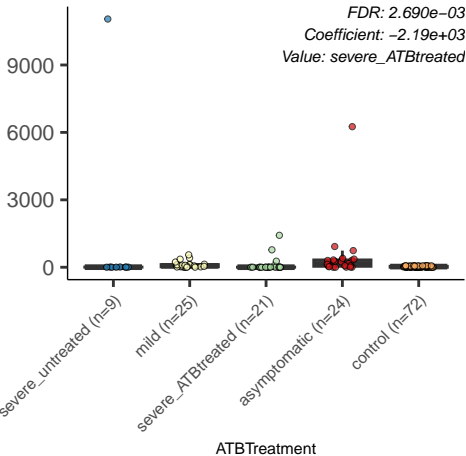




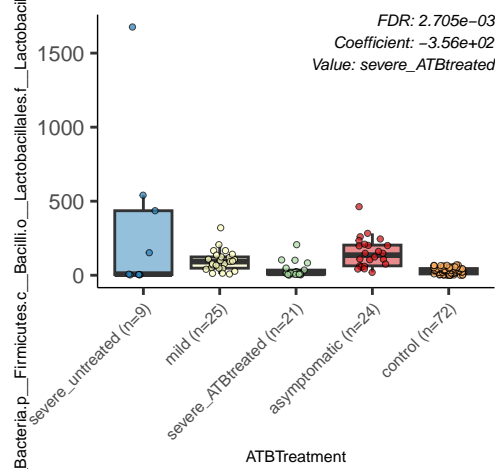


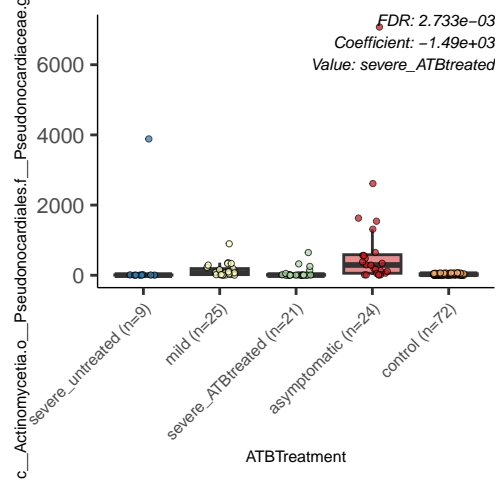


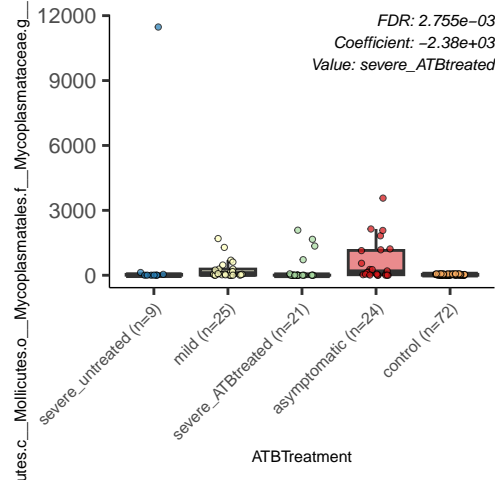
cutes.c__Clostridia.o__Eubacteriales.f__Peptostreptococaceae.g__

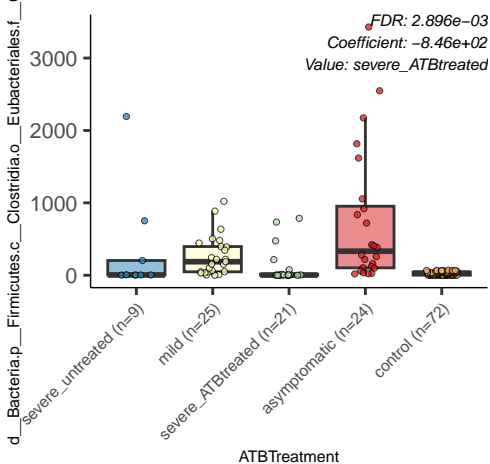


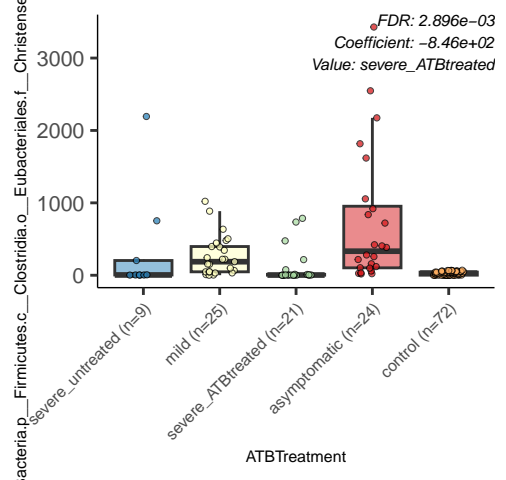
Bacteria.p_Firmicutes.c_Bacilli.o_Lactobacillales.f_Lactobacilli

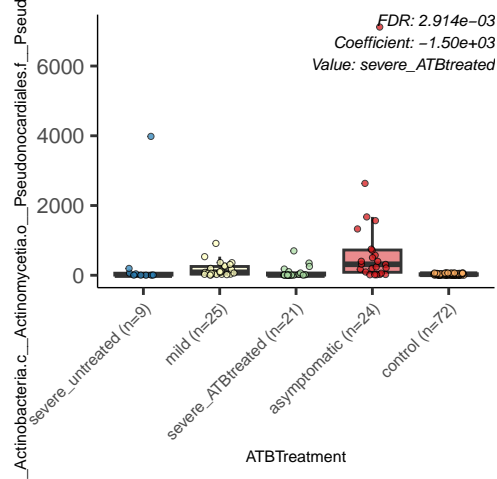


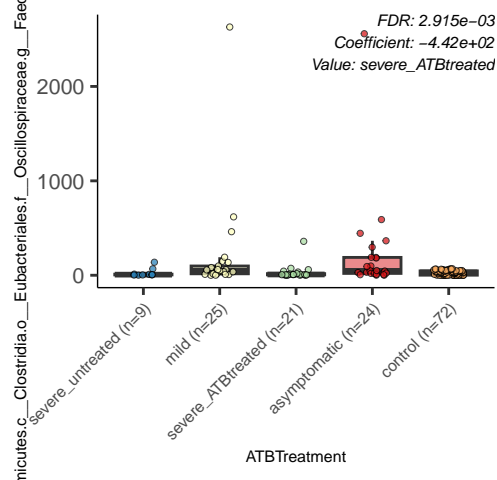


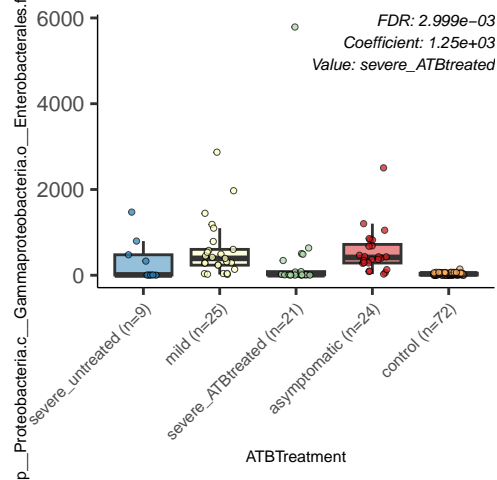


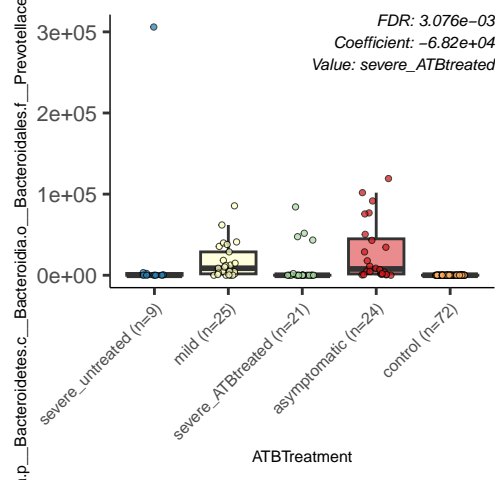


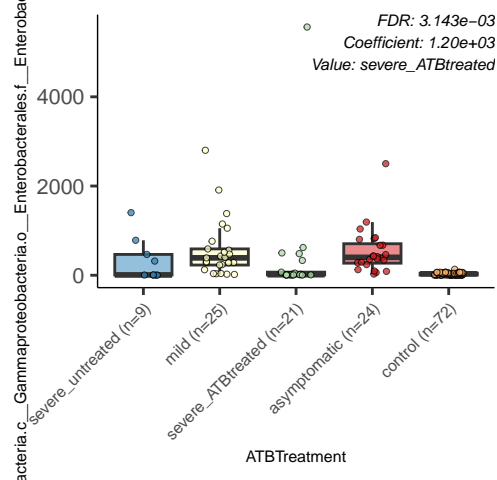


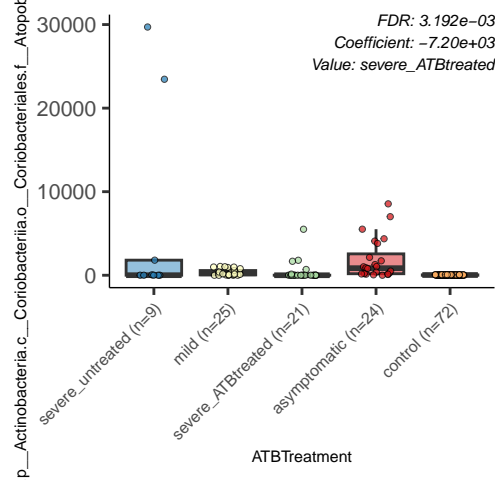


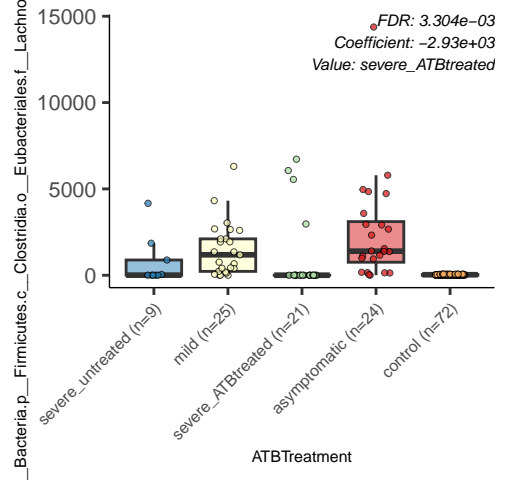


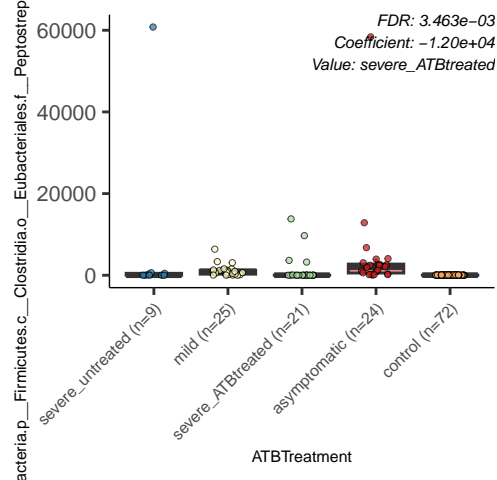




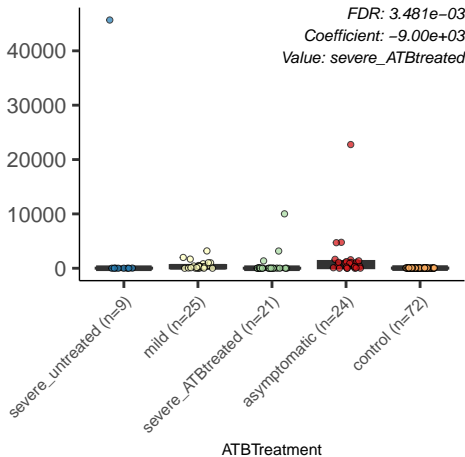


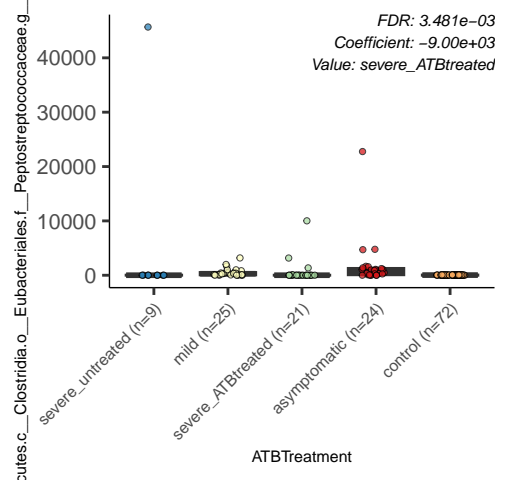


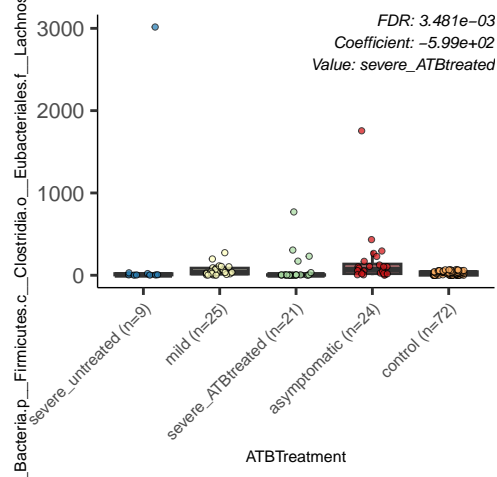


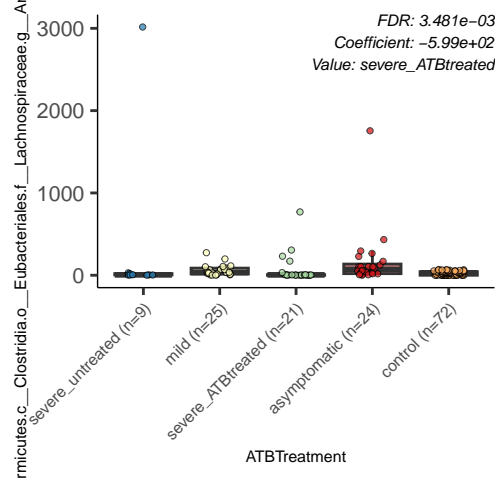


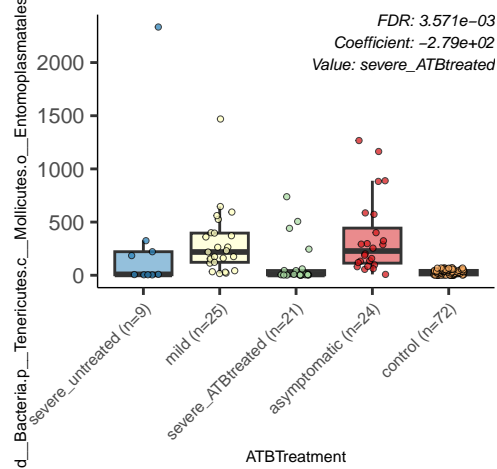
bacteria.p__Firmicutes.c__Clostridia.o__Eubacteriales.f__Peptostreptococcus

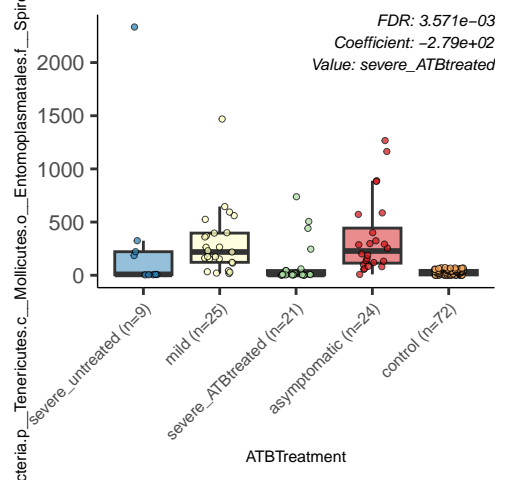


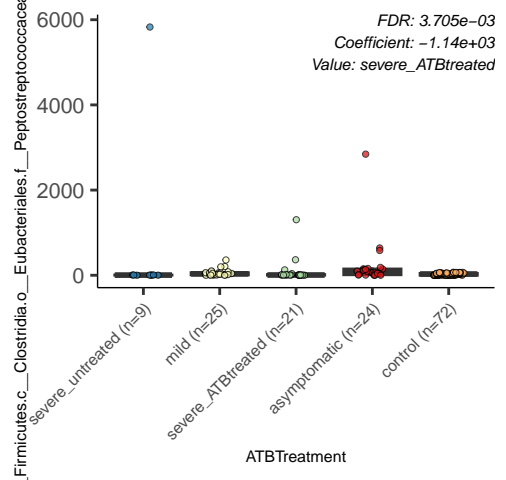




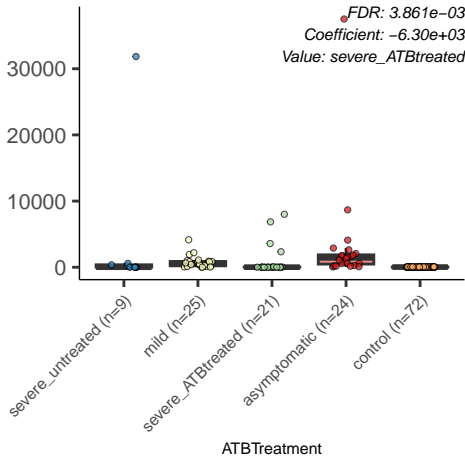


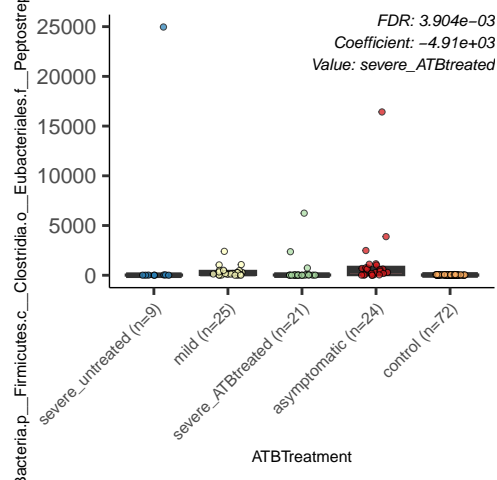






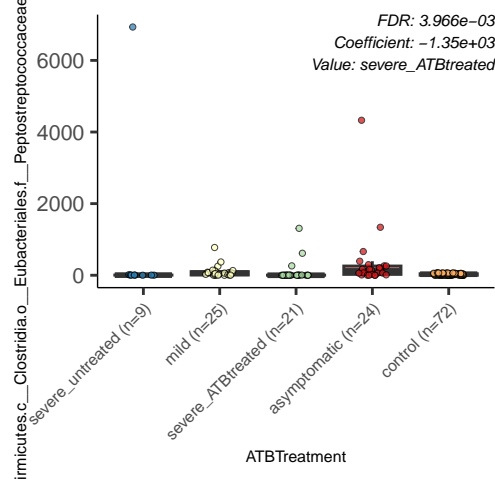
irmicutes.c__Clostridia.o__Eubacteriales.f__Peptostreptococcaceae

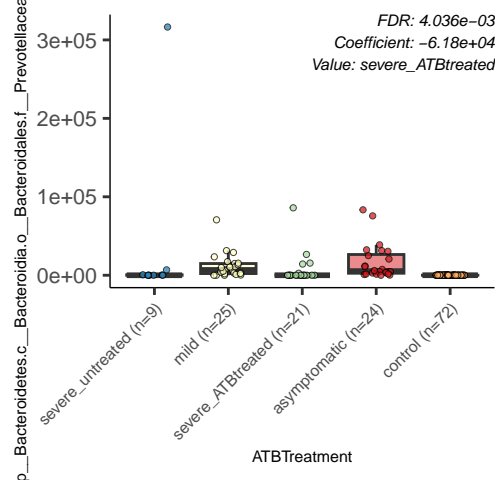


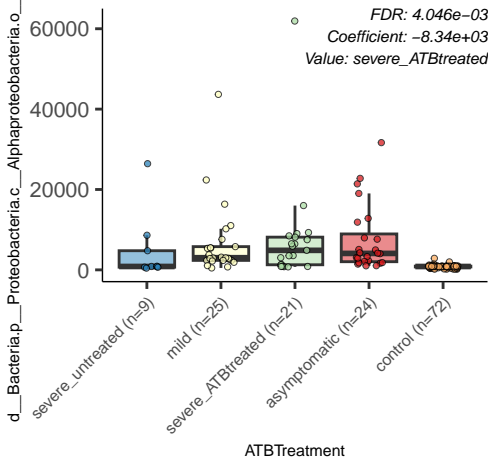


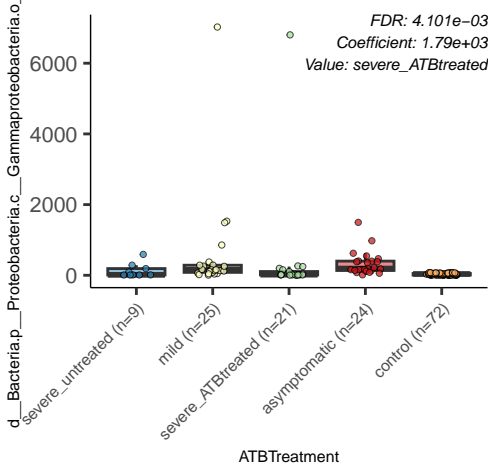
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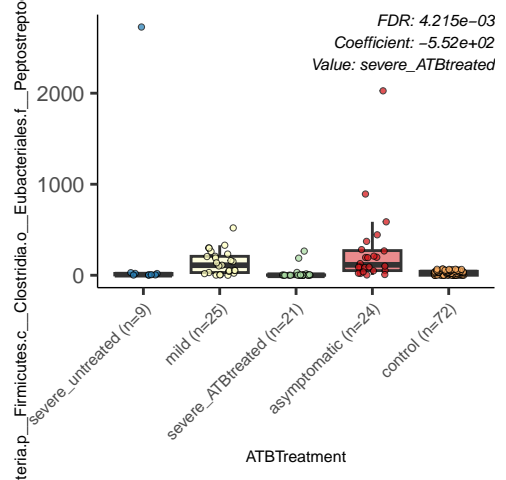
FDR: $3.966e-03$
Coefficient: $-1.35e+03$
Value: severe_ATBtreated

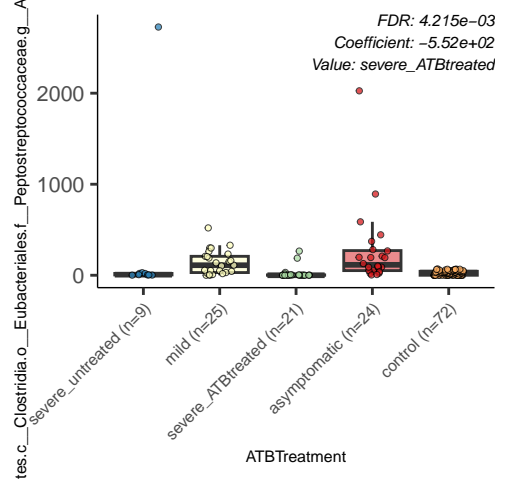


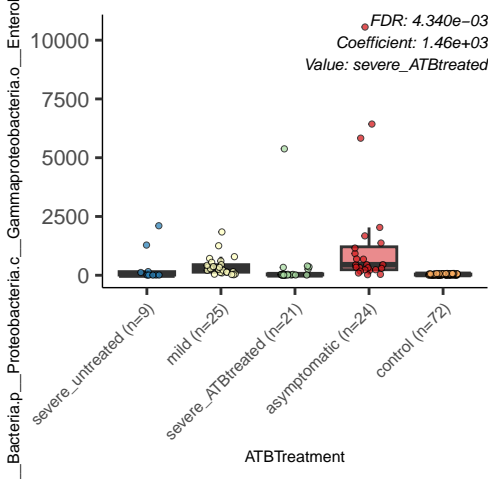


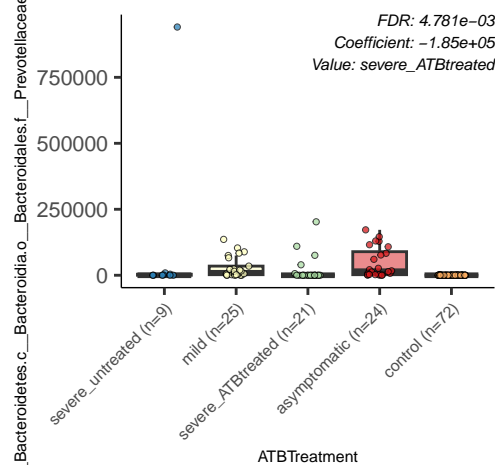


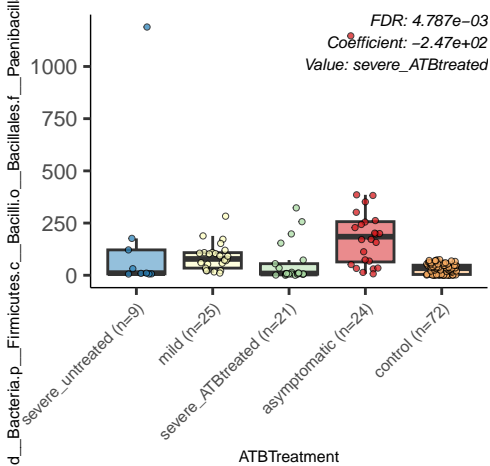


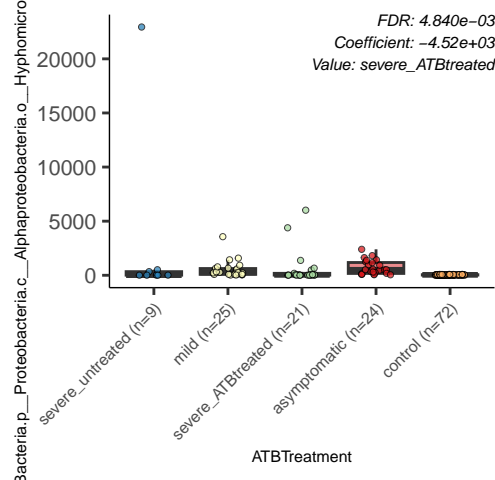


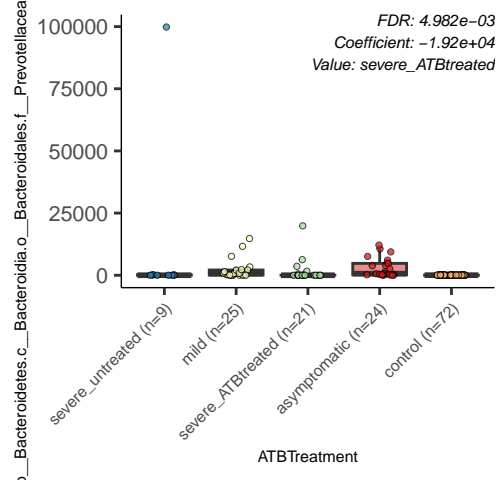


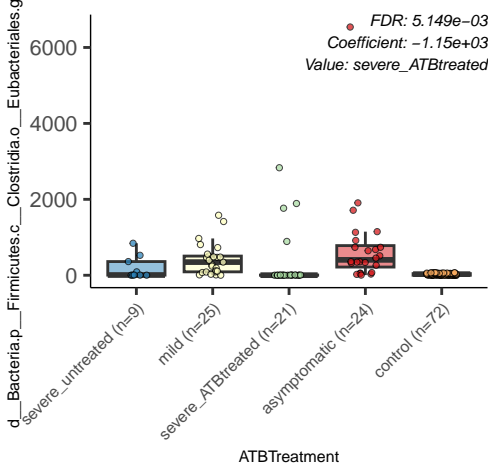


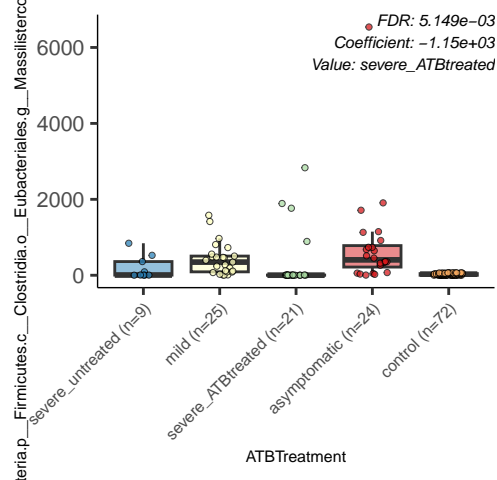


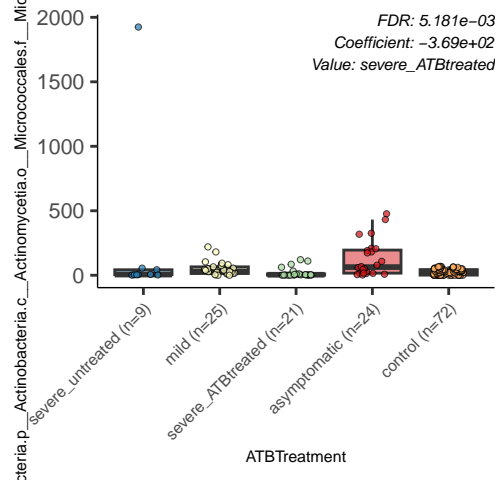


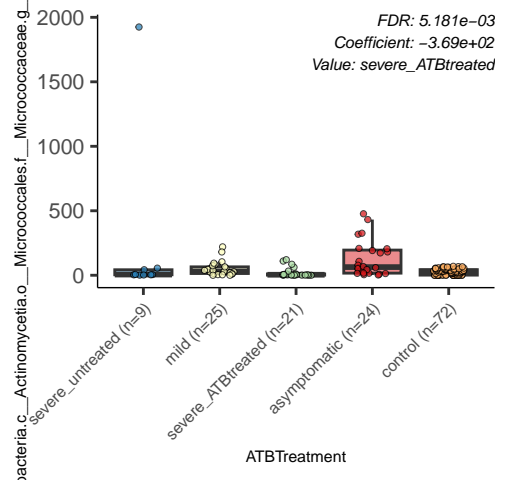


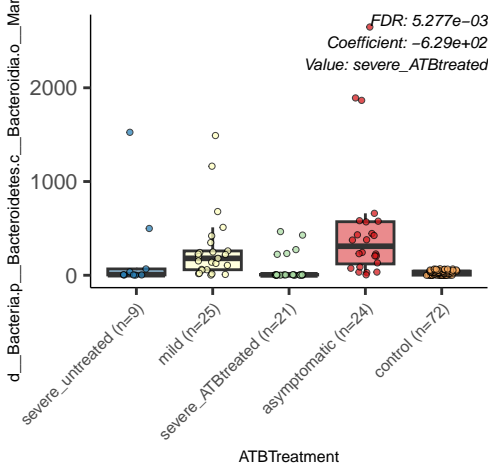


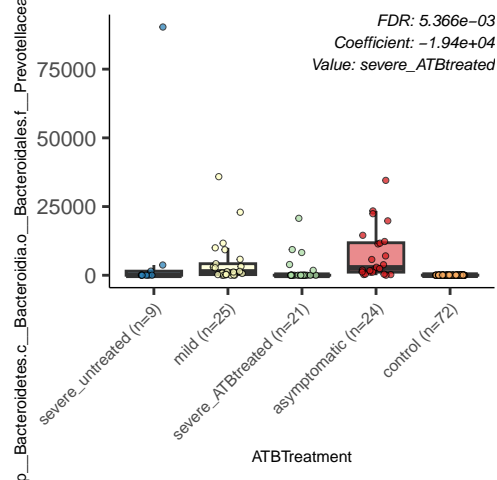


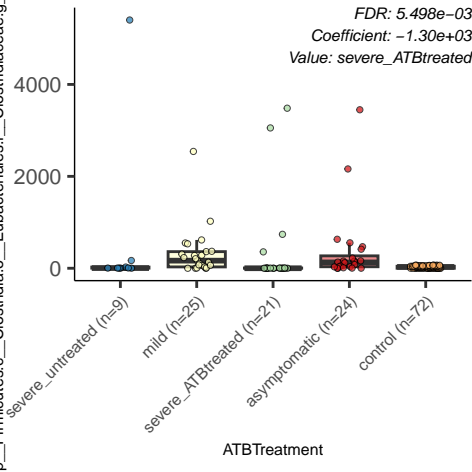


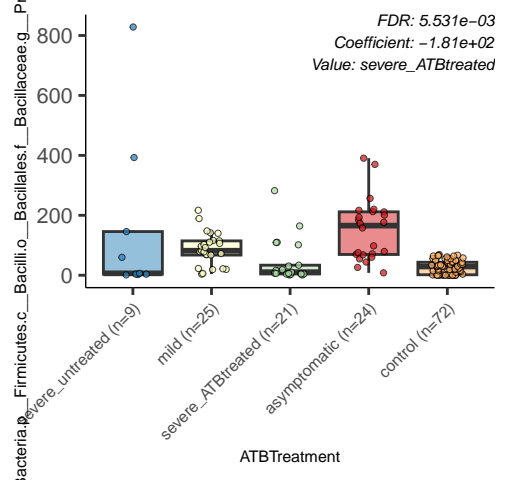


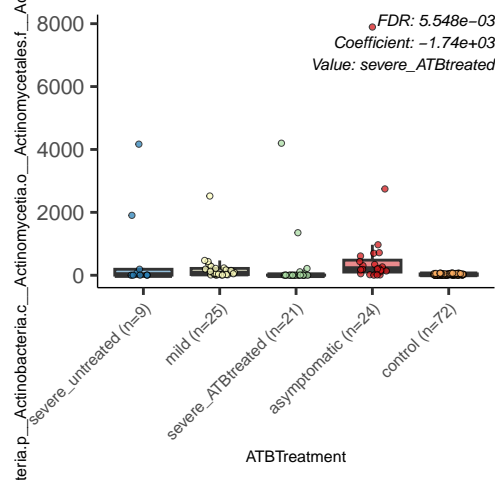


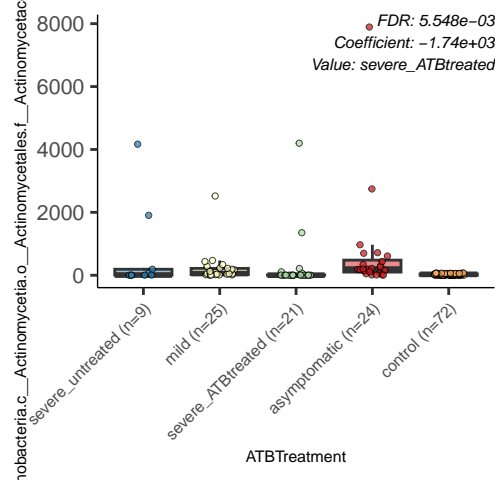






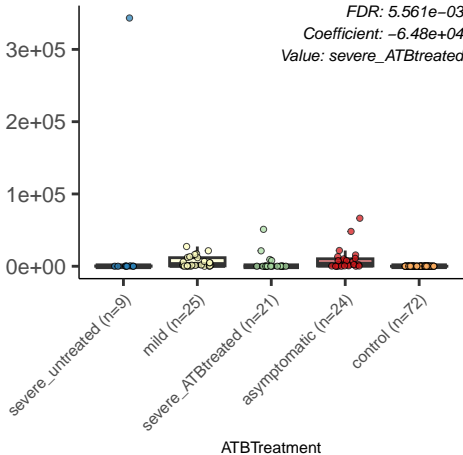


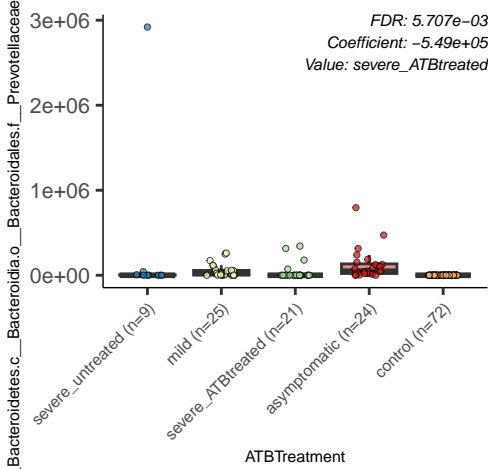




p__Bacteroidetes.c__Bacteroidia.o__Bacteroidales.f__Prevotellaceae

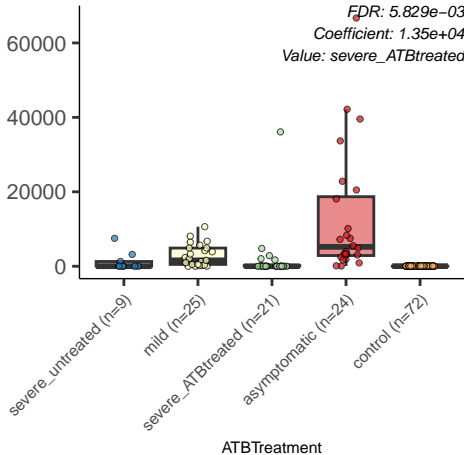
FDR: $5.561e-03$
Coefficient: $-6.48e+04$
Value: severe_ATBtreated

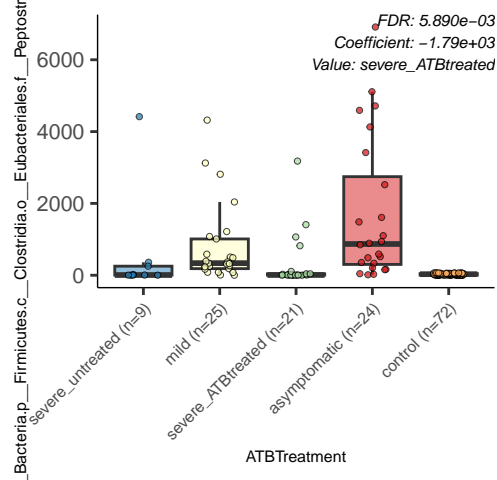


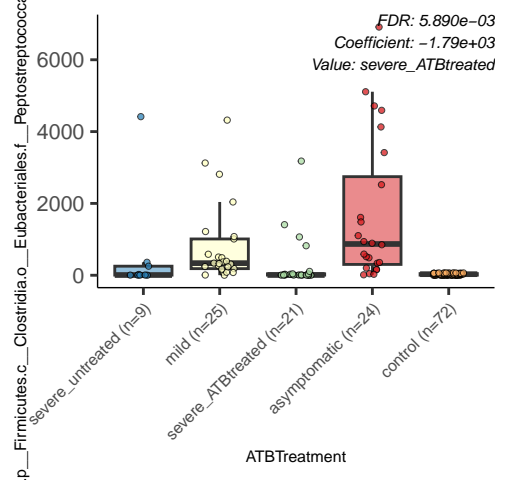


Firmicutes.c__Bacilli.o__Lactobacillales.f__Streptococcaceae.g__Str

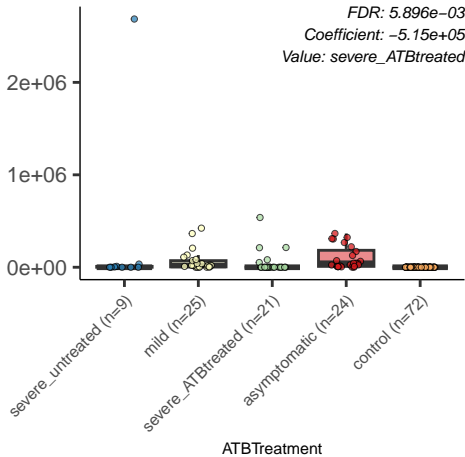
FDR: 5.829×10^{-3}
Coefficient: 1.35×10^4
Value: severe_ATBtreated



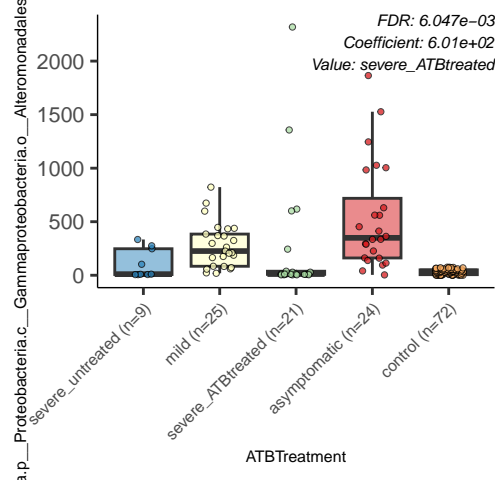


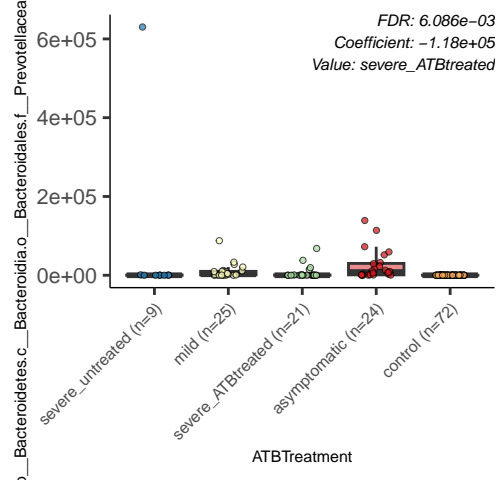


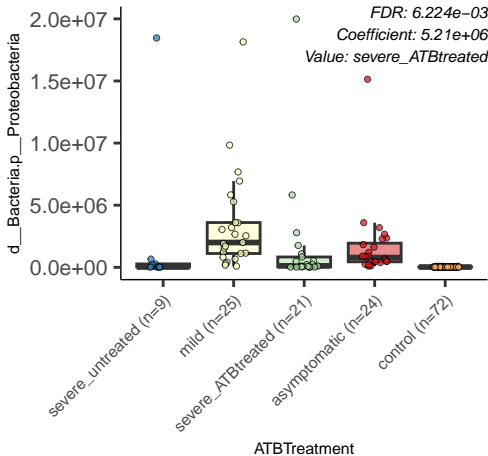
a.p._Bacteroidetes.c._Bacteroidia.o._Bacteroidales.f._Prevotellac

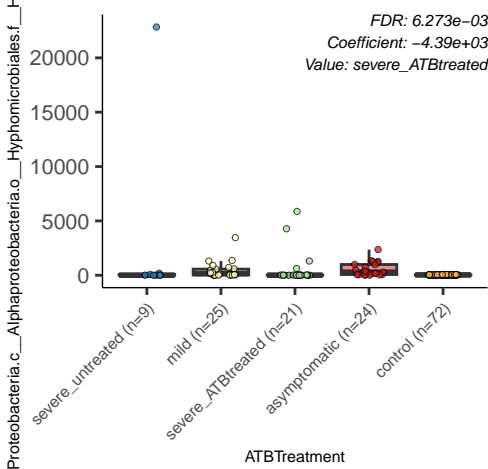


a.p__Proteobacteria.c__Gammaproteobacteria.o__Alteromonadales



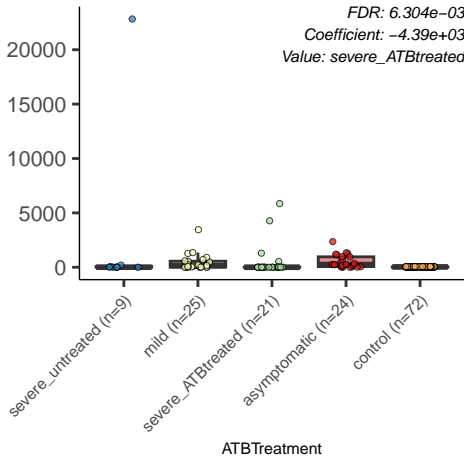


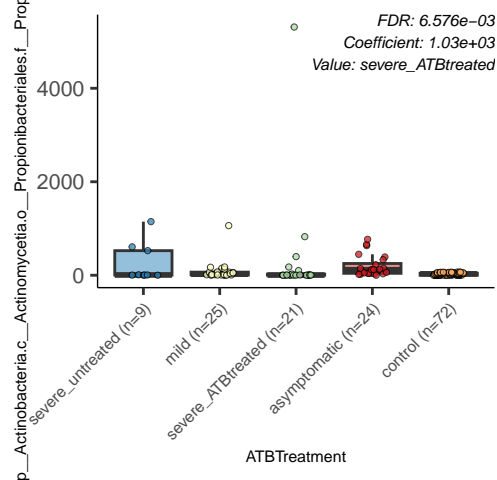


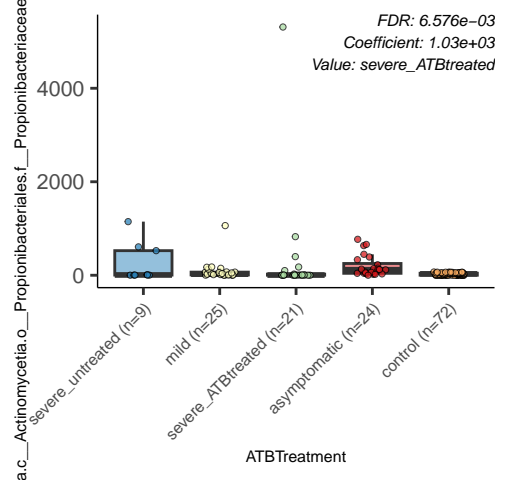


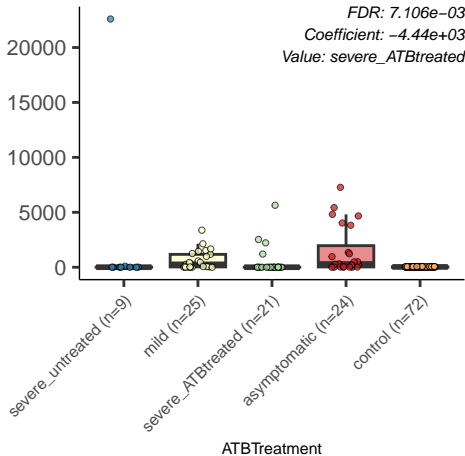
c__Alphaproteobacteria.o__Hyphomicrobiales.f__Hyphomicrobiaceae

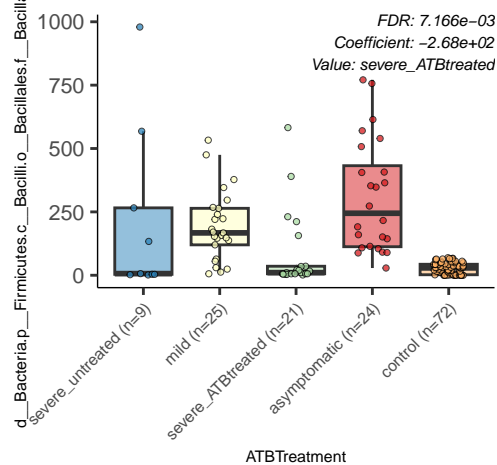
FDR: $6.304e-03$
Coefficient: $-4.39e+03$
Value: severe_ATBtreated

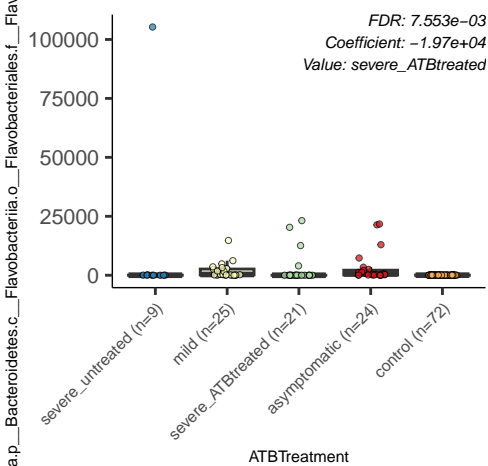




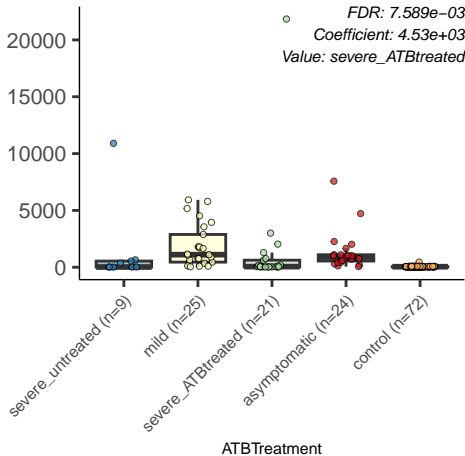


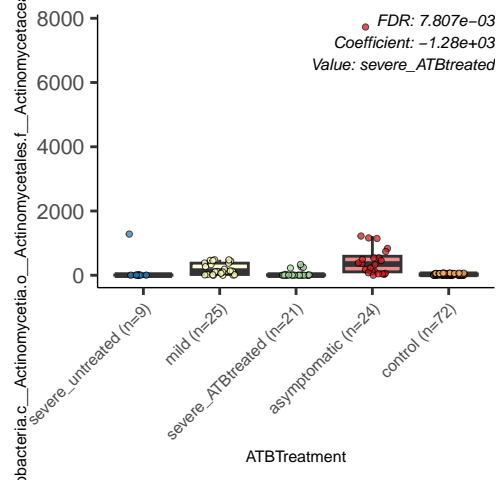




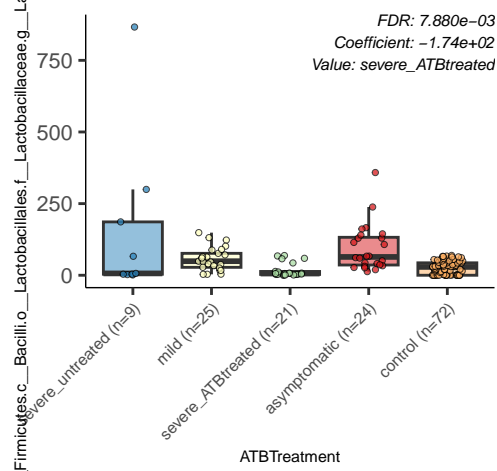


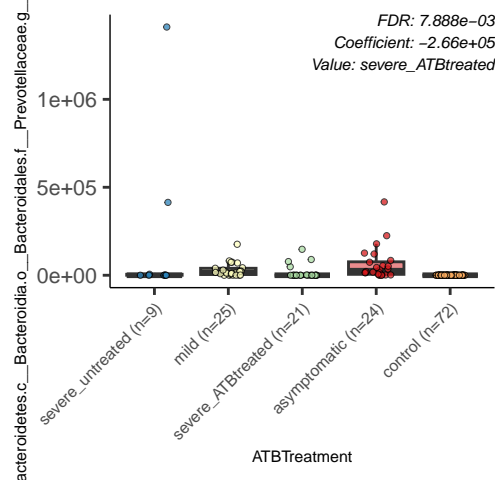
p__Proteobacteria.c__Gammaproteobacteria.o__Enterobacterales.f

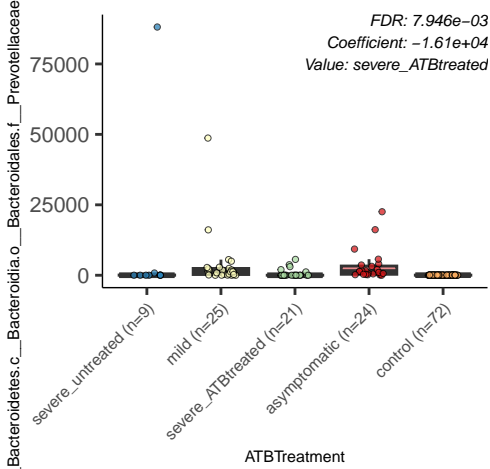


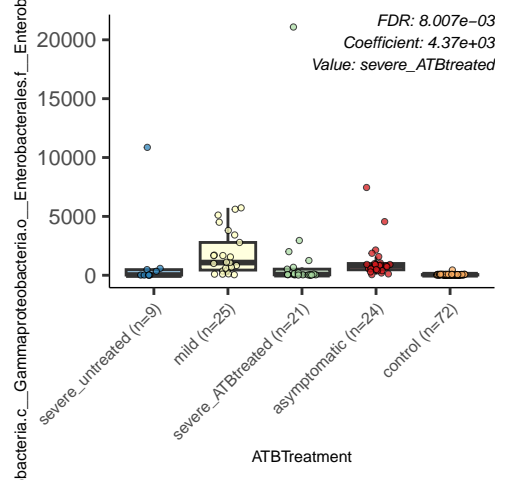


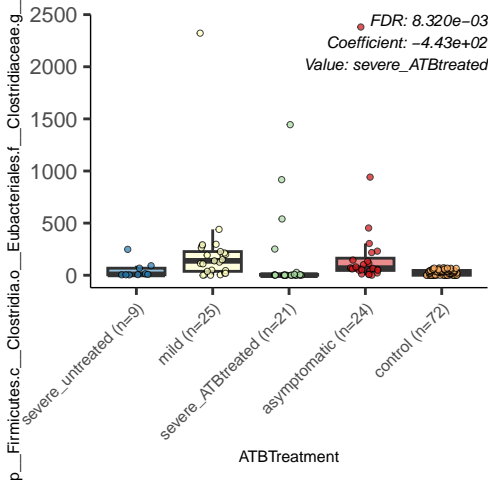
Firmicutes.c_Bacilli.o_Lactobacillales.f_Lactobacillaceae.g_La

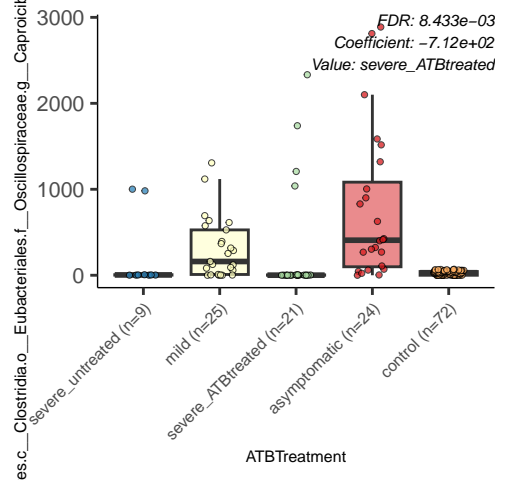


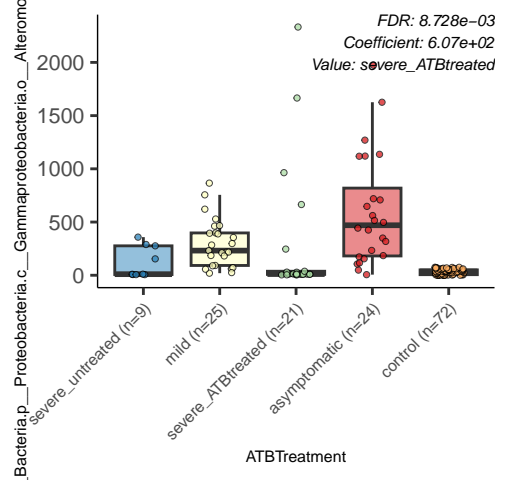




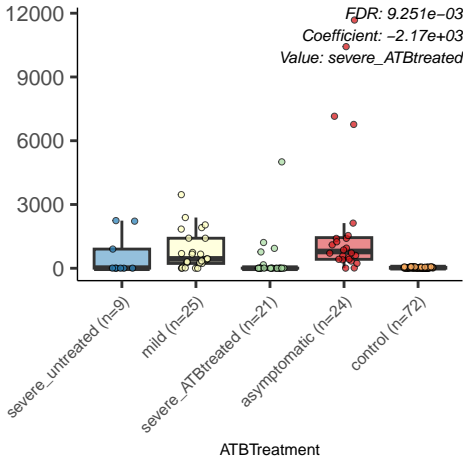


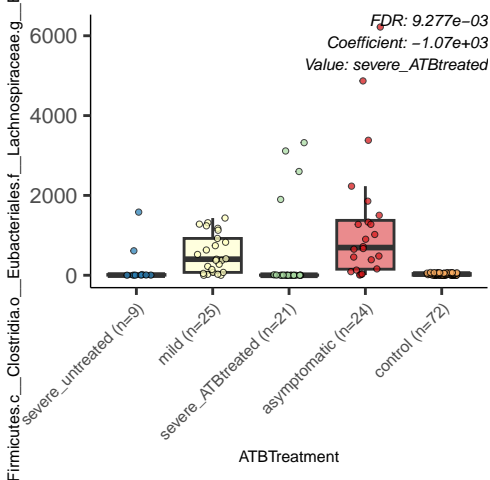


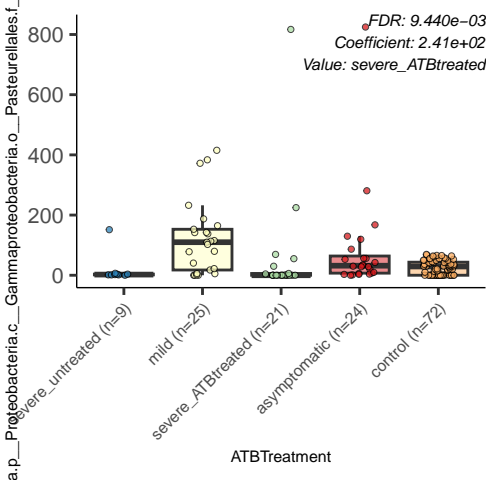


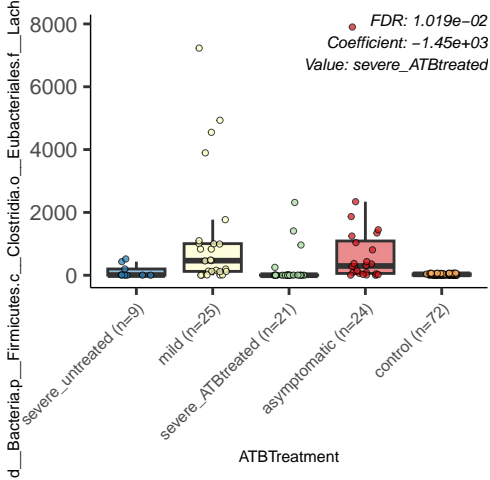


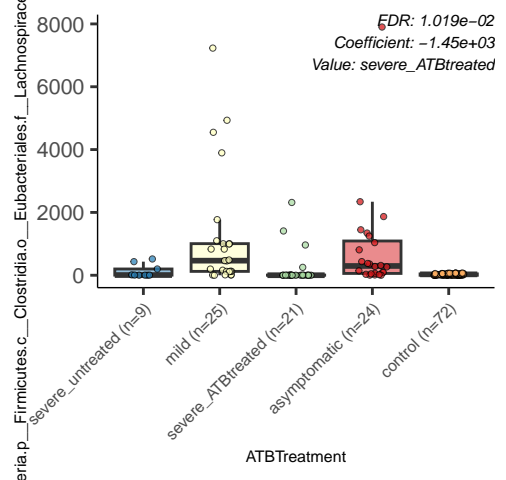
Actinobacteria.c_Actinomyces.f_Actinomyces

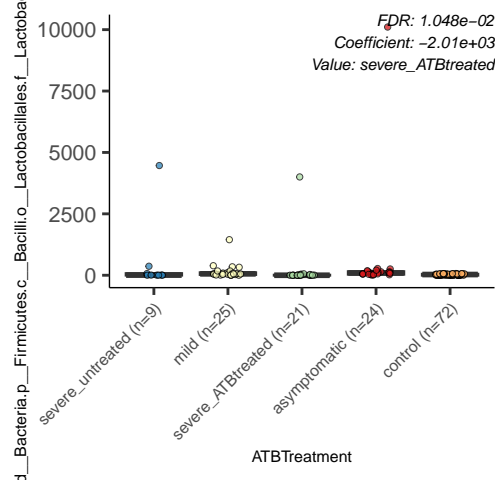


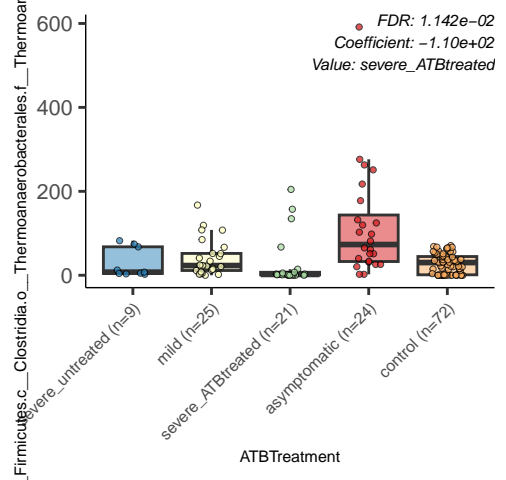


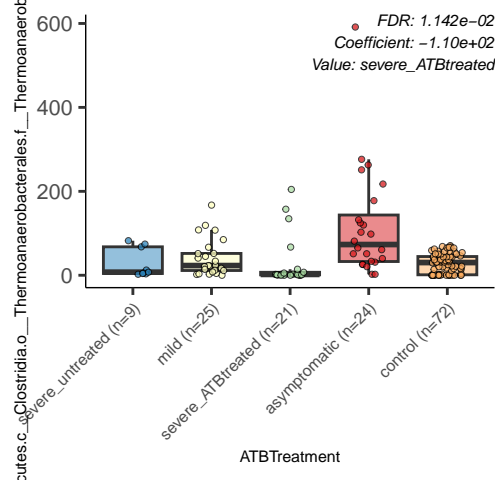


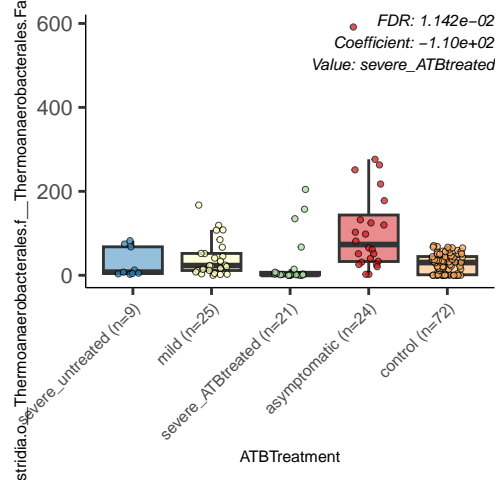




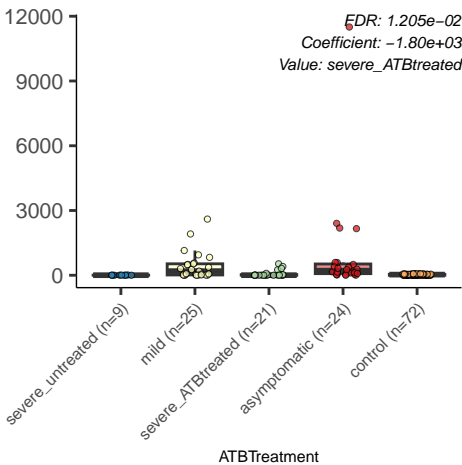


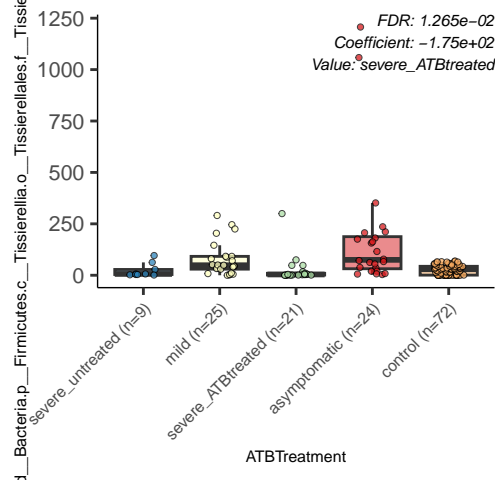


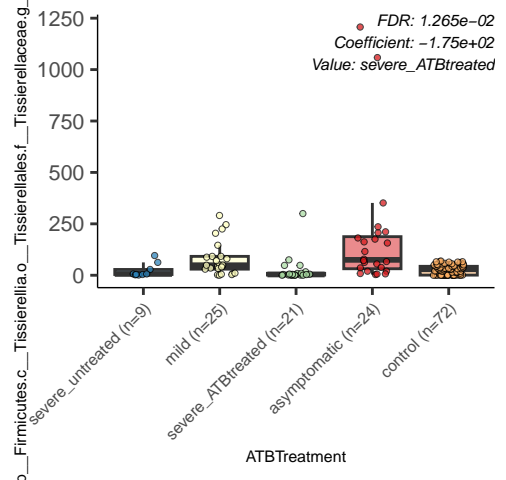


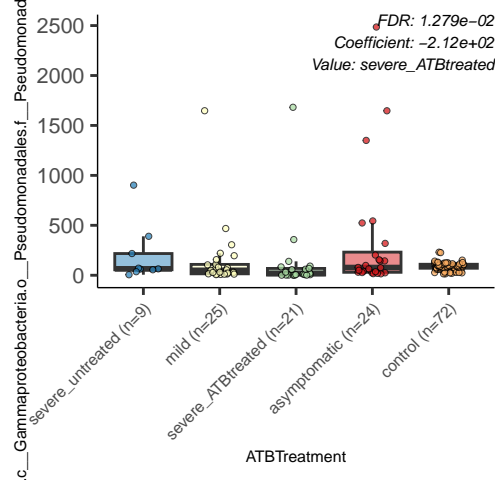


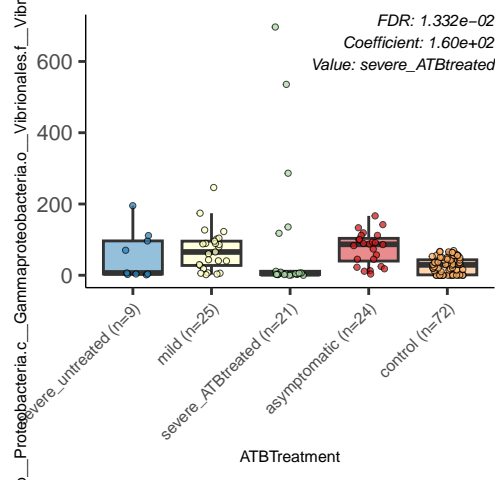
Gammaproteobacteria.o__Cardiobacteriales.f__Cardiobacteriaceae.

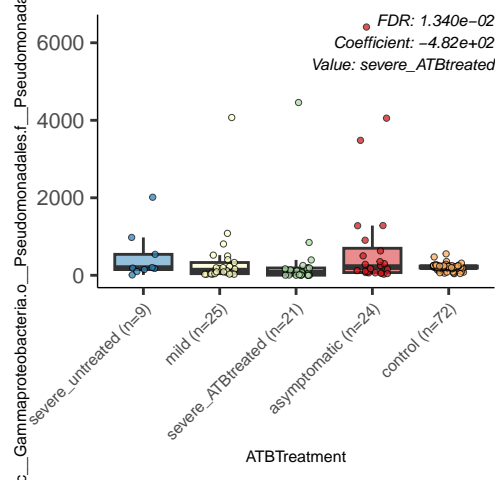


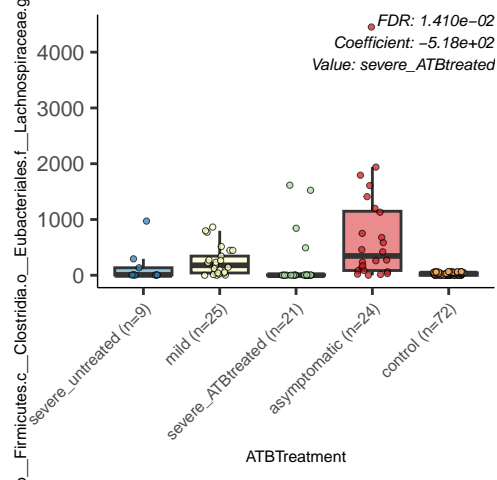


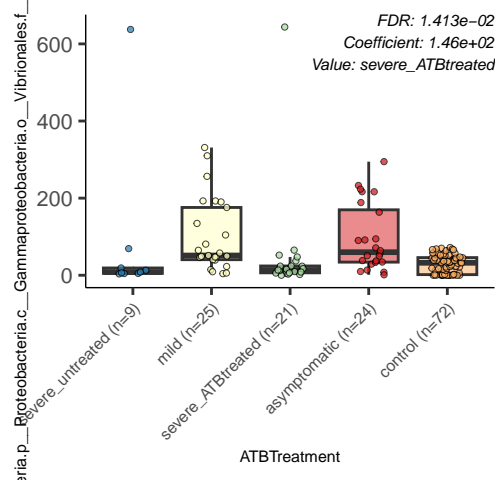


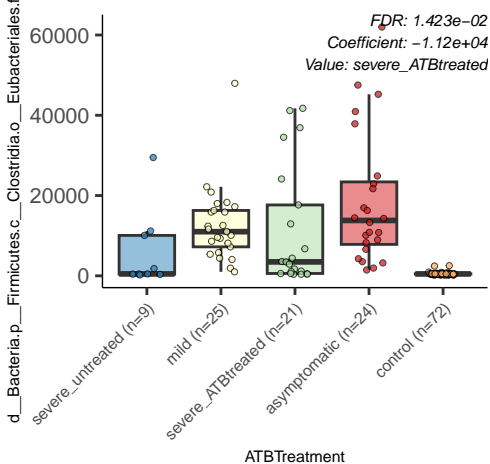


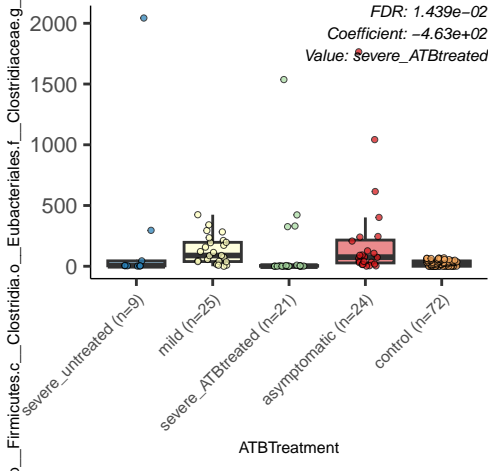


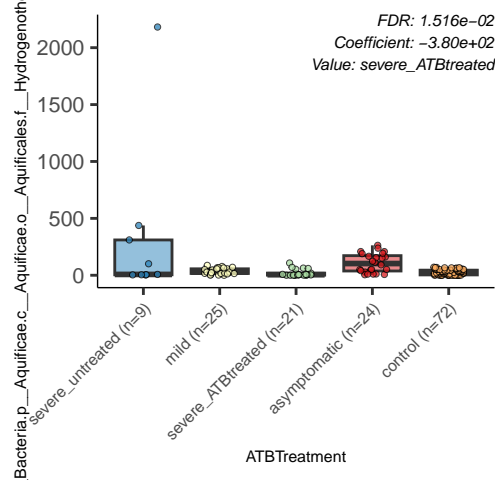


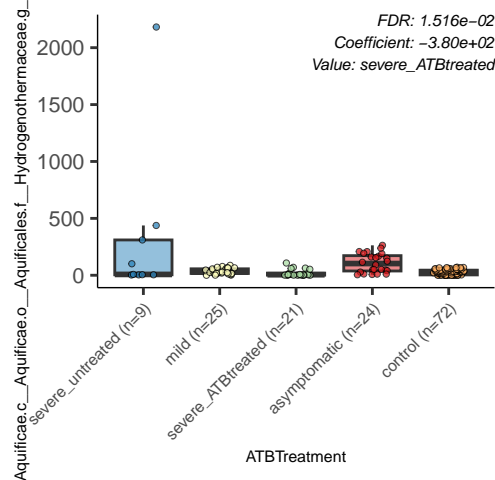


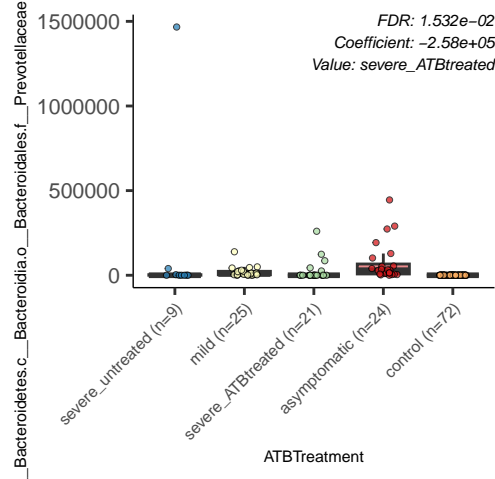


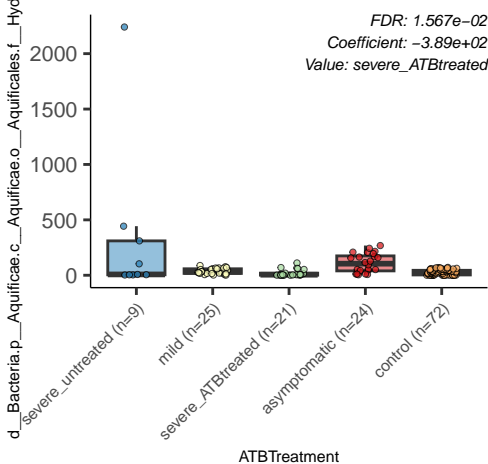


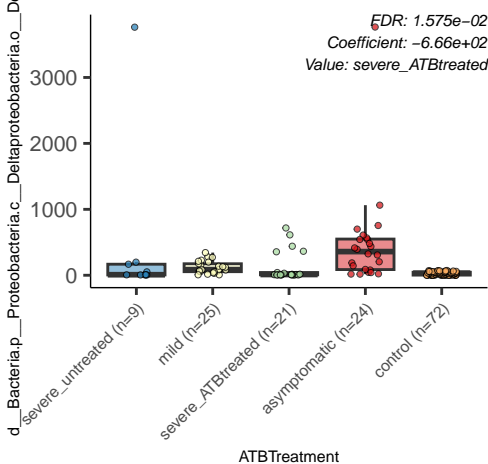


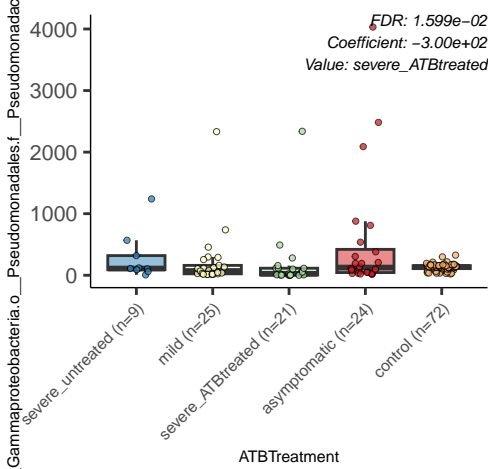


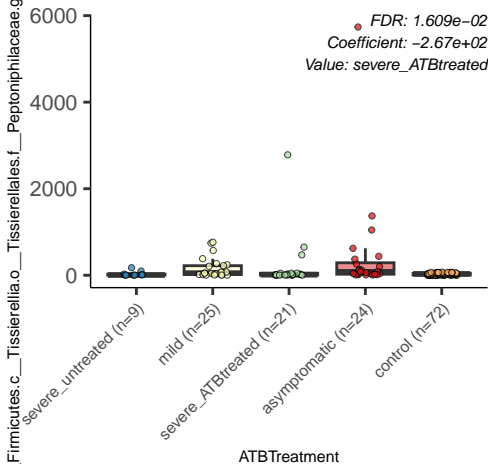


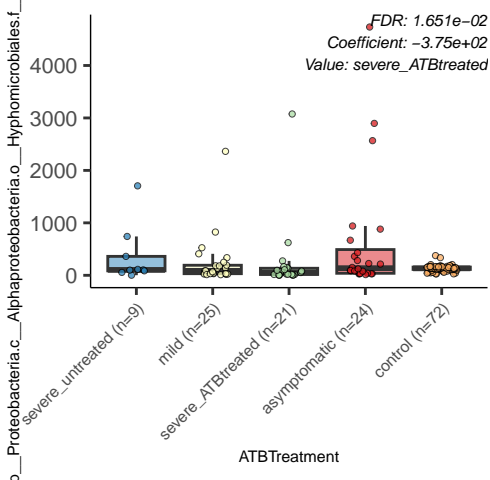


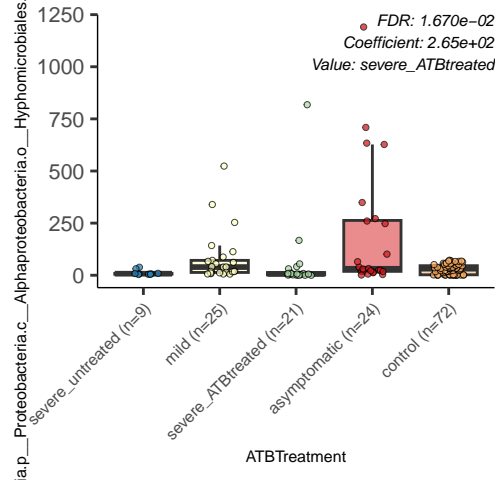


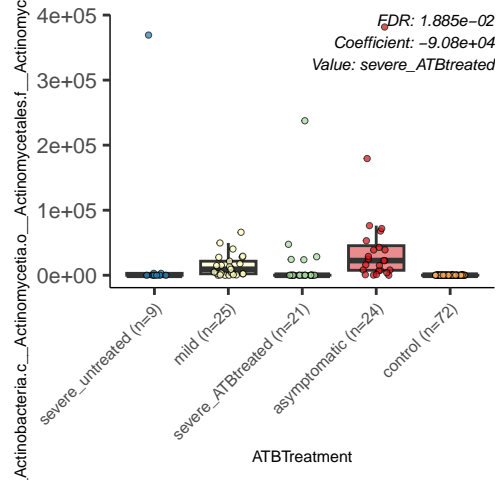




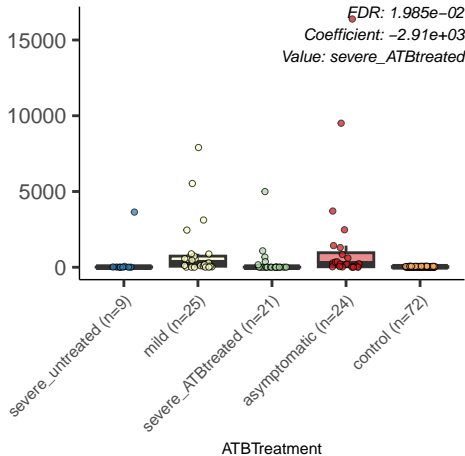


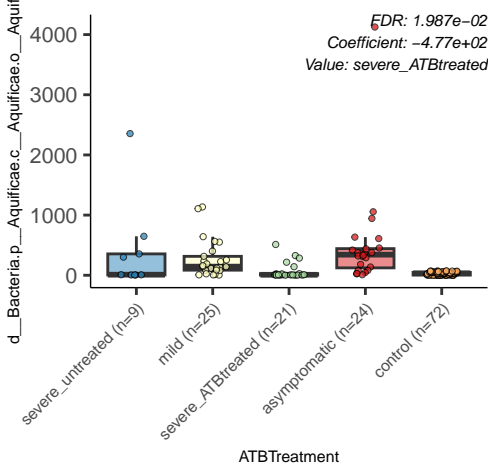


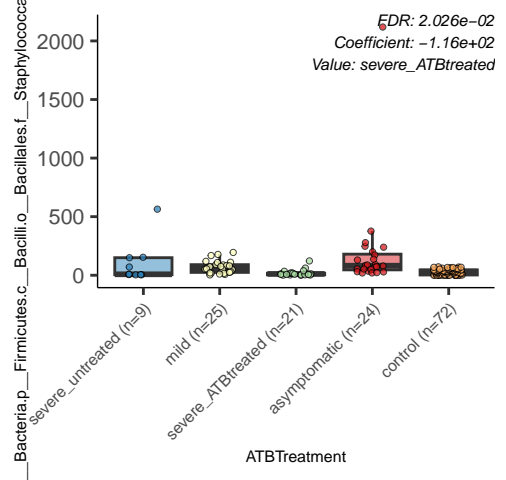


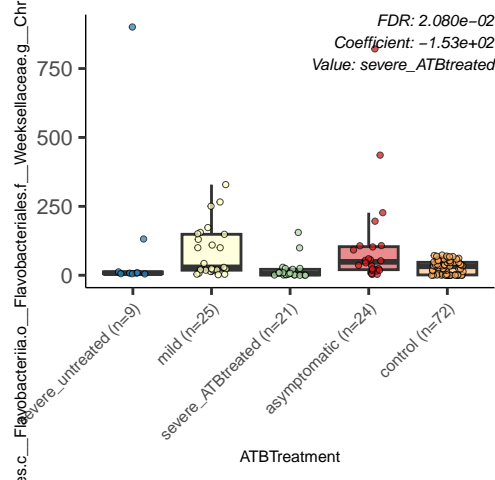


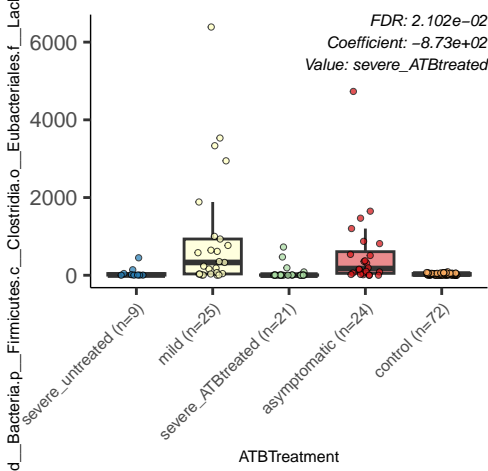
Bacteroidetes.c__Bacteroidia.o__Bacteroidales.f__Bacteroidaceae.

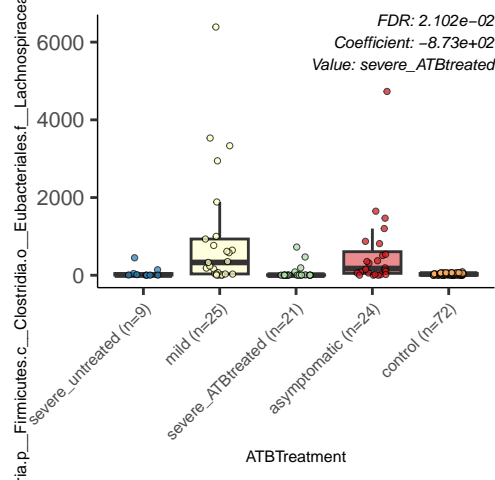






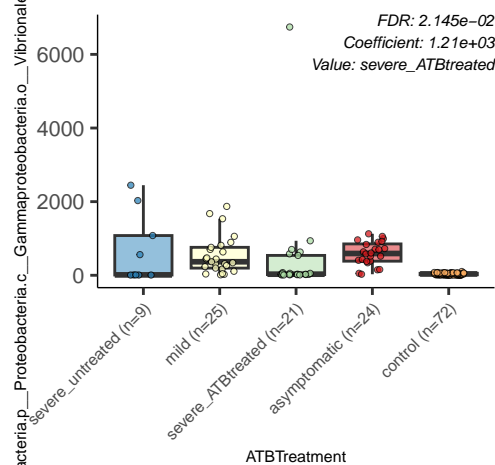


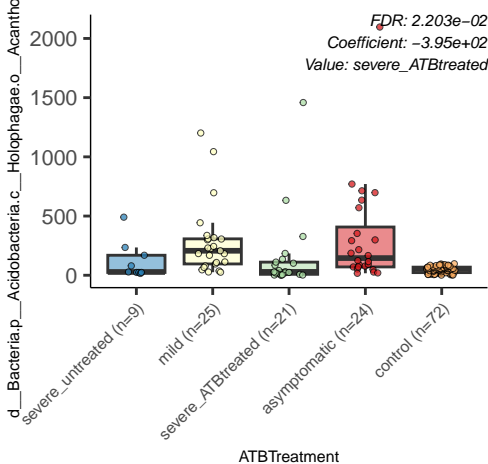


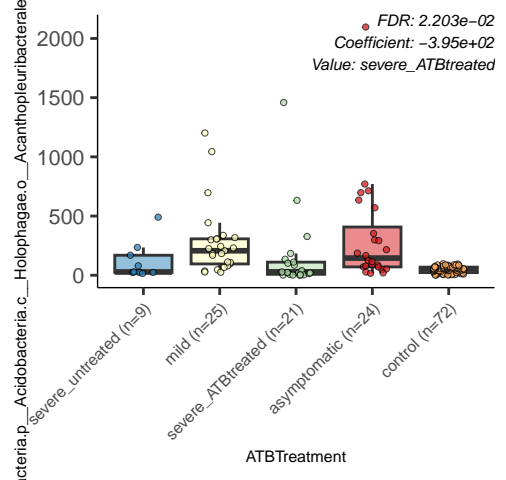


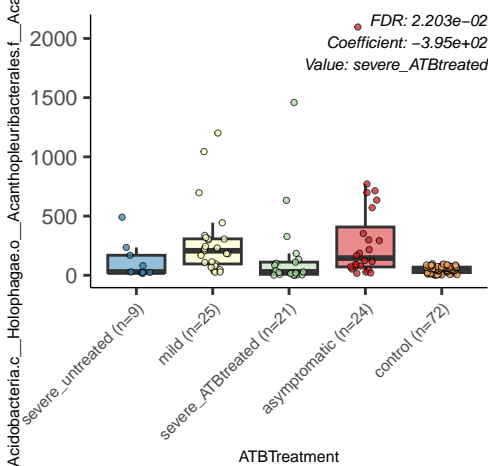
Bacteria.p__Proteobacteria.c__Gammaproteobacteria.o__Vibrionaceae

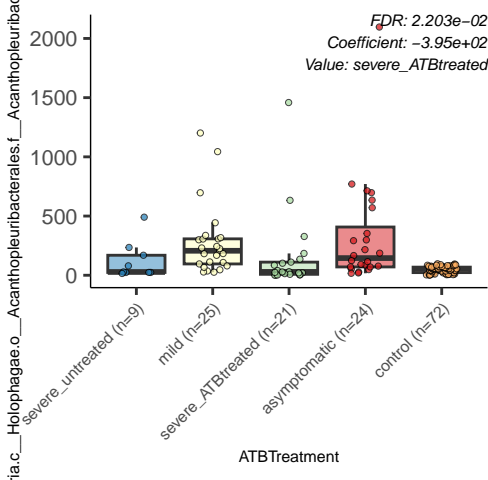
FDR: 2.145×10^{-2}
Coefficient: 1.21×10^3
Value: severe_ATBtreated

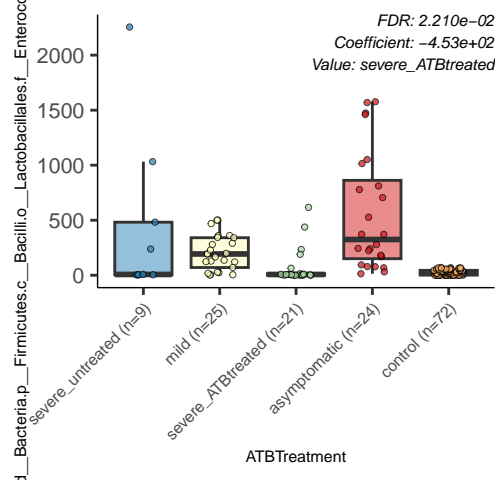


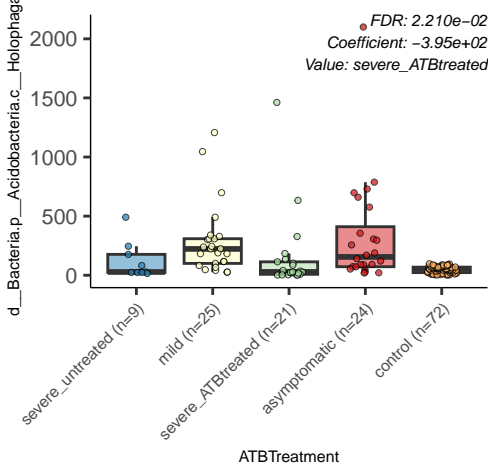


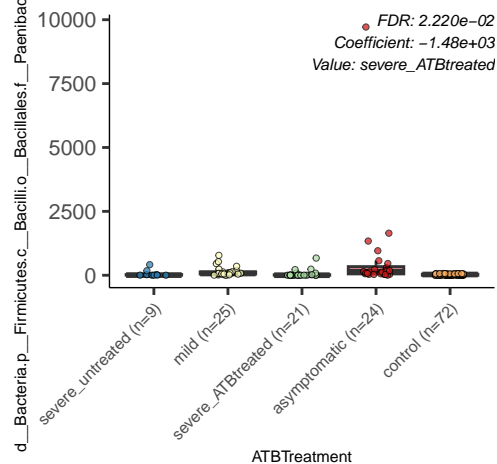


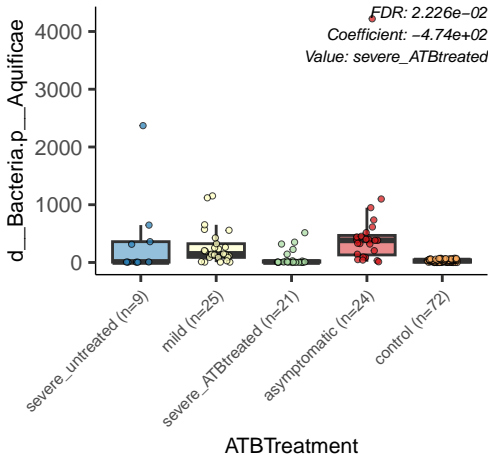


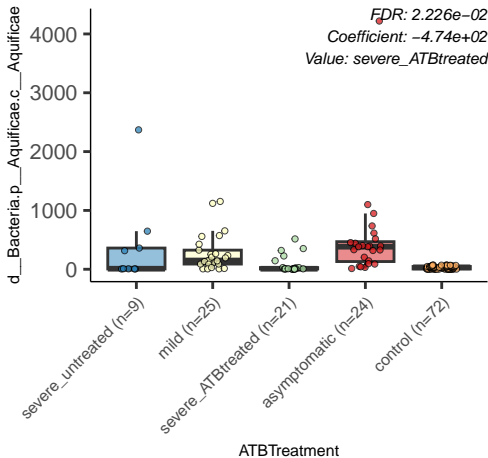


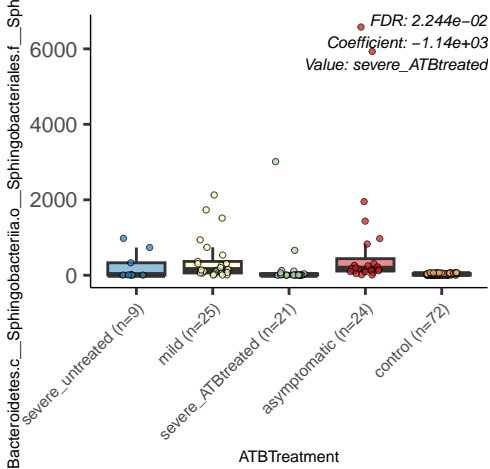


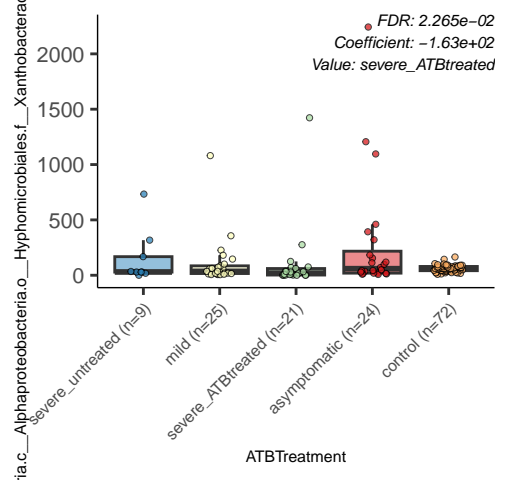


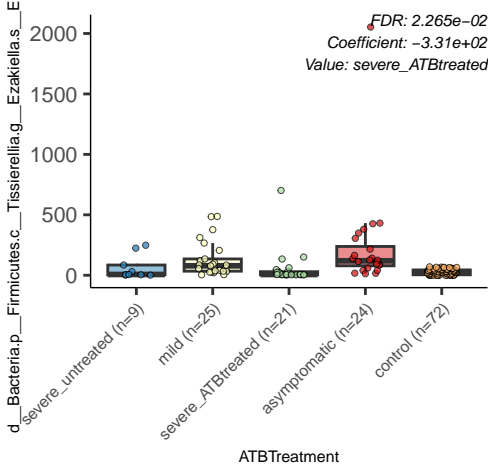


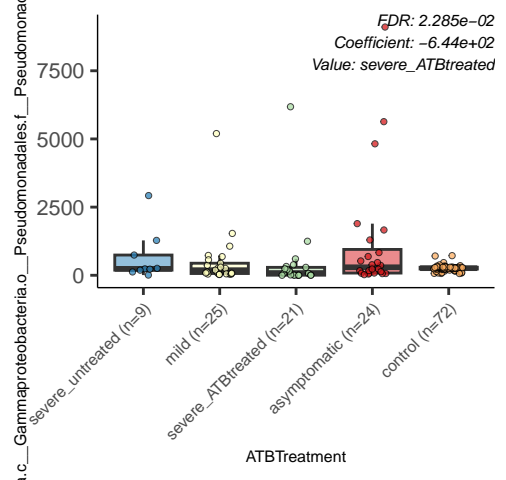


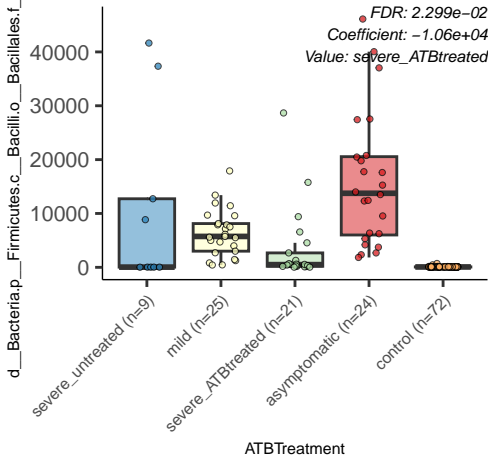






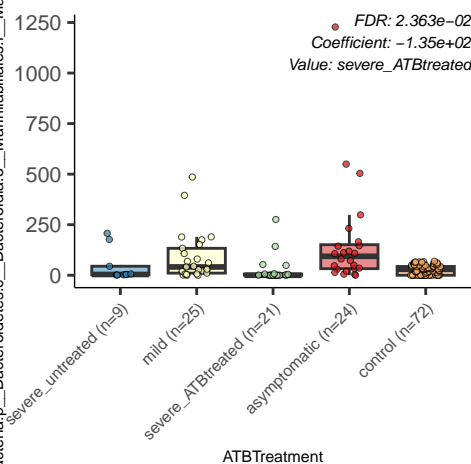




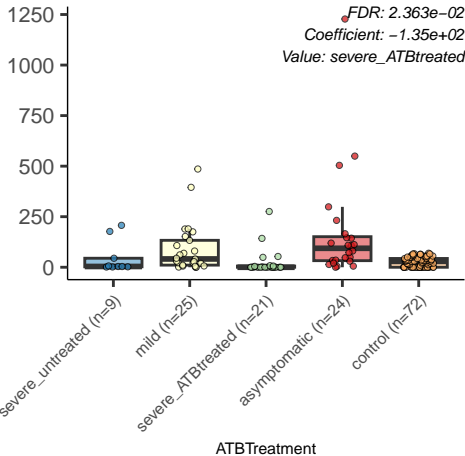


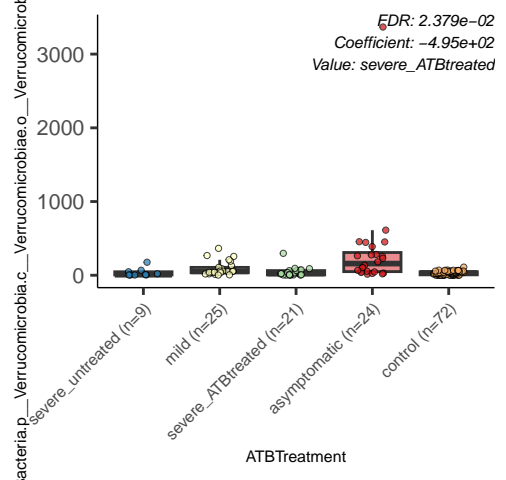
Bacteria.p_Bacteroidetes.c_Bacteroidia.o_Marinilabiales.f_Ma

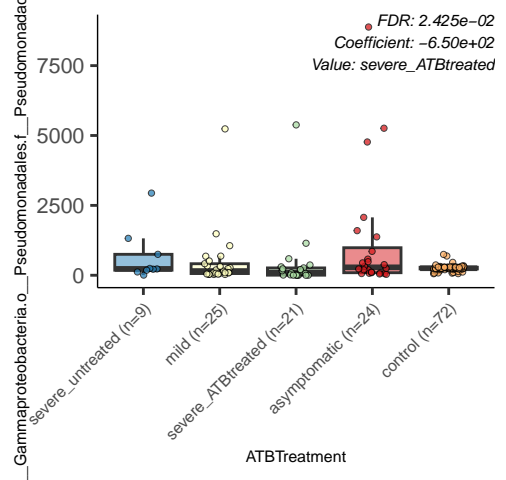
FDR: 2.363×10^{-2}
Coefficient: -1.35×10^2
Value: severe_ATBtreated

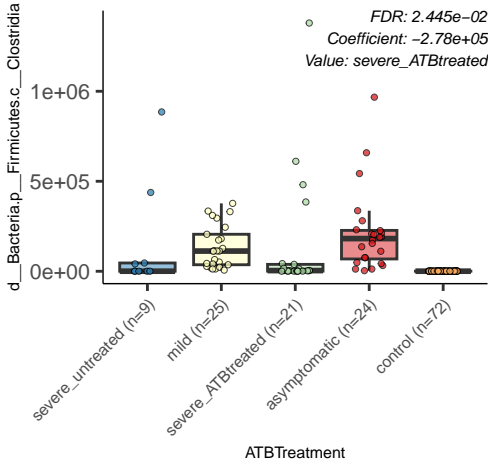


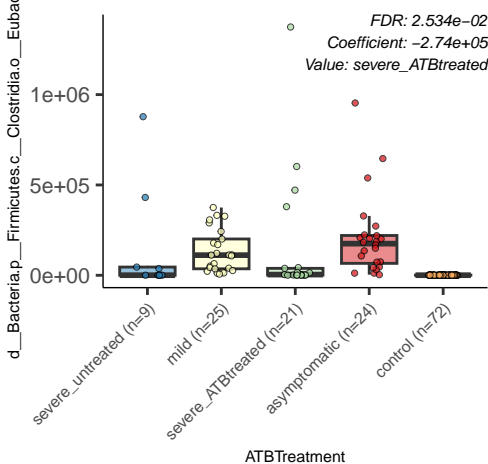
Bacteroidetes.c__Bacteroidia.o__Marinilabiales.f__Marinilabiales

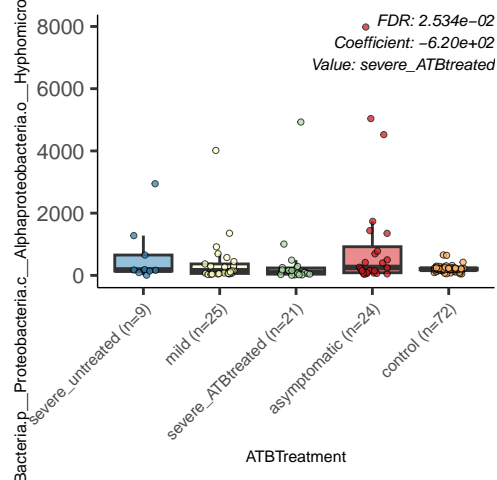


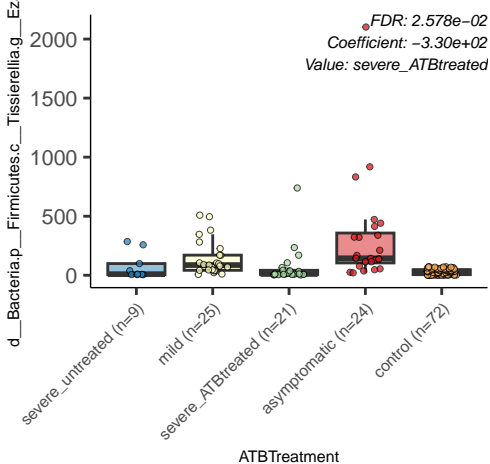


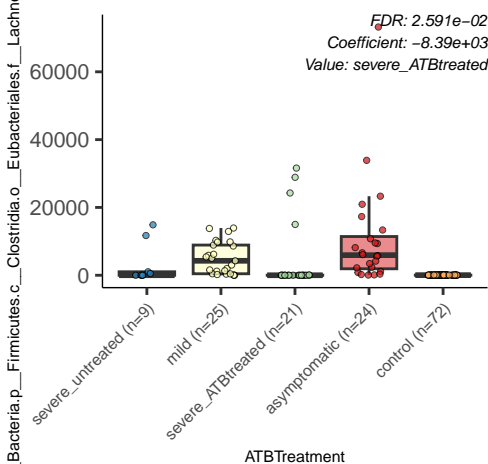


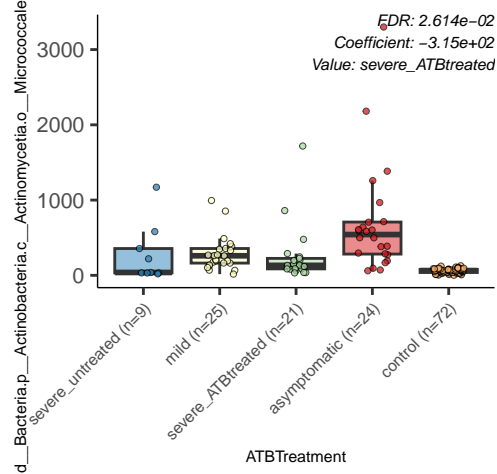


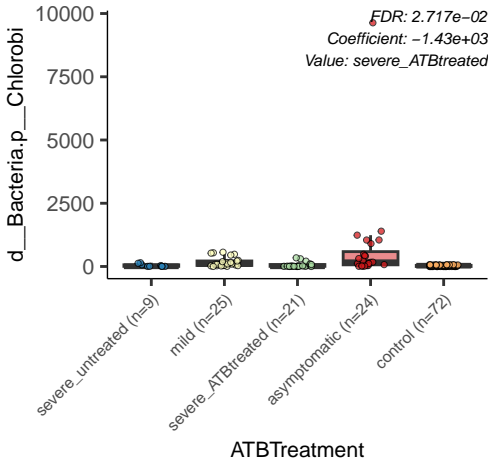


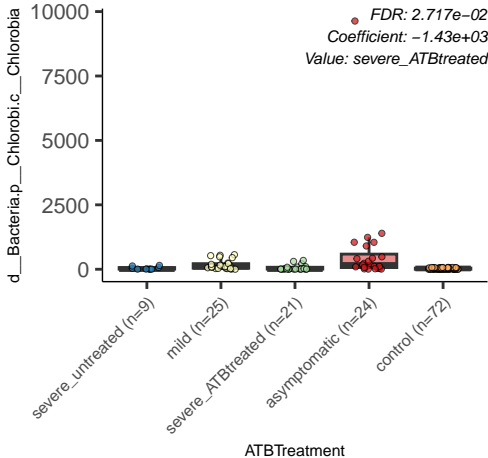


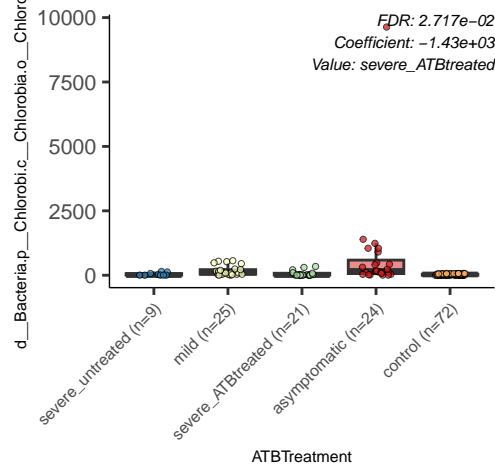


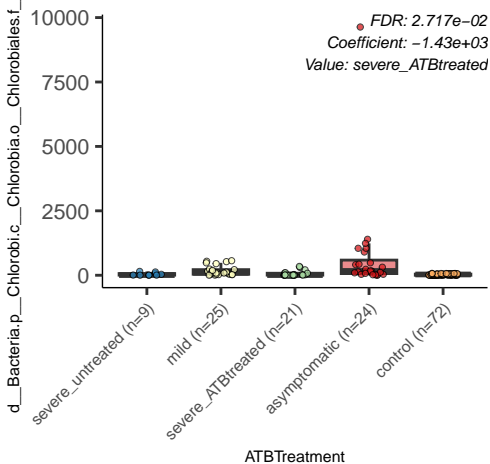


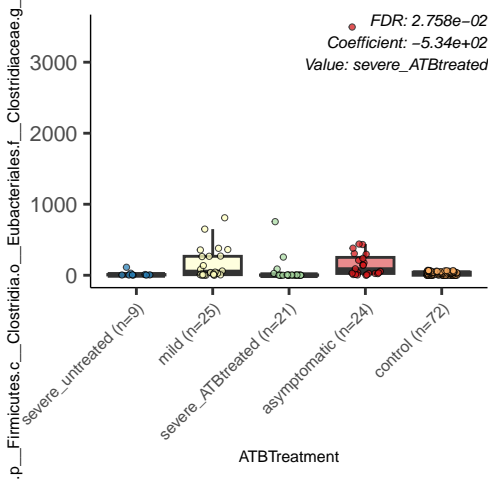


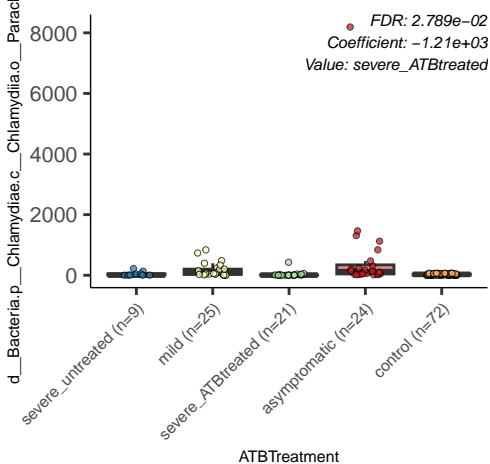


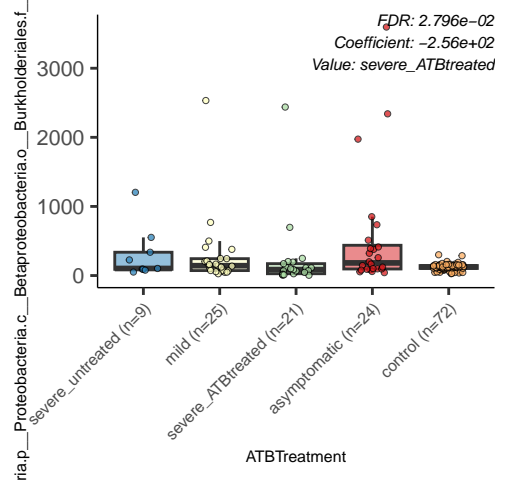


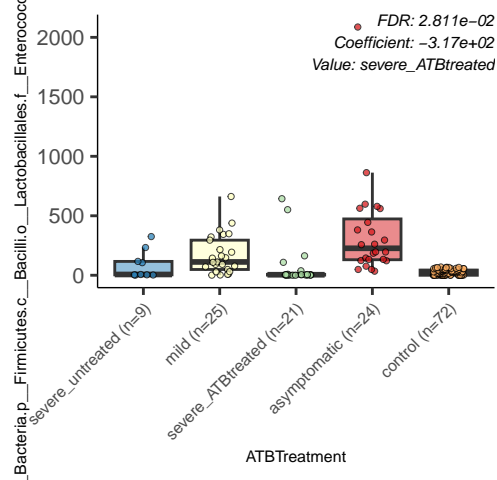


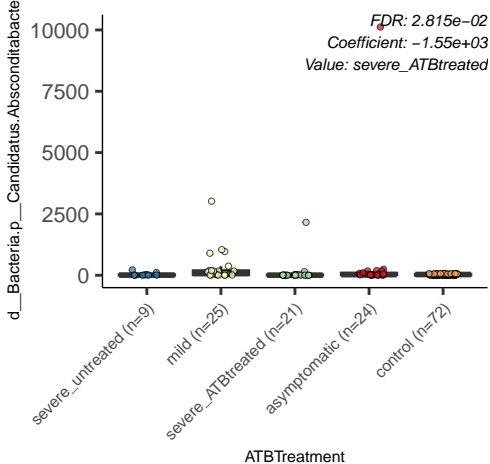


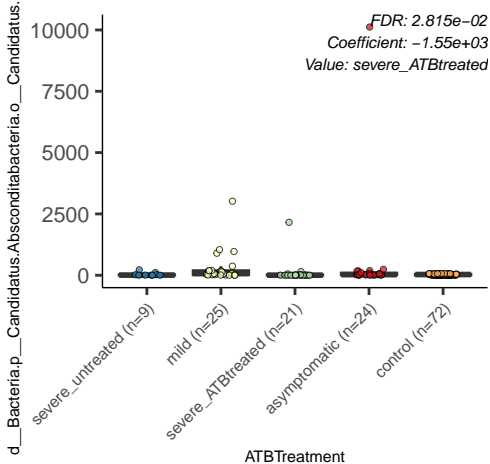


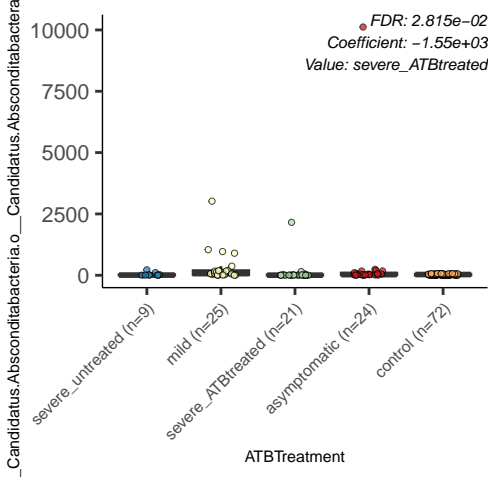


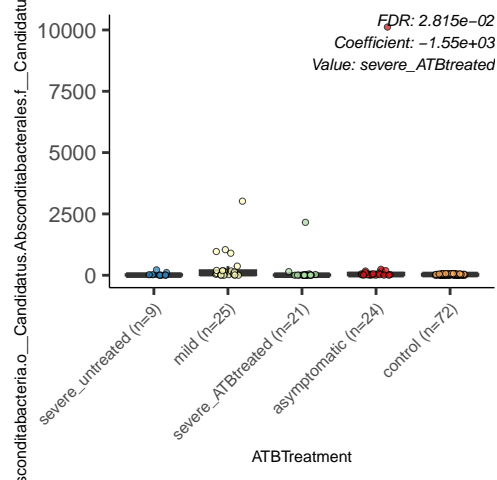


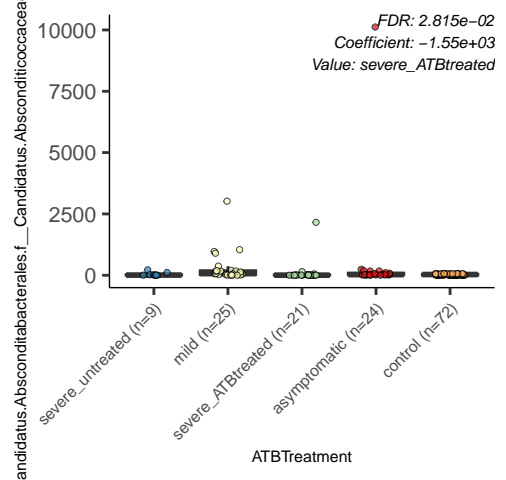


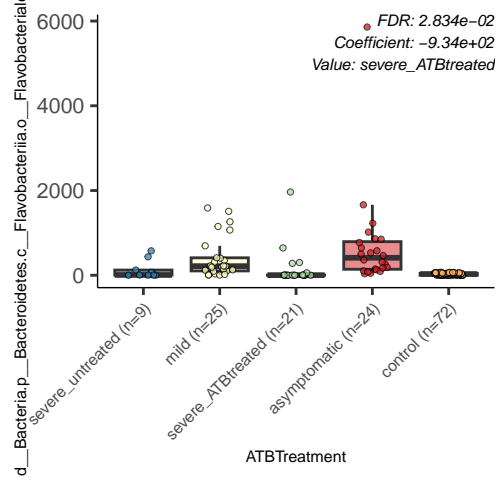


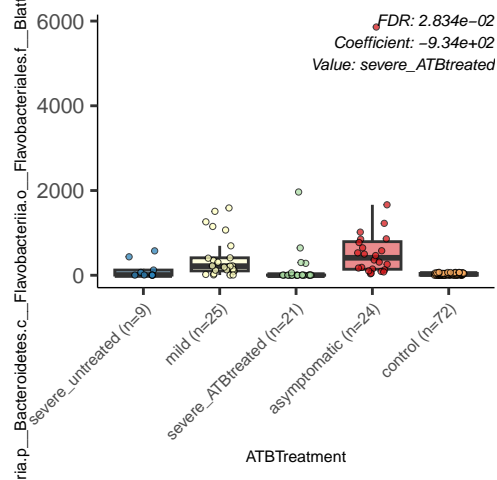


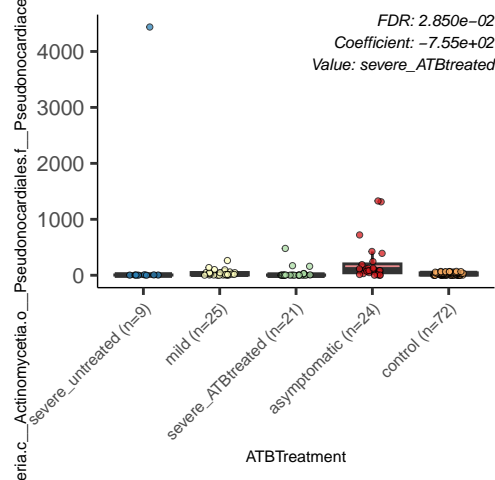


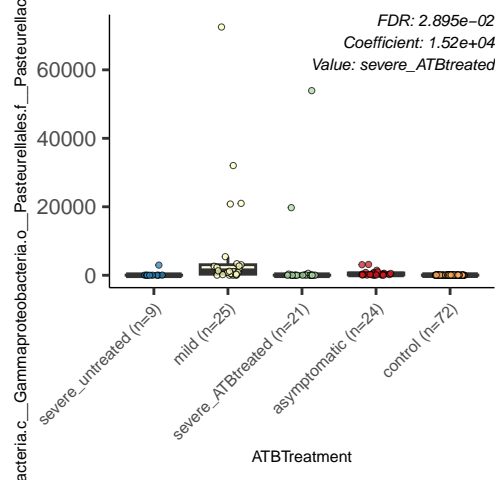


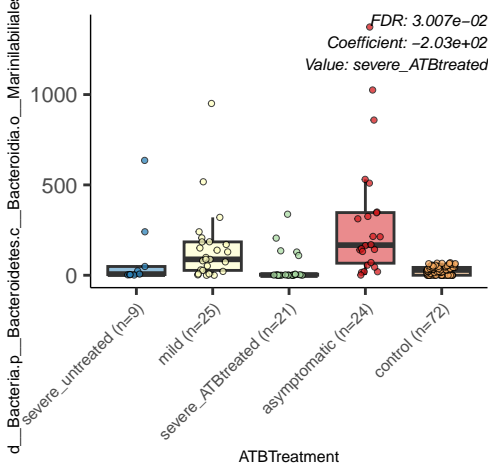


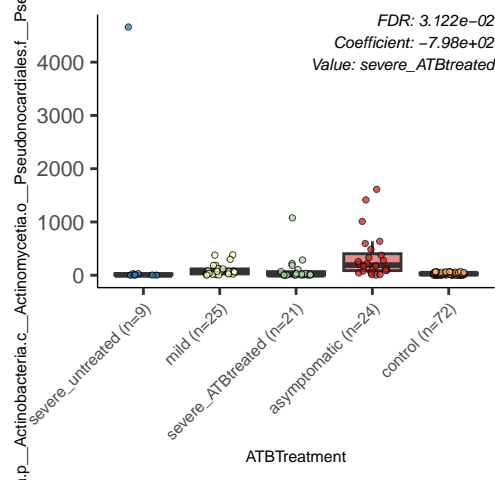


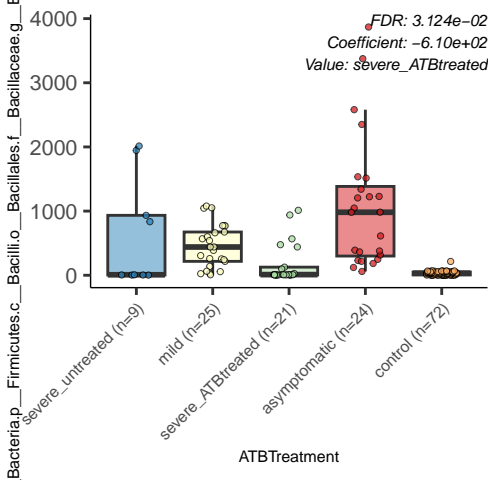






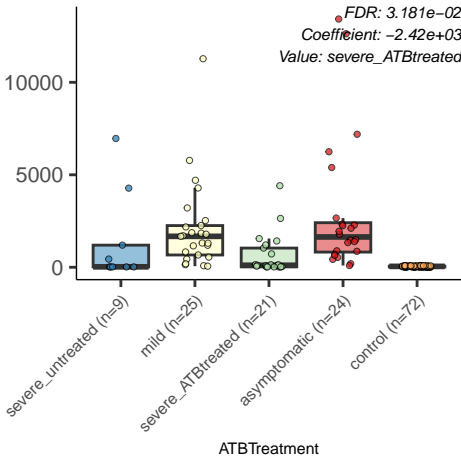


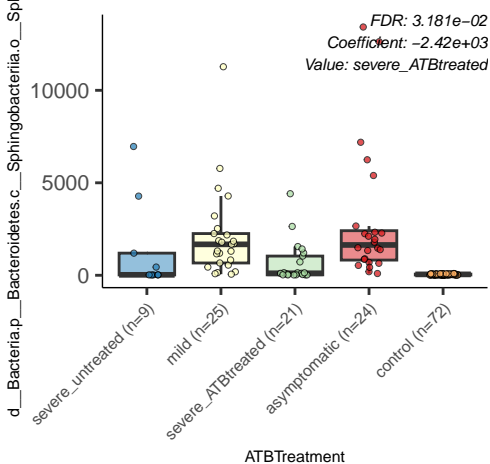




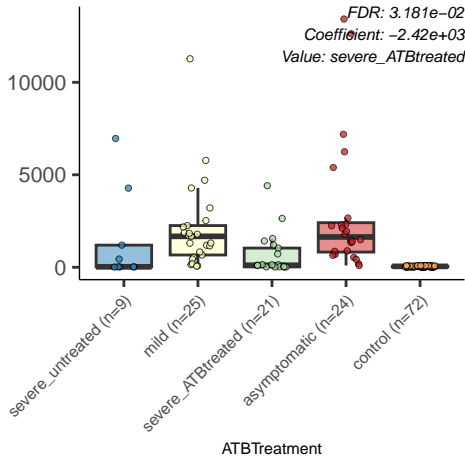
d__Bacteria.p__Bacteroidetes.c__Sphingobact

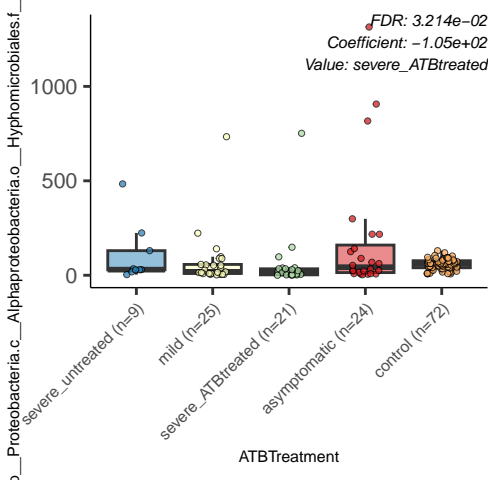
FDR: 3.181×10^{-2}
Coefficient: -2.42×10^3
Value: severe_ATBtreated





Bacteria.p_Bacteroidetes.c_Sphingobacteriia.o_Sphingobacteriia





ria.c__Alphaproteobacteria.o__Hyphomicrobiales.f__Xanthobacteria

FDR: 3.214×10^{-2}
Coefficient: -1.05×10^2
Value: severe_ATBtreated

1000
500
0

severe_unreated (n=9)

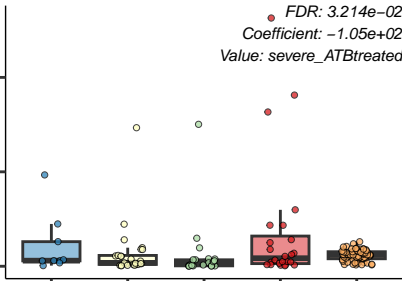
mild (n=25)

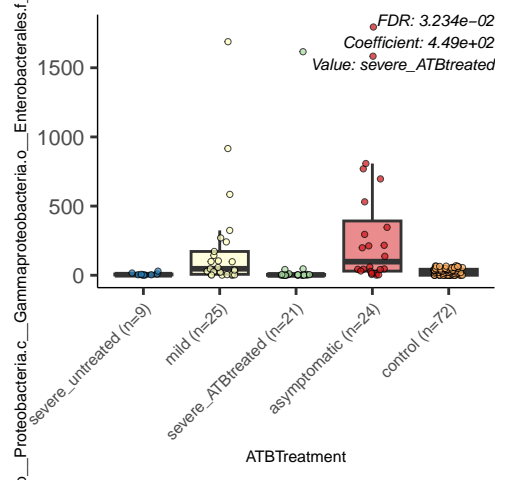
severe_ATBtreated (n=21)

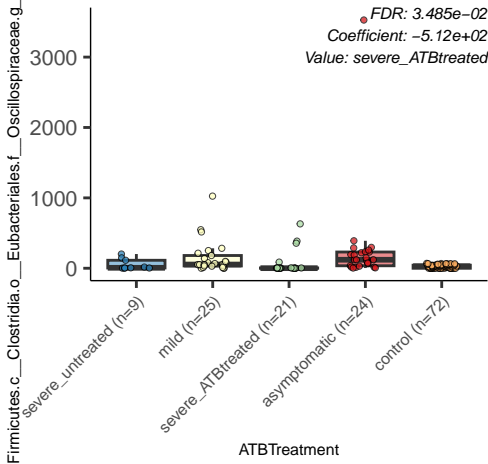
asymptomatic (n=24)

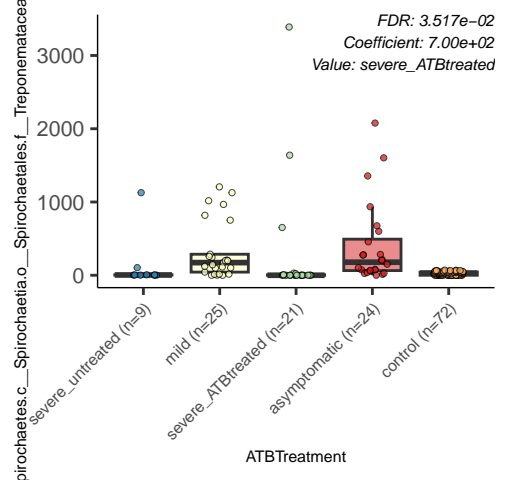
control (n=72)

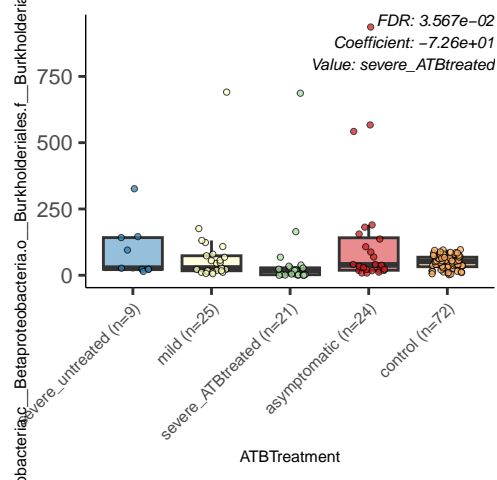
ATBTreatment

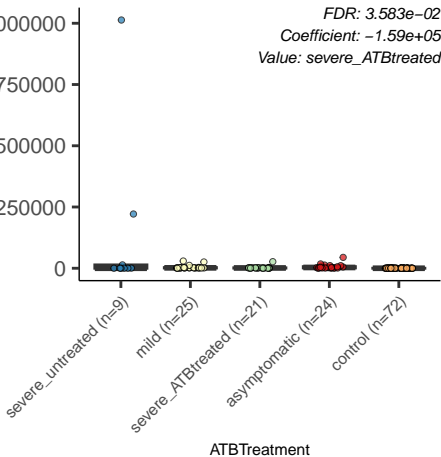


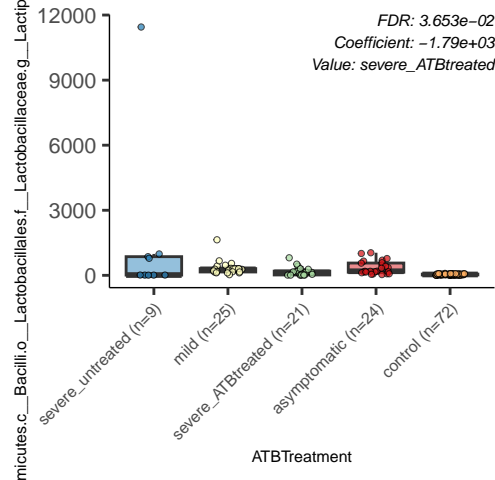


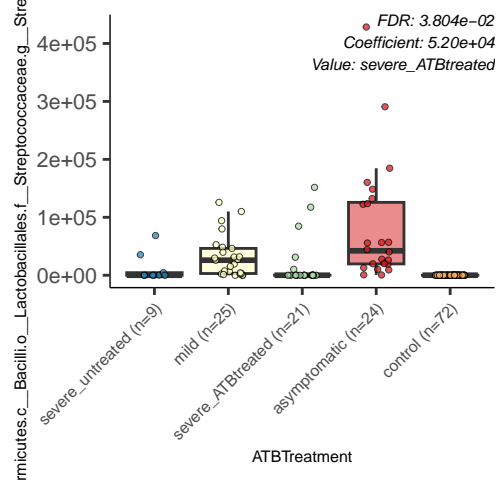




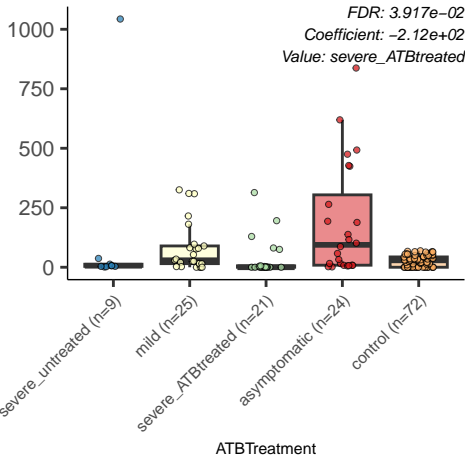


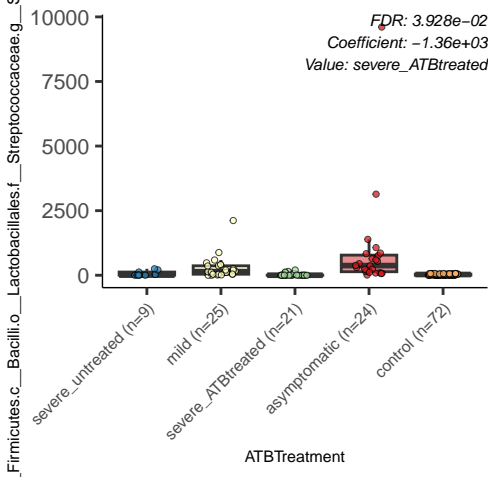


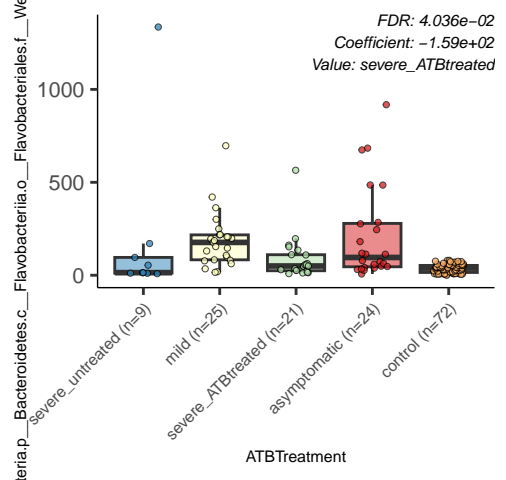


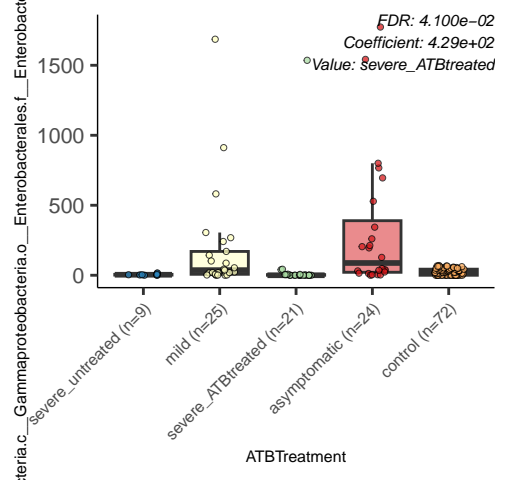


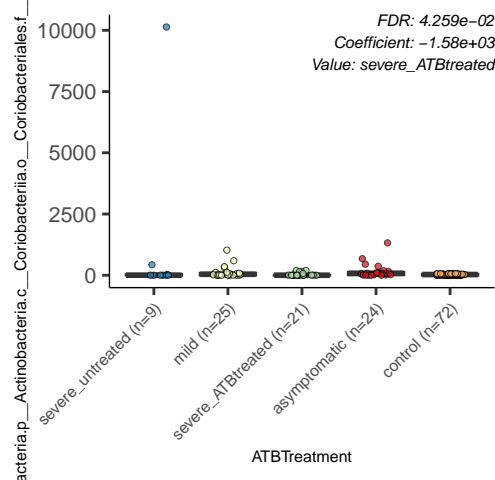
Bacteroidetes.c_Bacteroidia.o_Bacteroidales.f_Bacteroidaceae.



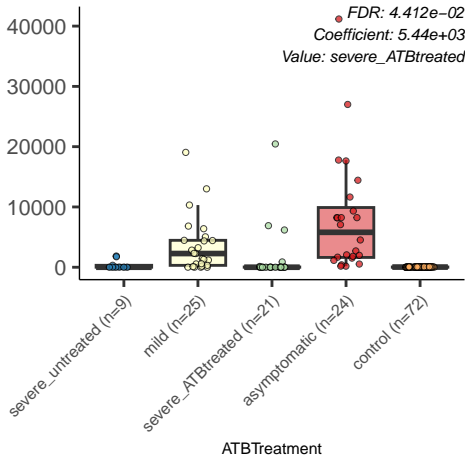




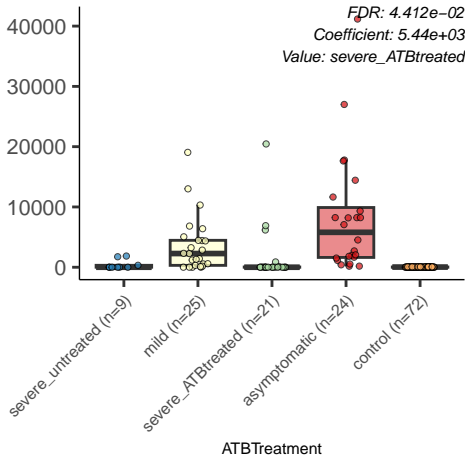


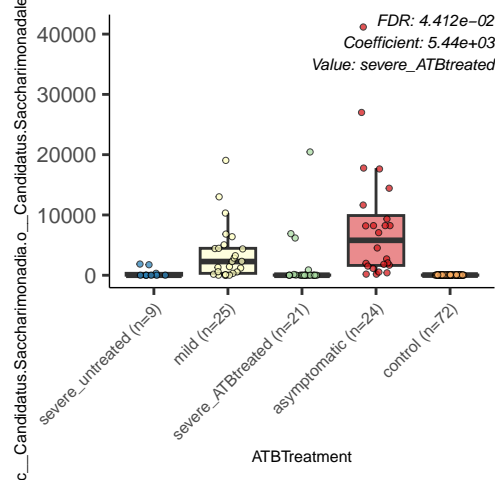


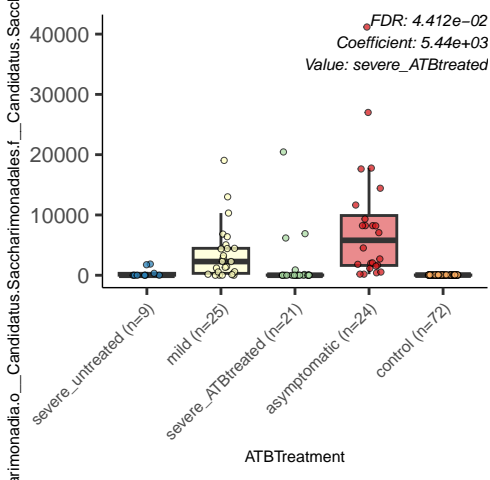
a.p__Candidatus.Saccharibacteria.c__Candidatus.Saccharimonadia

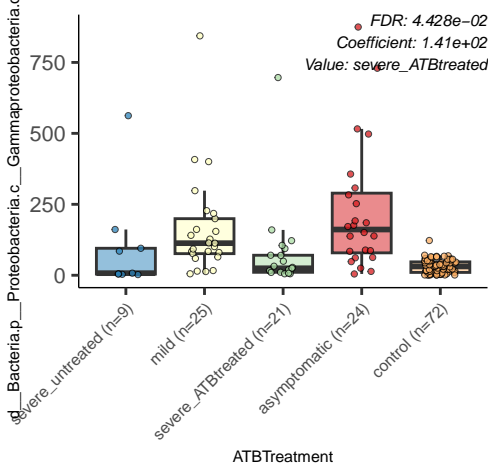


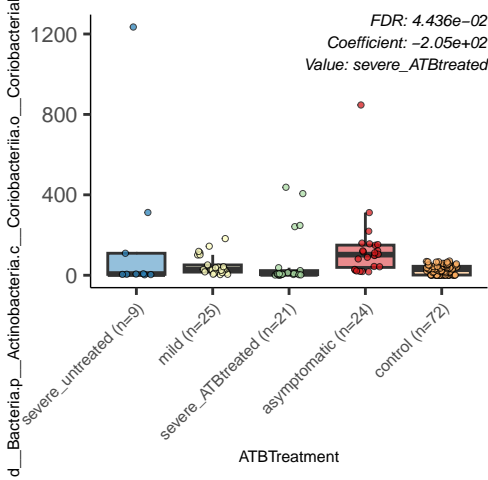
Saccharibacteria.c__Candidatus.Saccharimonadia.o__Candidatus.Sa

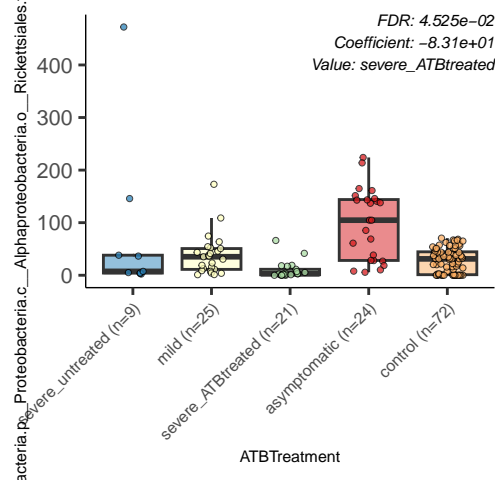










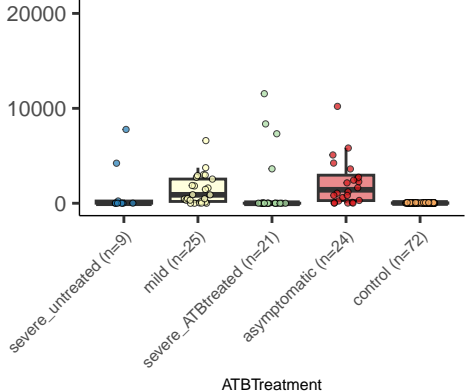


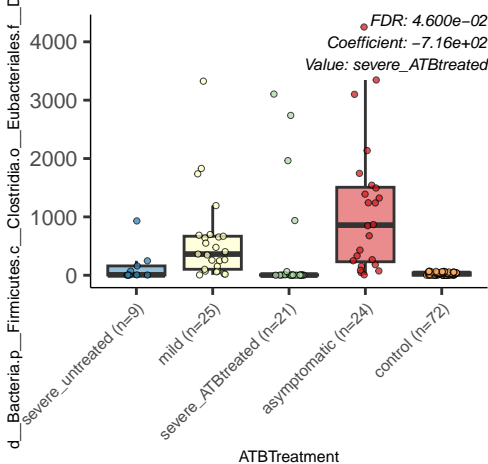
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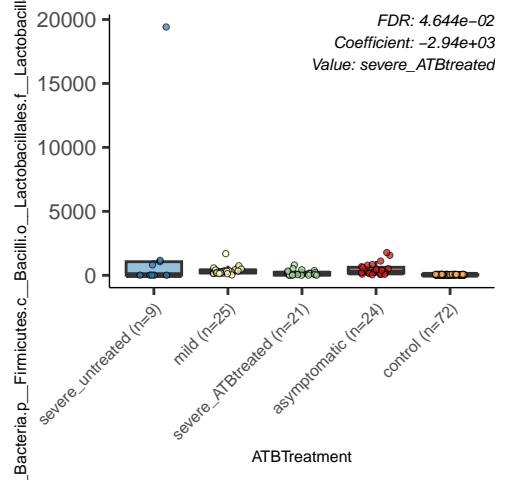
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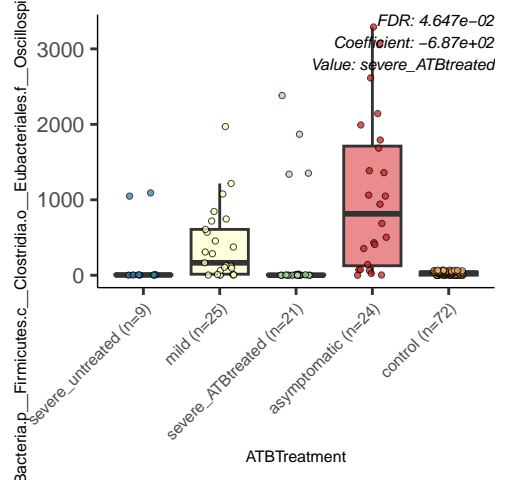
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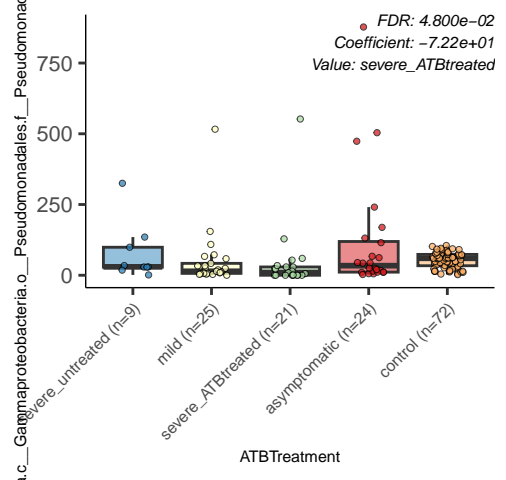
Value: severe_ATBtreated

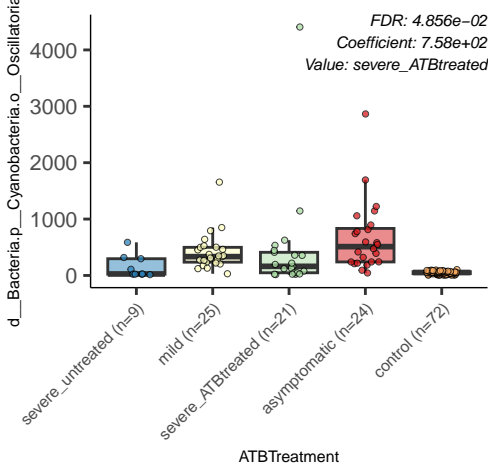




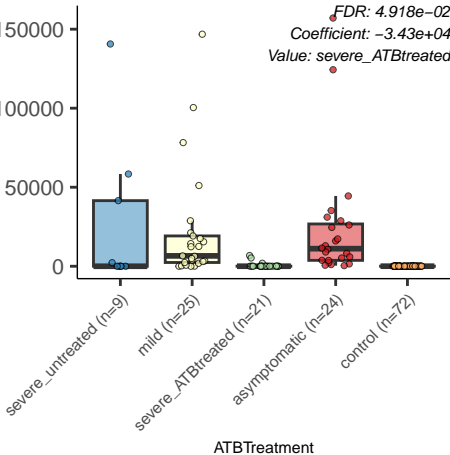


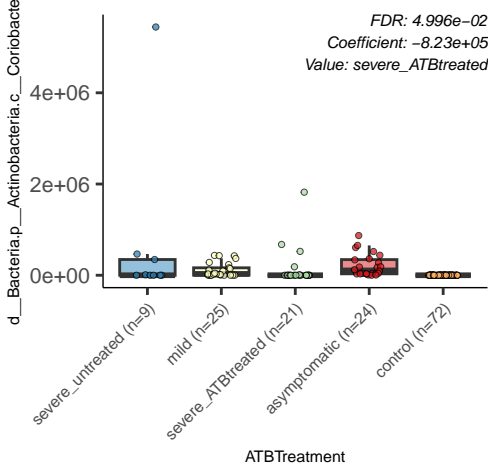


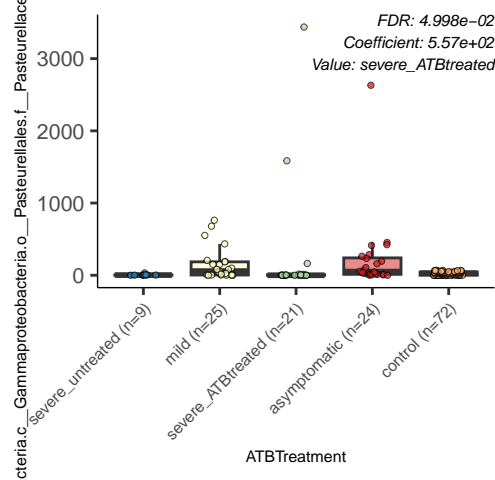


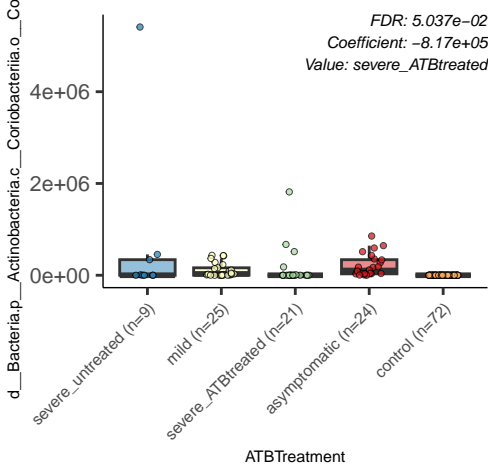


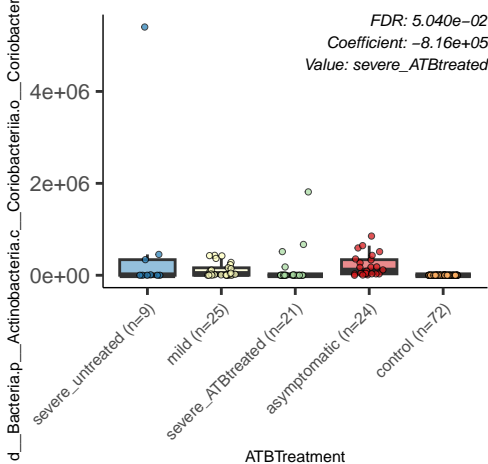
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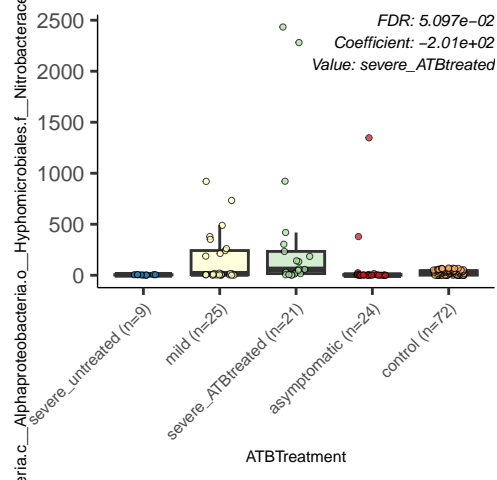


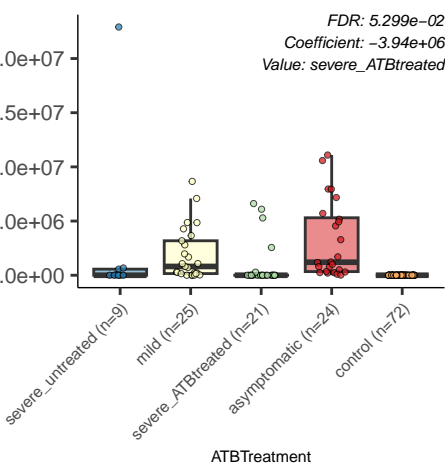


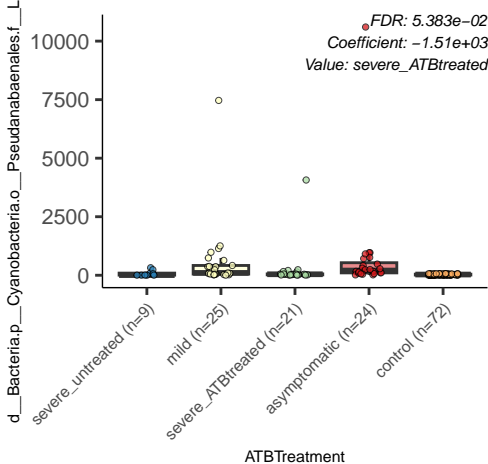


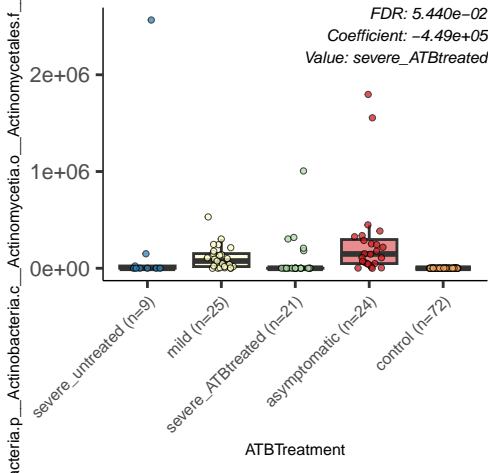




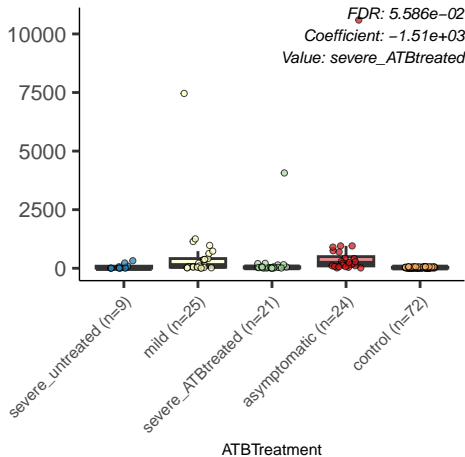


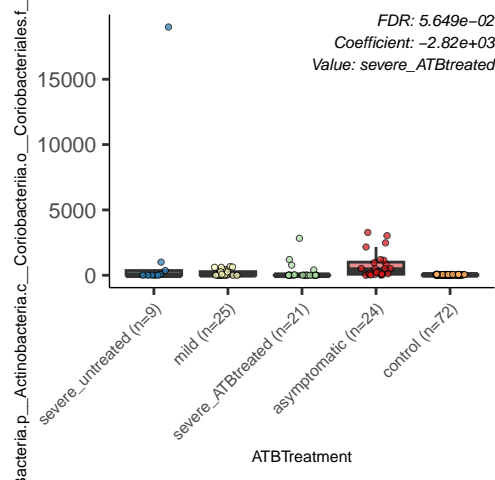




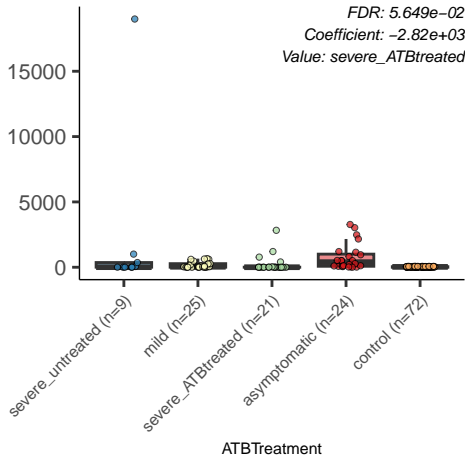


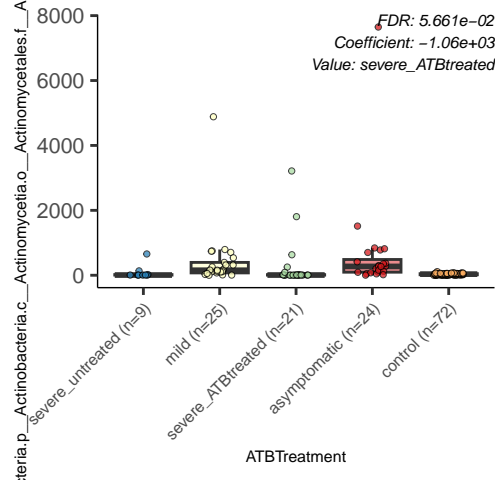
l_Bacteria.p__Cyanobacteria.o__Pseudanabaenales.f__Leptolyng

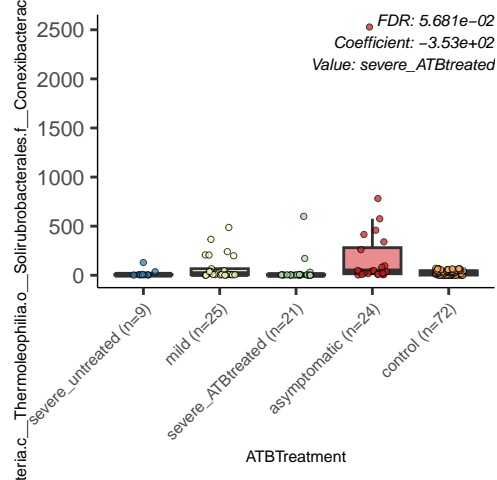


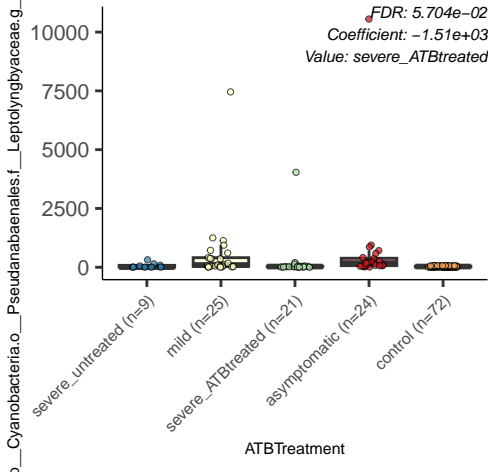


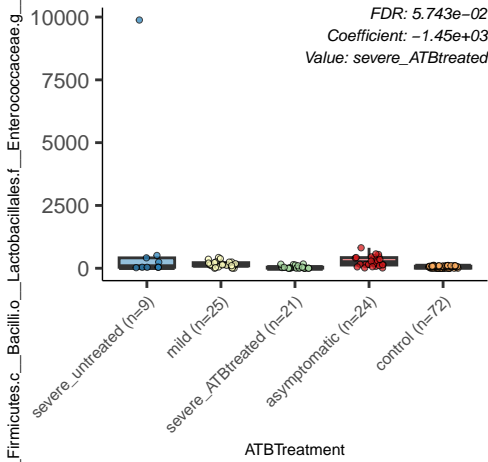
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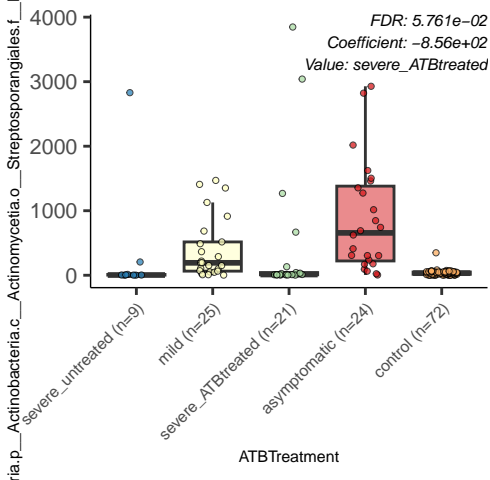


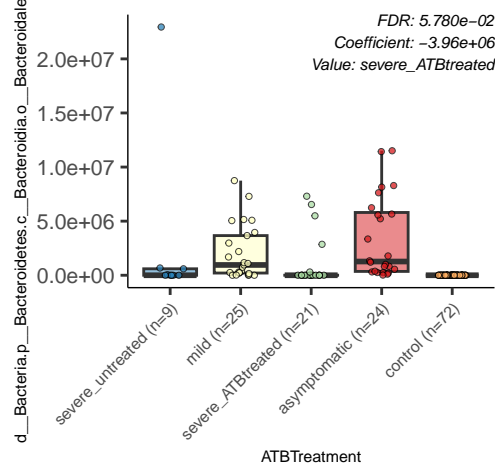


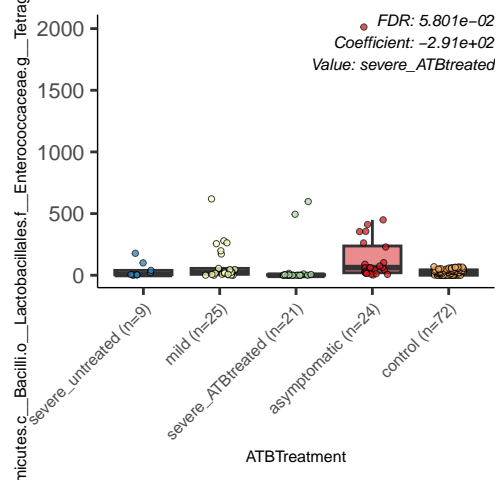


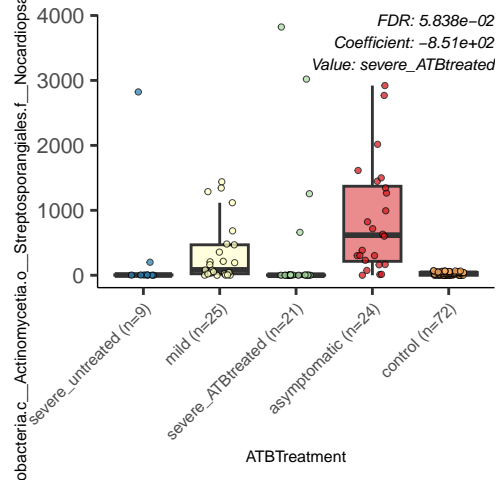


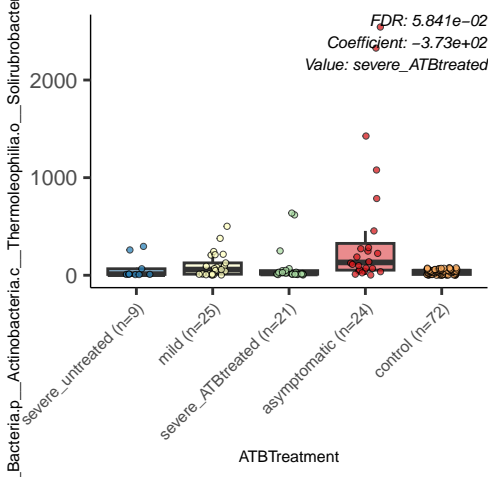


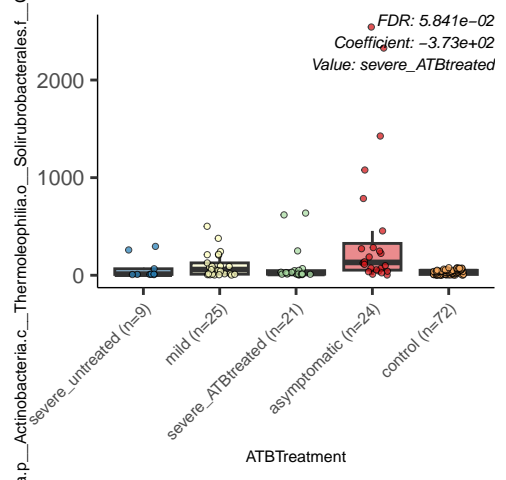




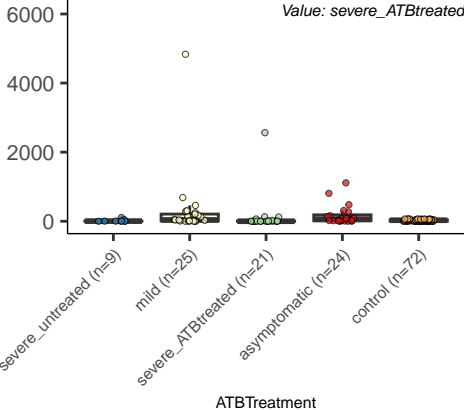
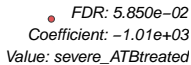


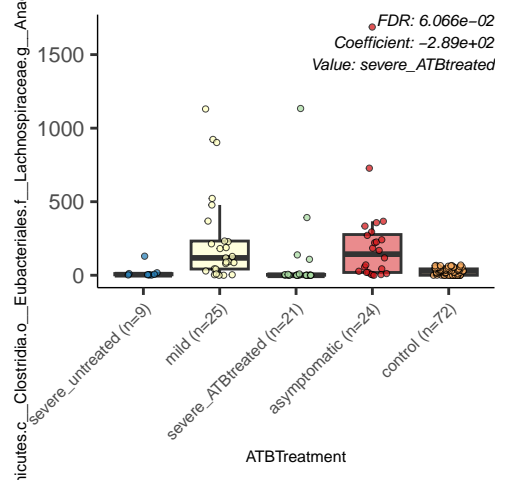


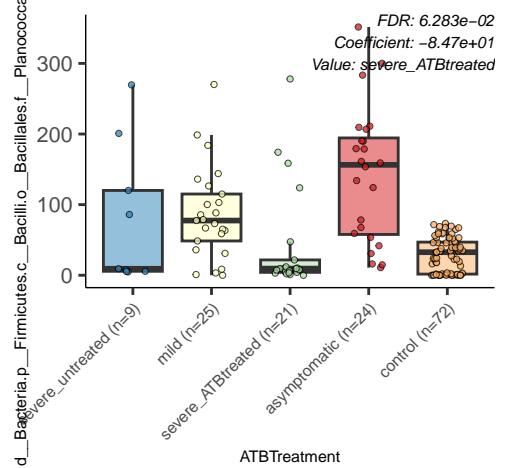


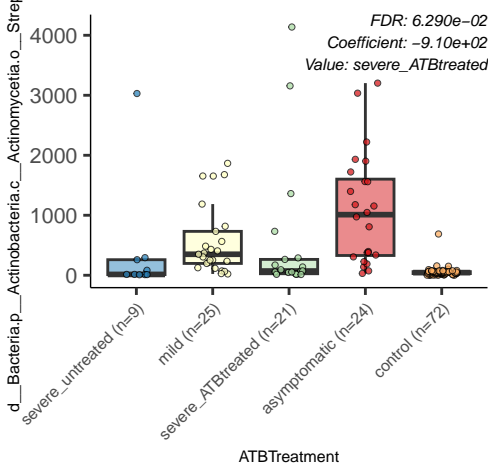


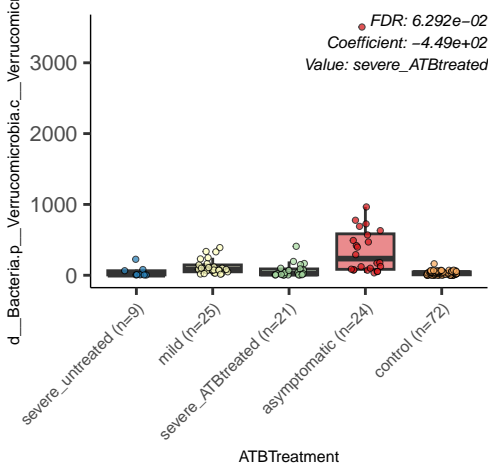
bacteria.c__Actinomycetia.o__Actinomycetales.f__Actinomycetaceae

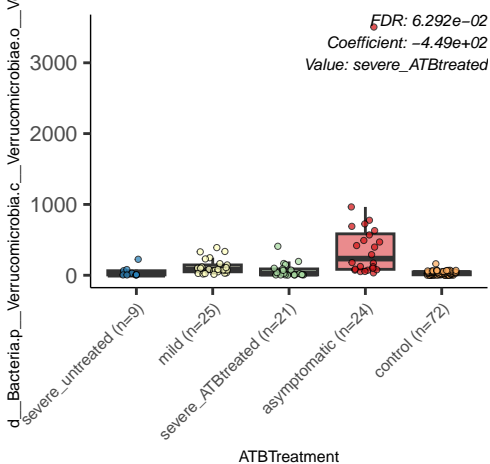


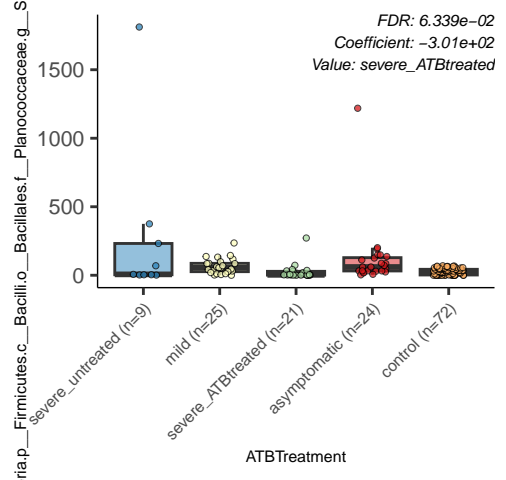


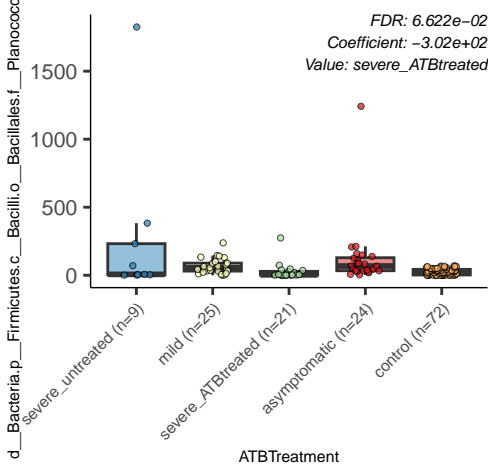


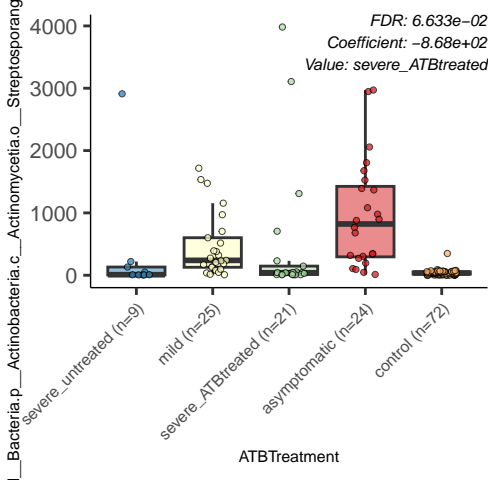


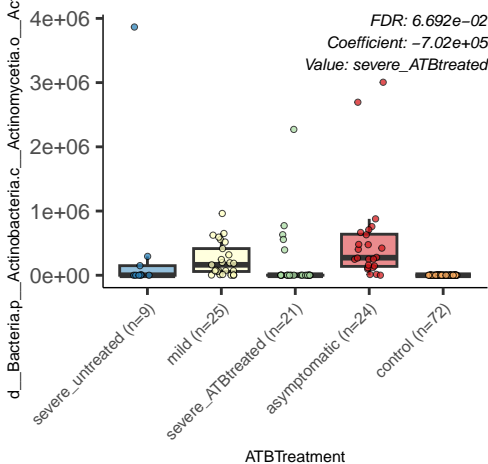


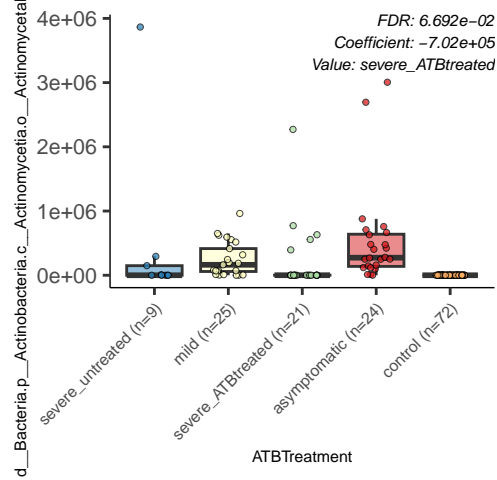


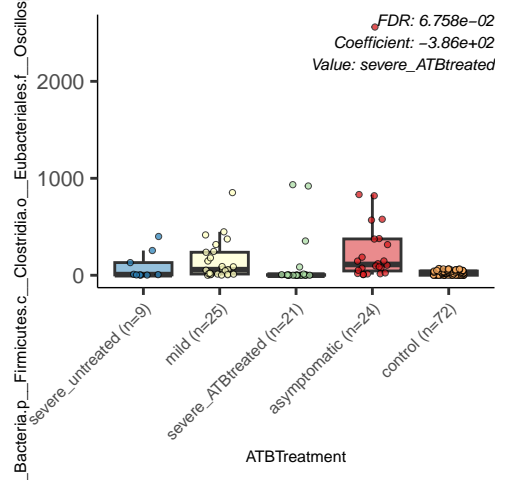


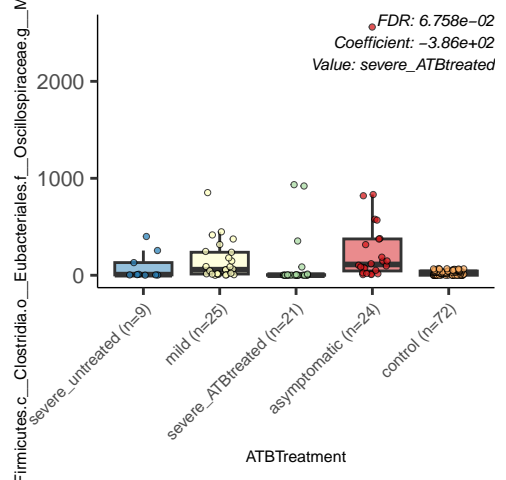


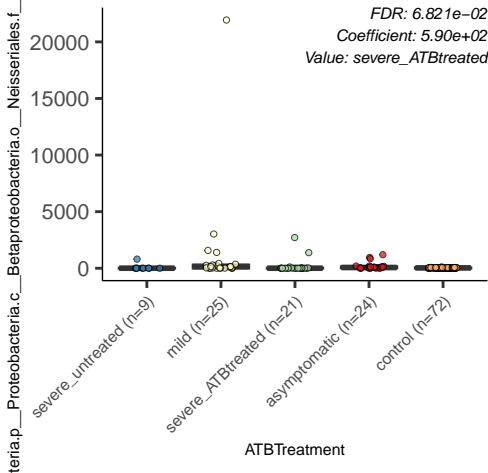


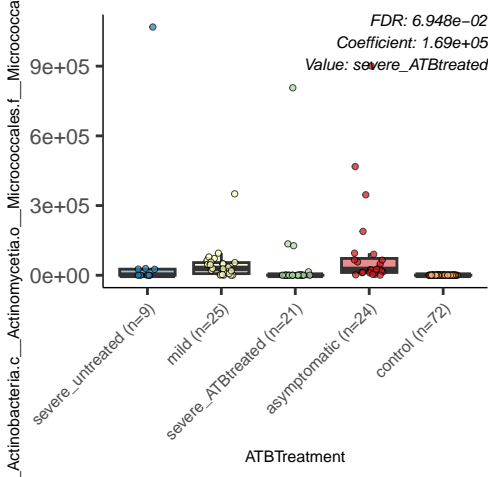


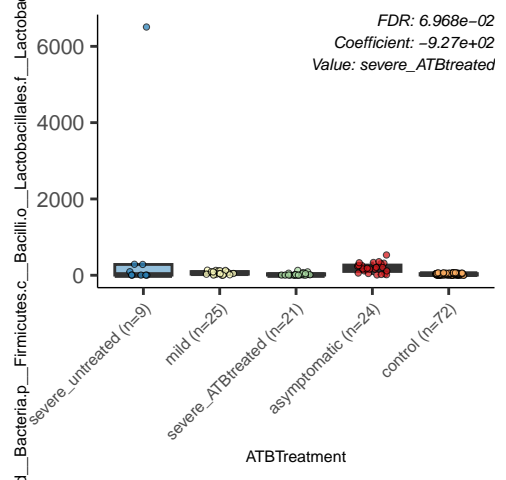


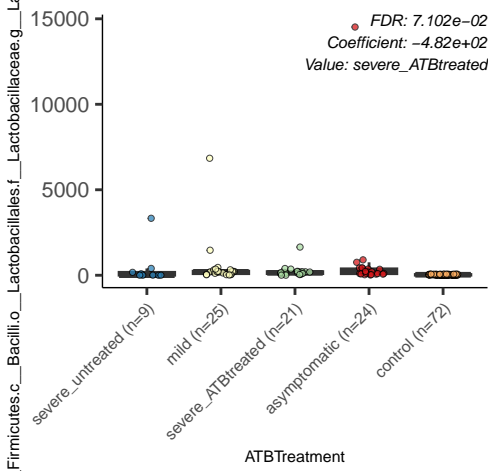


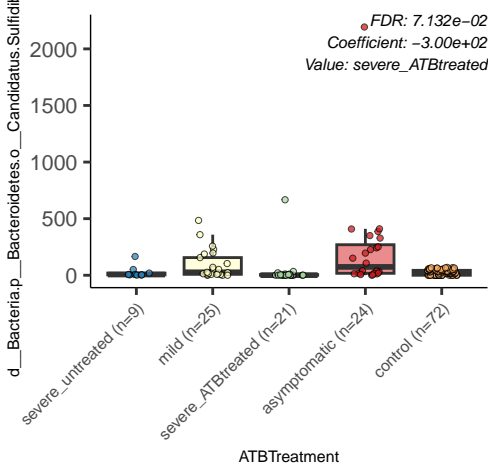


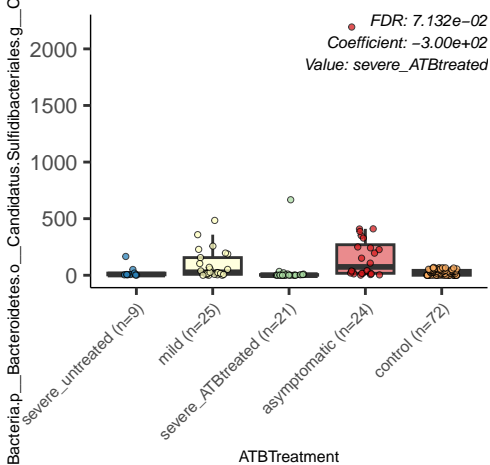


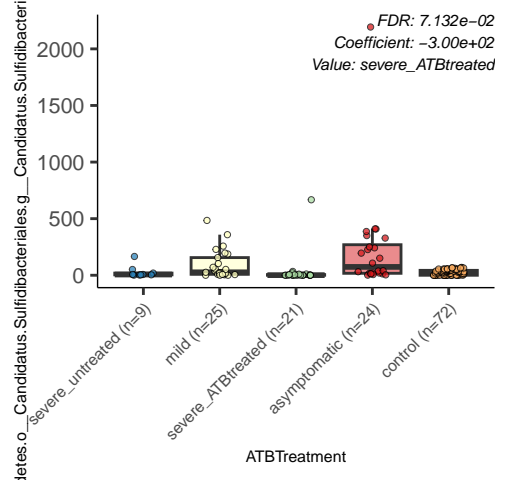


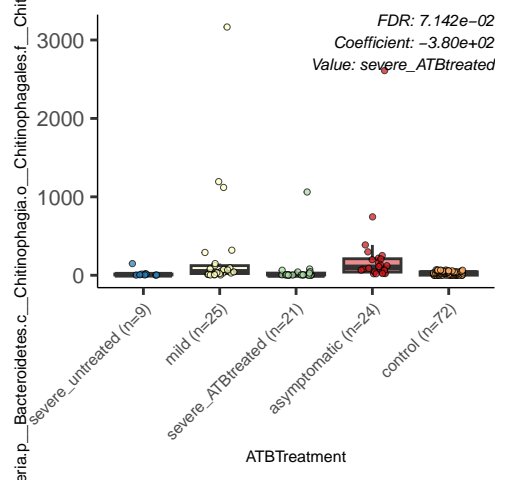


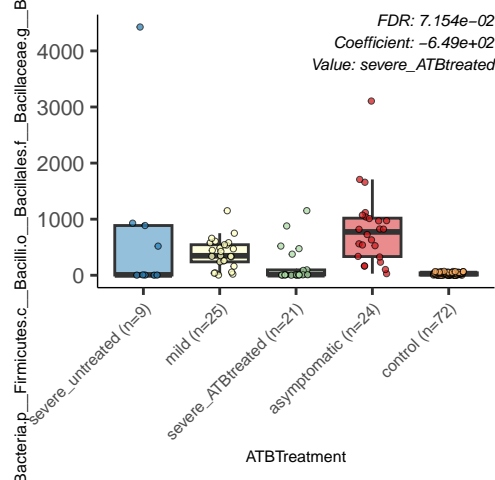


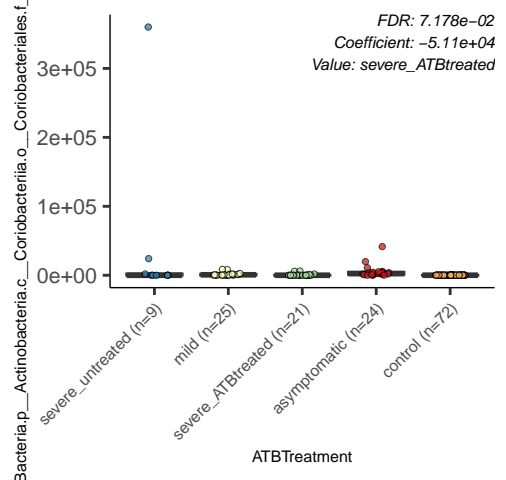


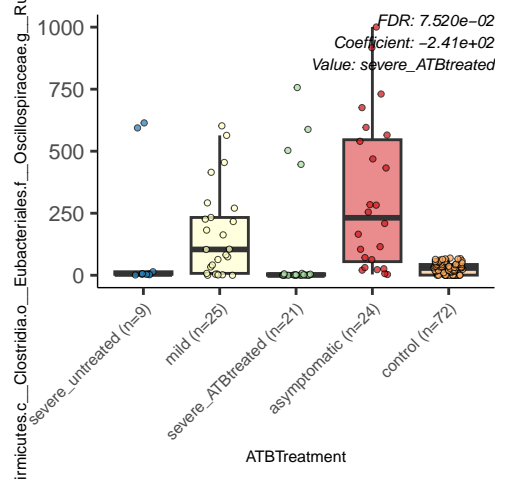


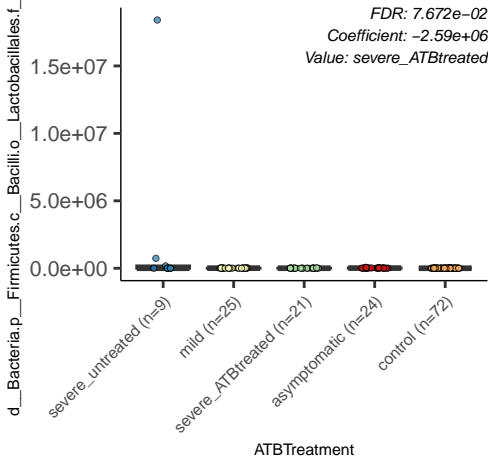


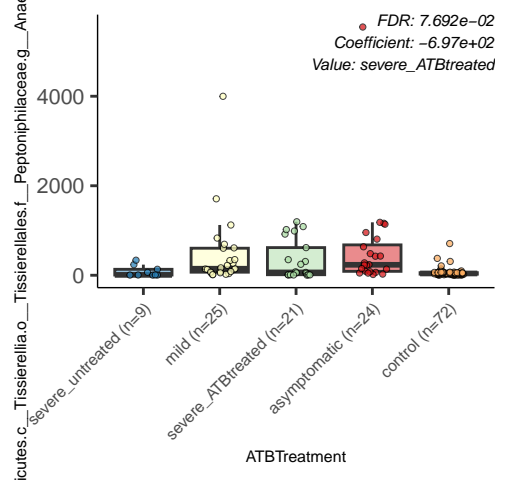


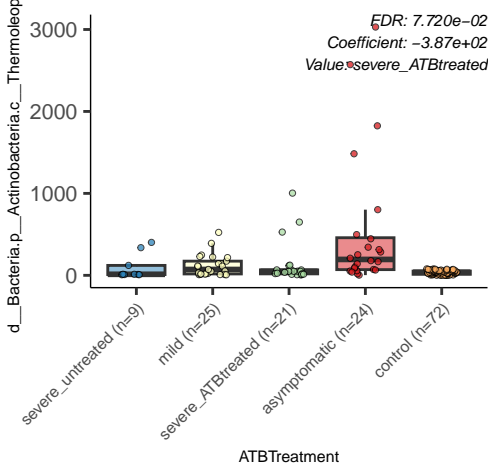




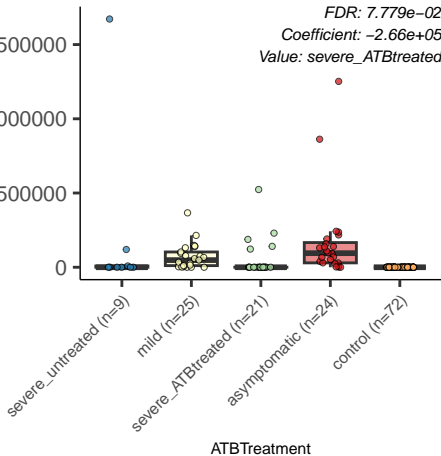






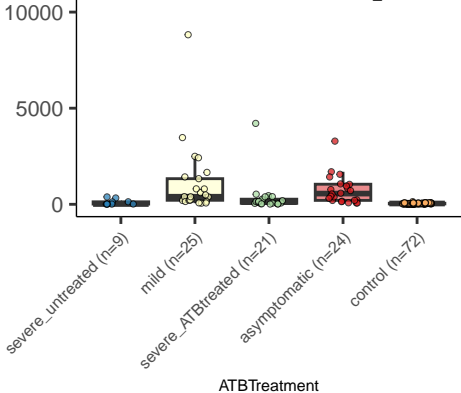


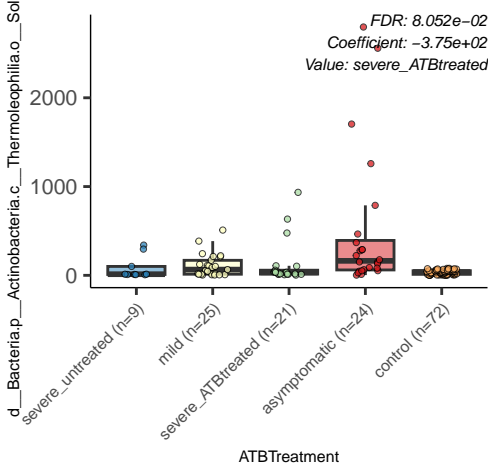
FDR: 7.779×10^{-2}
Coefficient: -2.66×10^5
Value: severe_ATBtreated

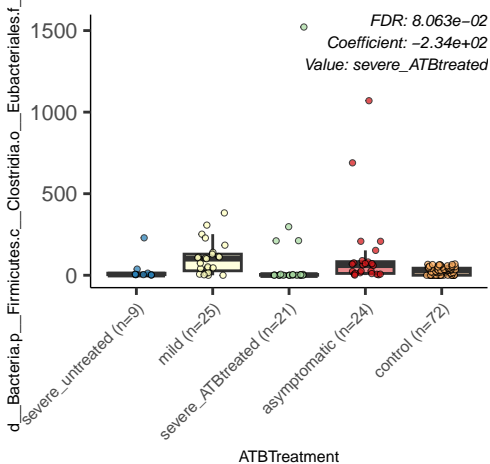


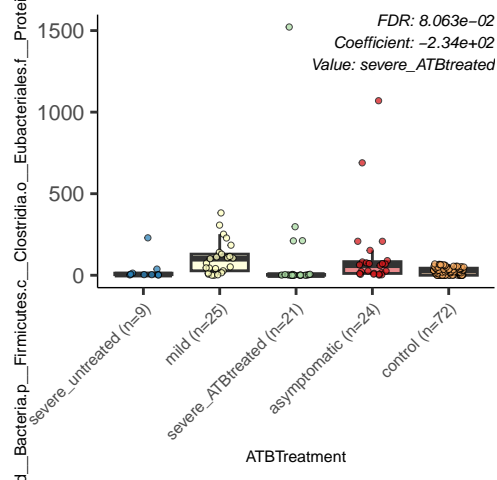
d__Bacteria.p__Cyanobacteria.o__Pseudanabaena

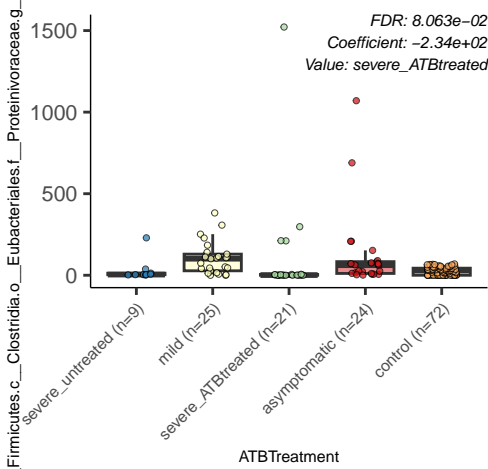
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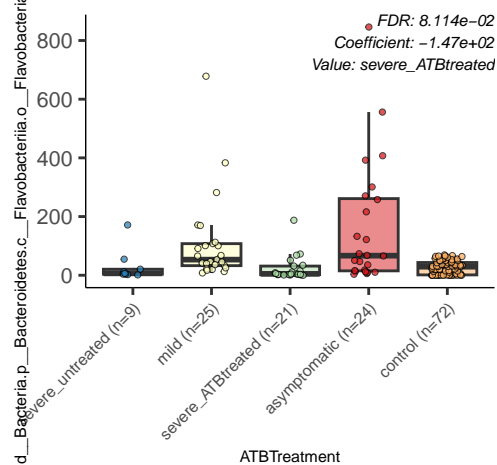


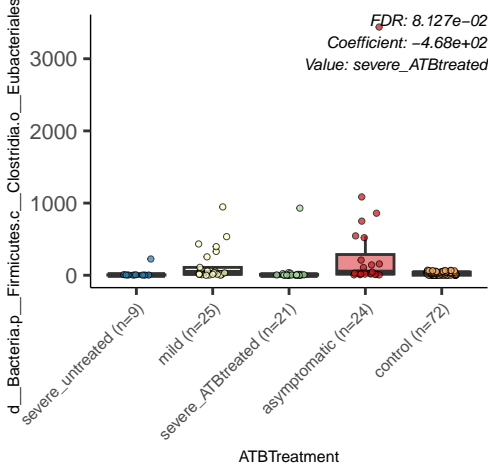


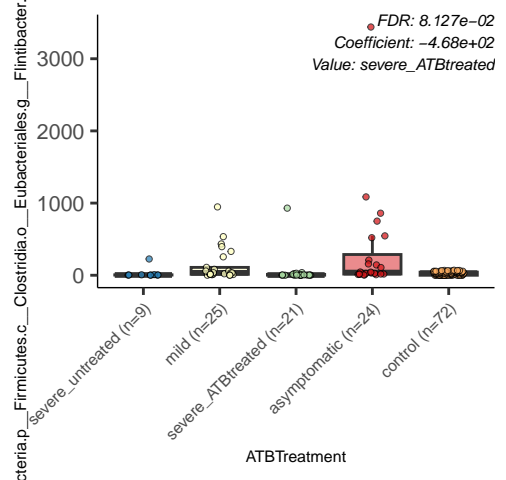


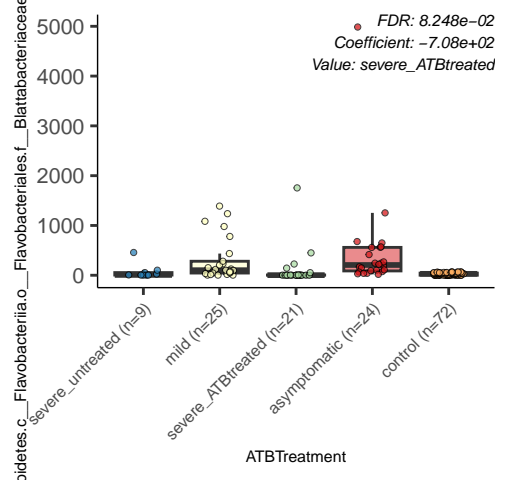


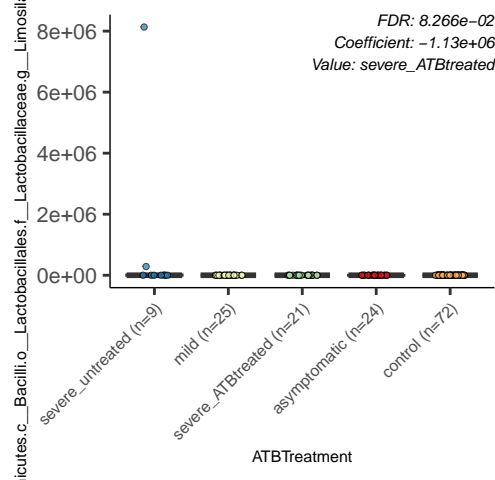


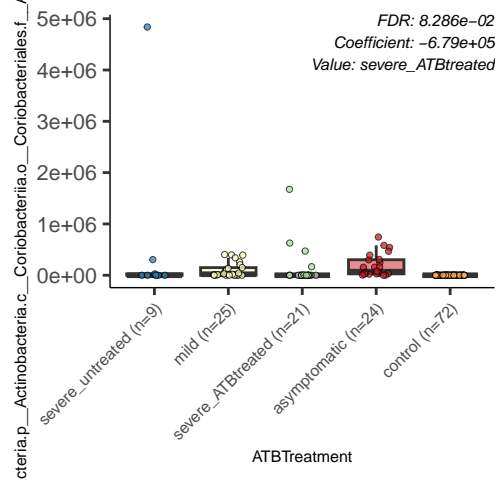


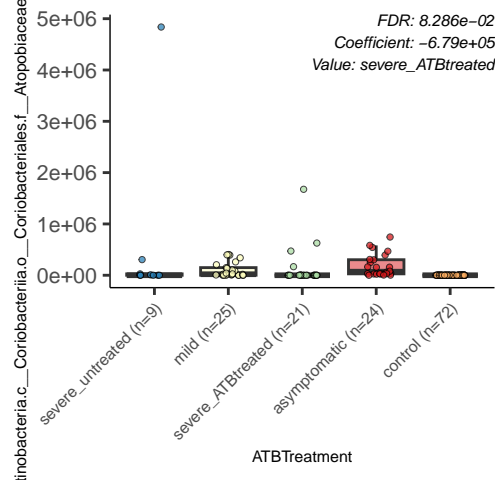


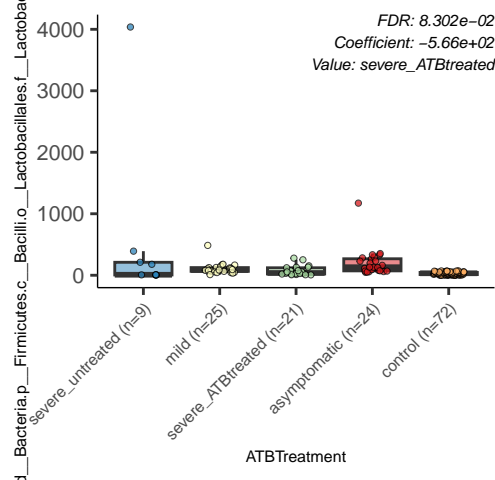


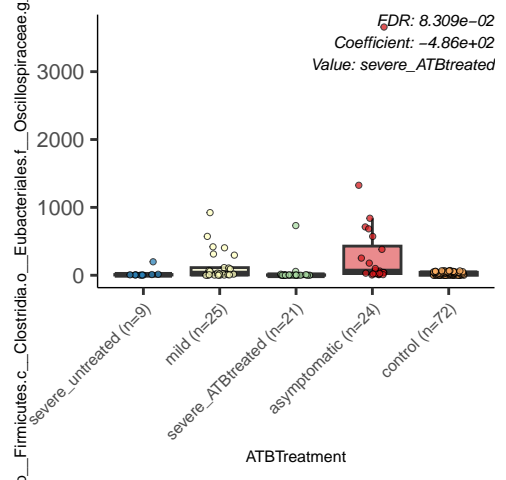


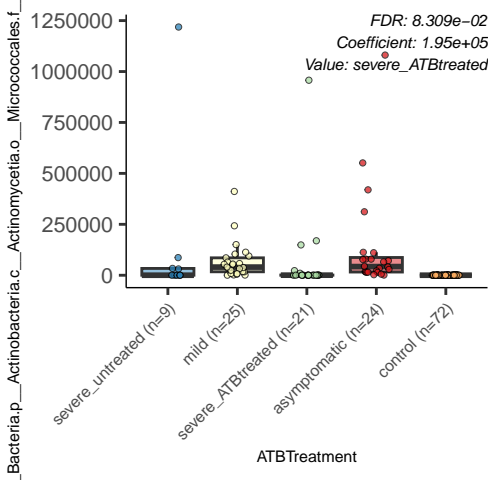


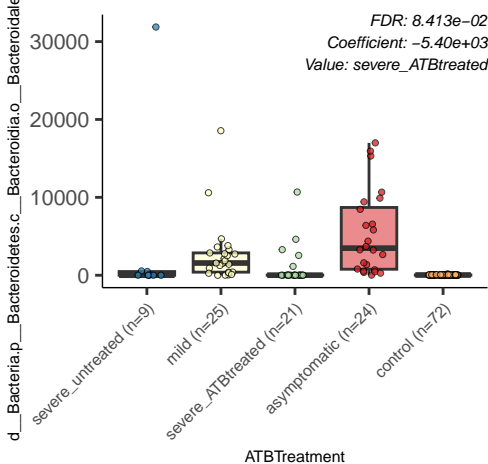


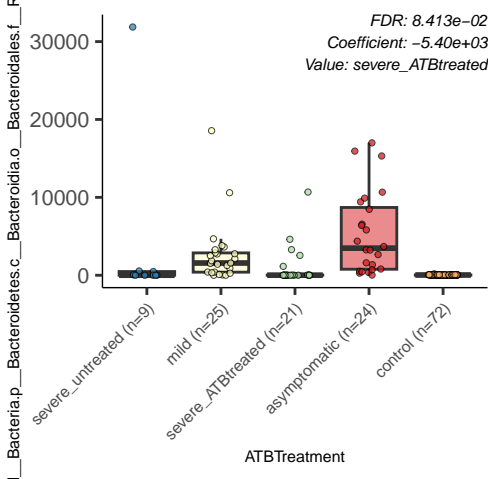


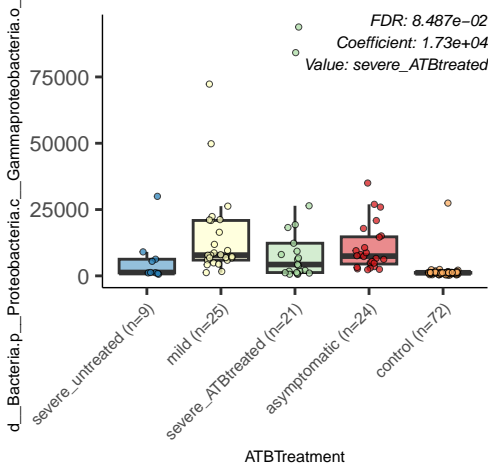


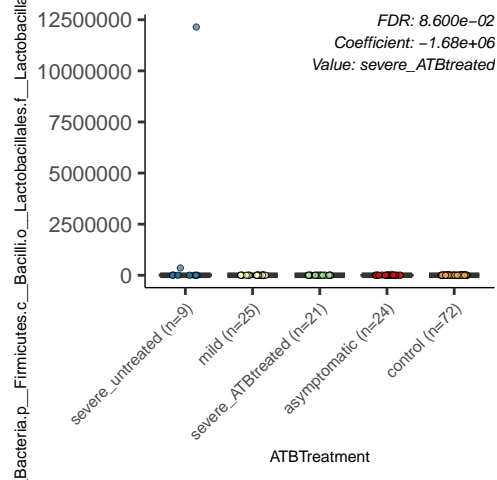




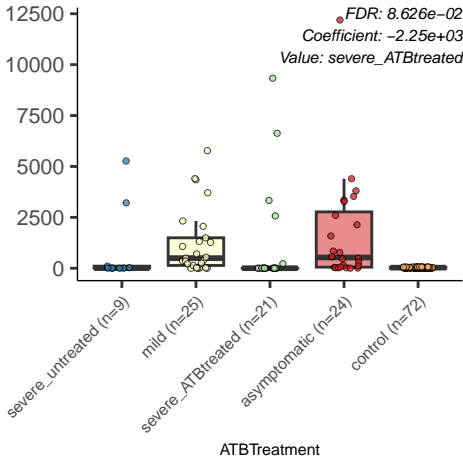


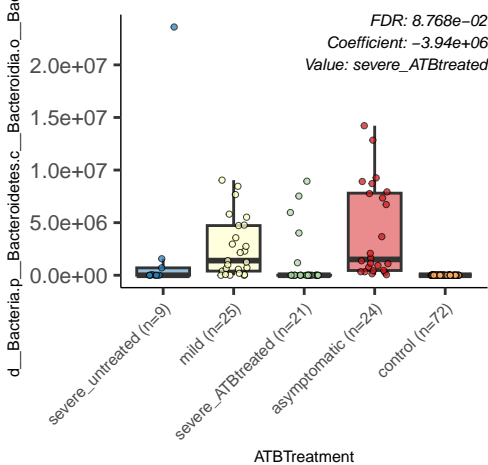


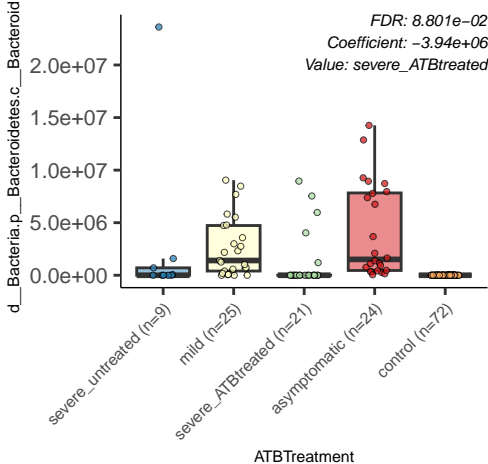


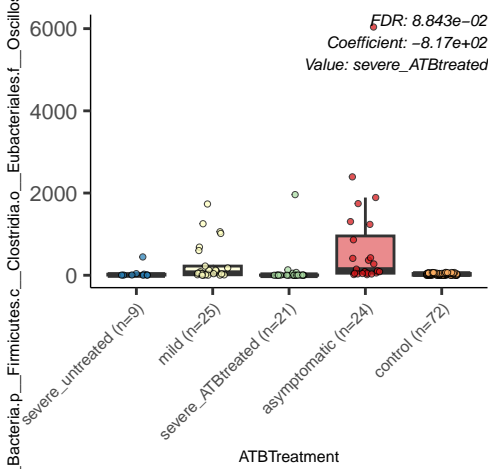


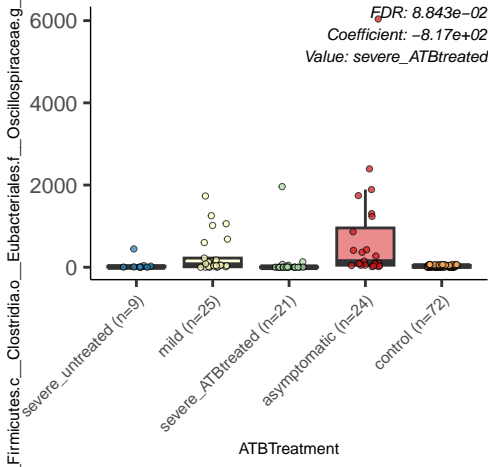
Bacteroidetes.c_Bacteroidia.o_Bacteroidales.f_Bacteroidaceae.g

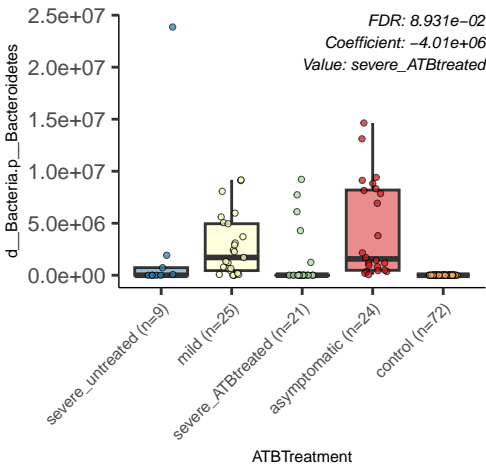


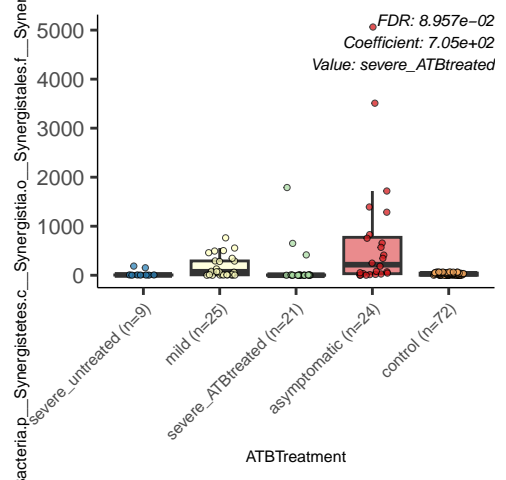


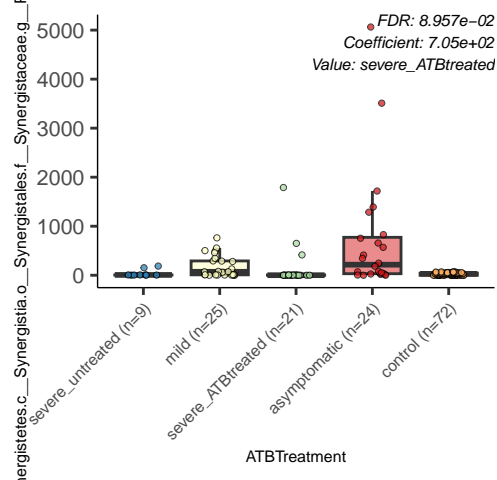


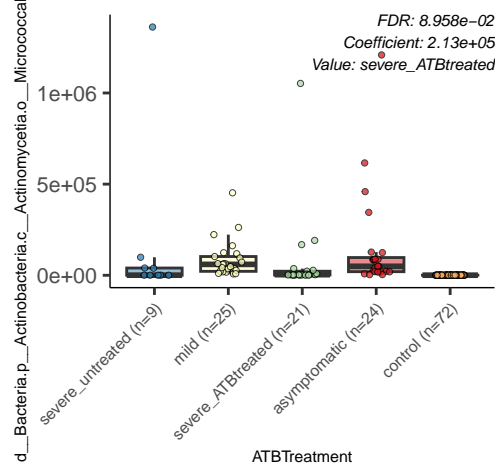


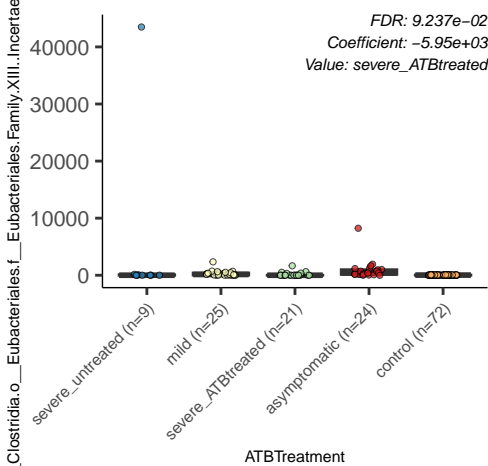


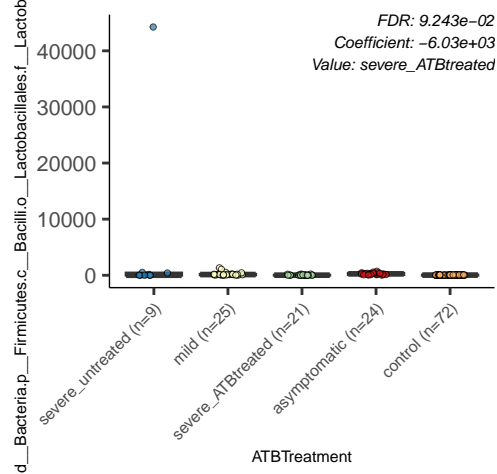


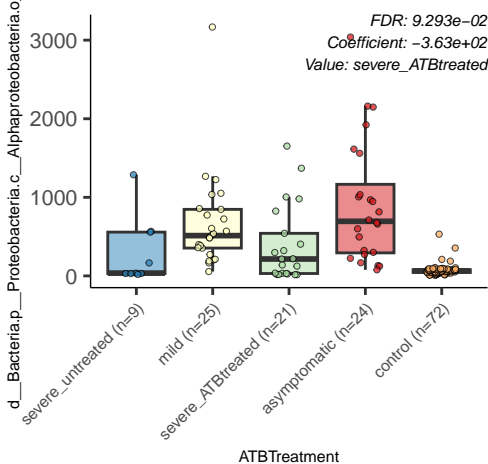


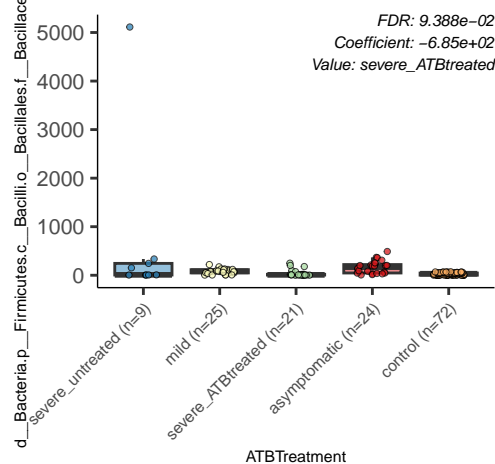


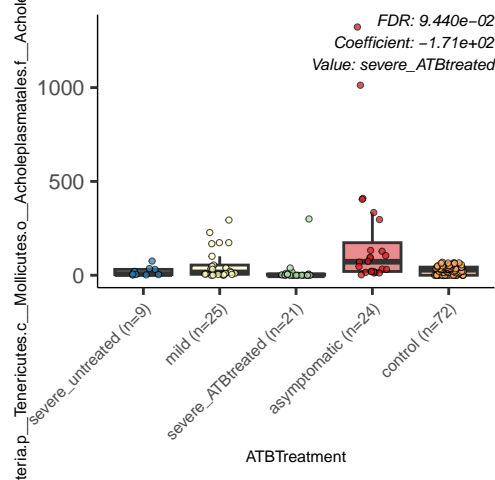


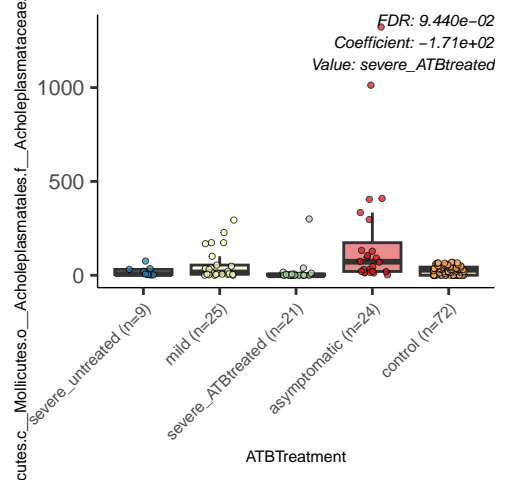


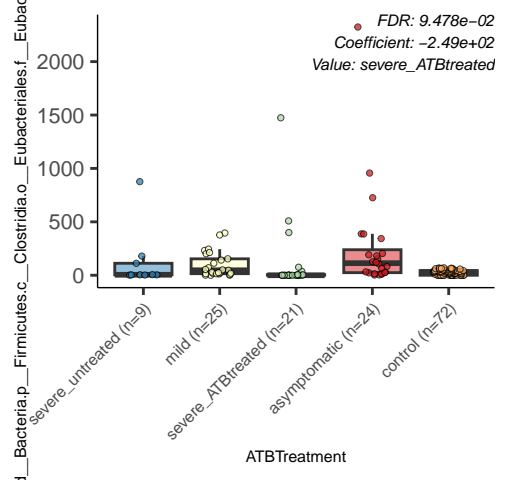




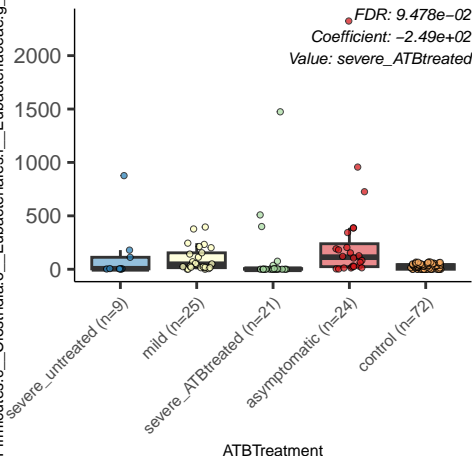


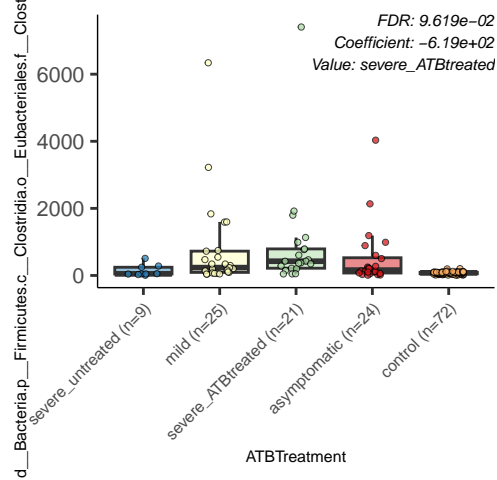


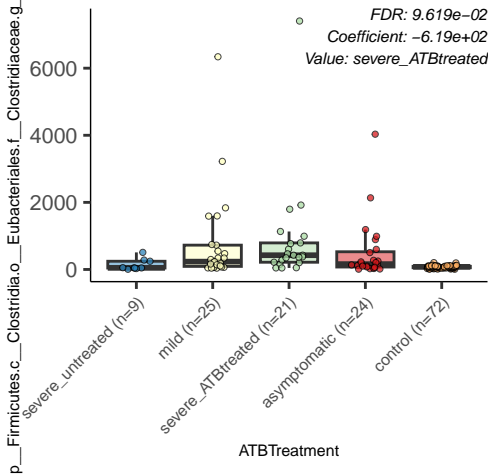


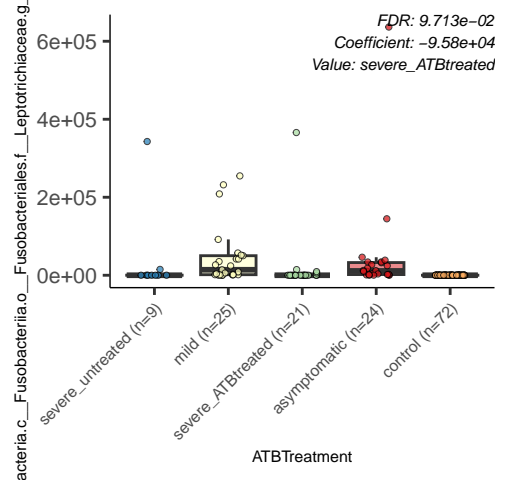


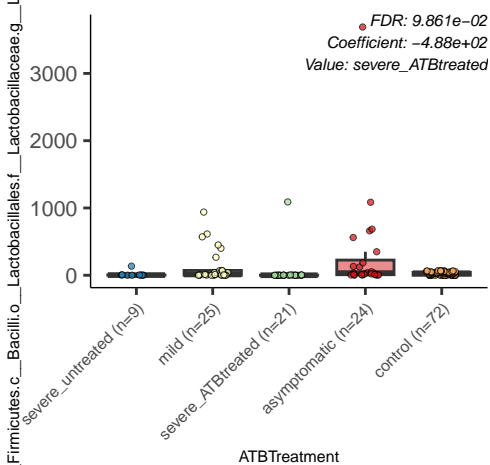
Firmicutes.c__Clostridia.o__Eubacteriales.f__Eubacteriaceae.g

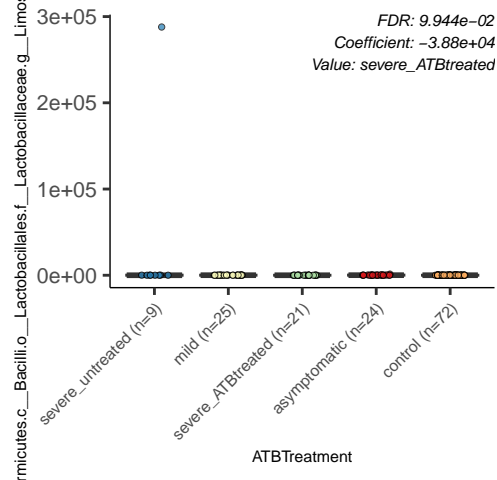


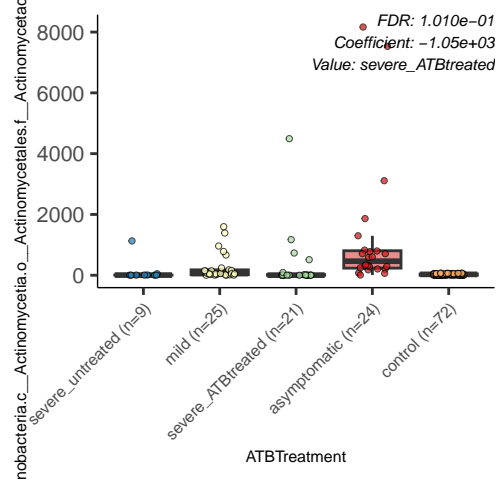


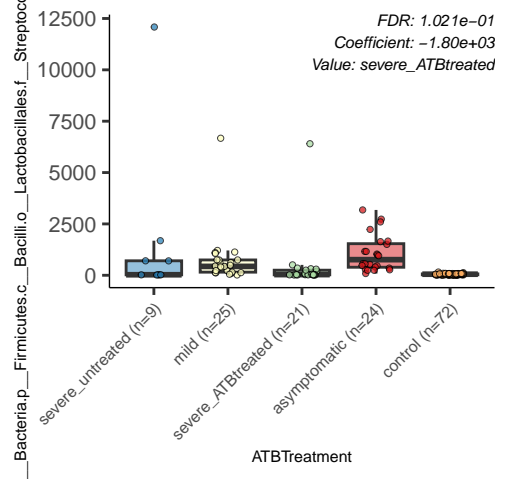


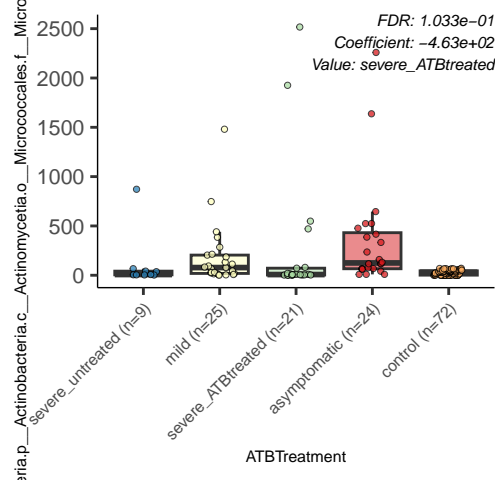




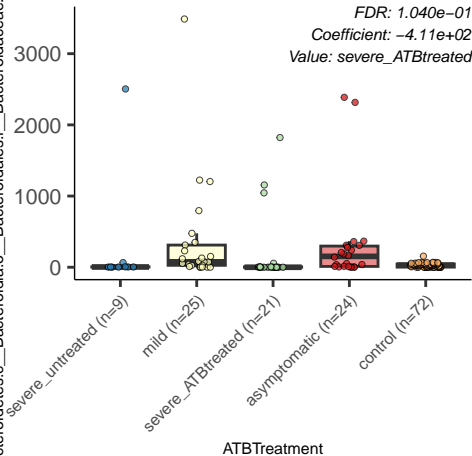


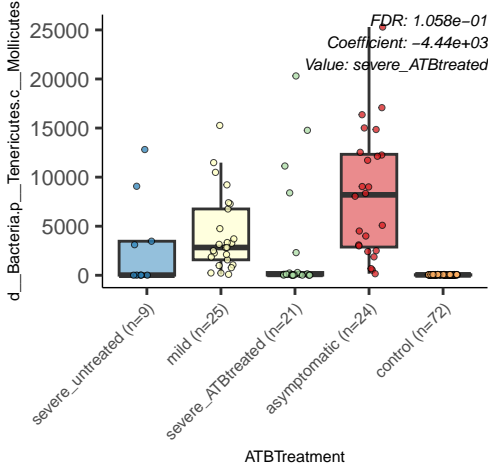


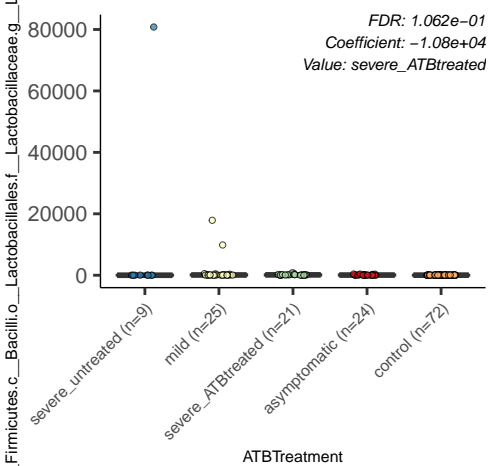


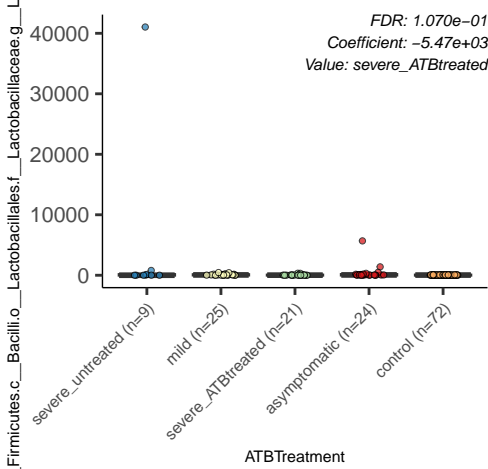


Bacteroidetes.c__Bacteroidia.o__Bacteroidales.f__Bacteroidaceae.g

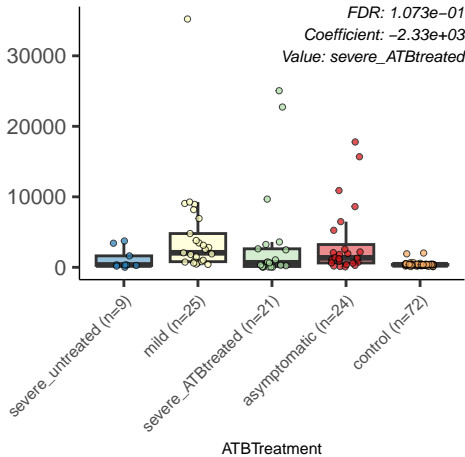




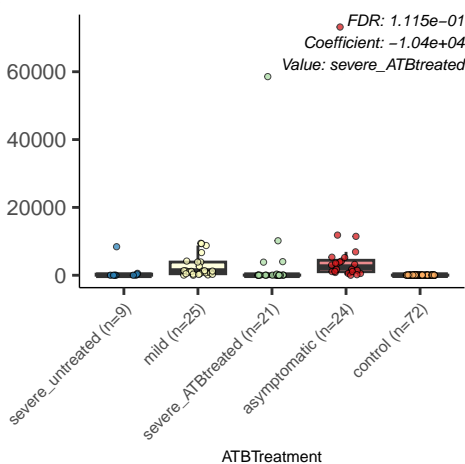


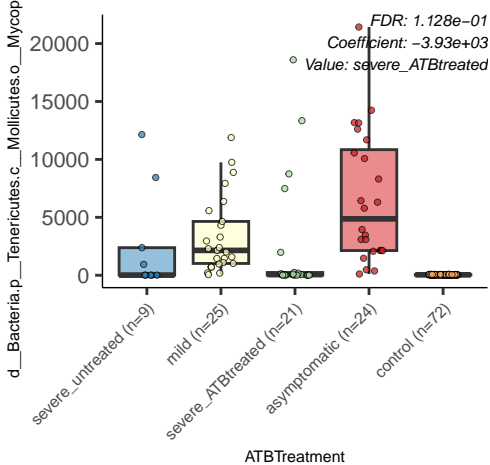


a.p_Firmicutes.c_Clostridia.o_Eubacteriales.f_Clostridiaceae.g

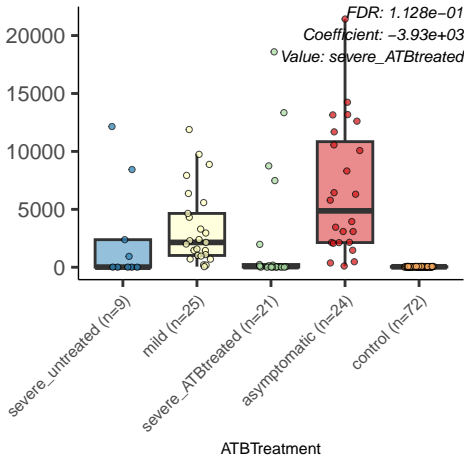


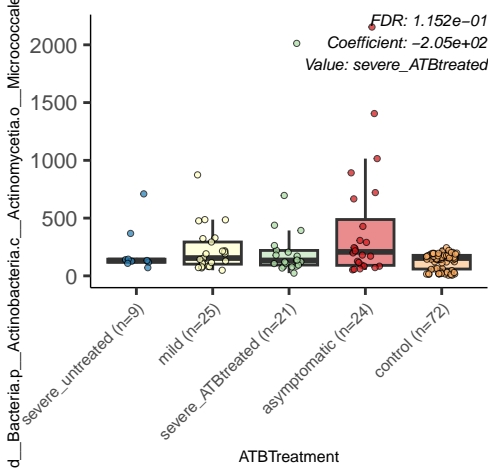
Actinomyces

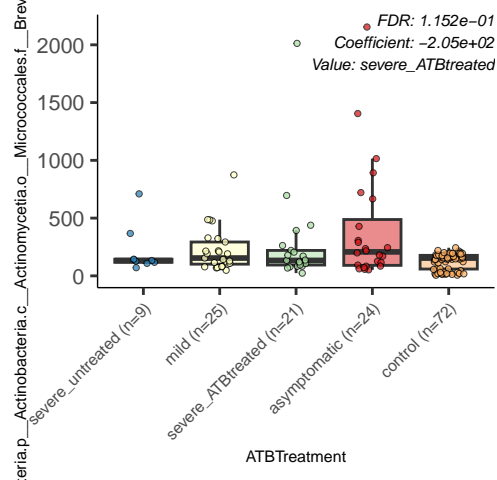




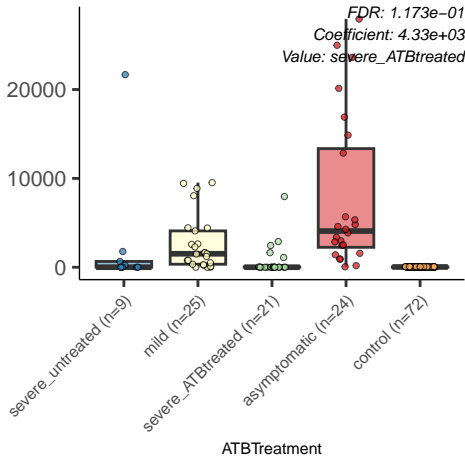
d_Bacteria.p_Tenericutes.c_Mollicutes.o_Mycoplasmales.

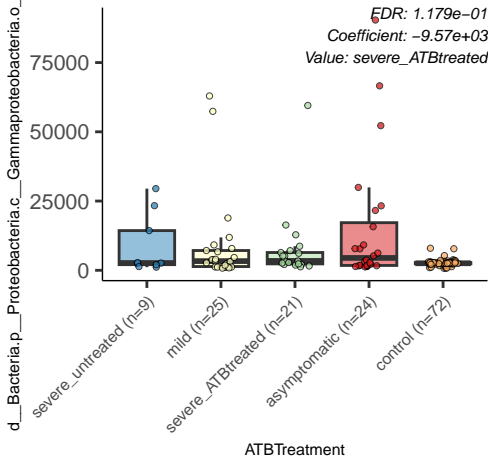


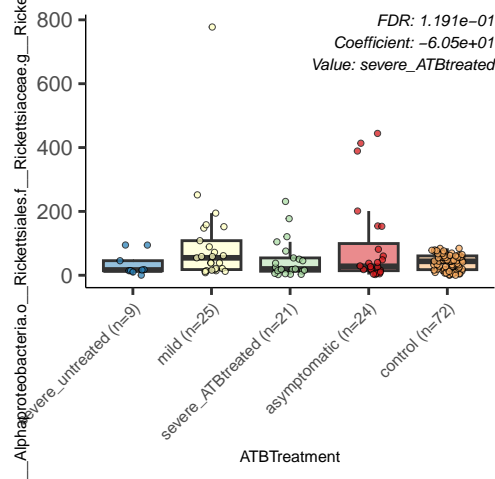


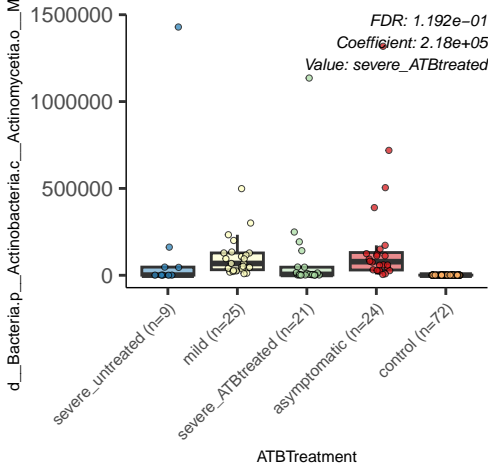


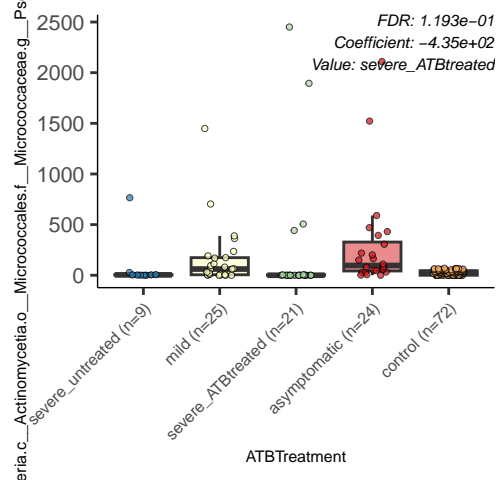
Firmicutes.c__Bacilli.o__Lactobacillales.f__Streptococcaceae.g__S

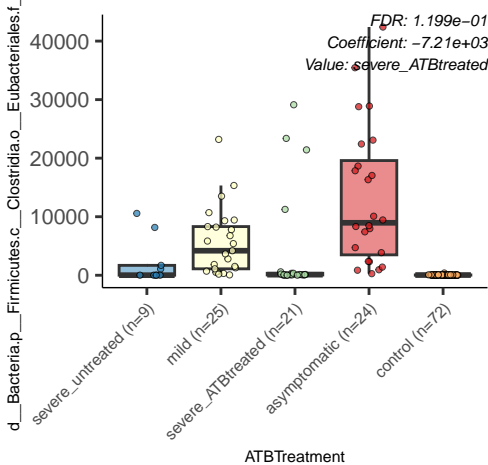




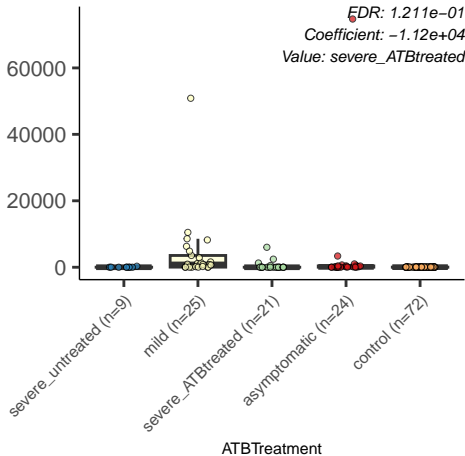


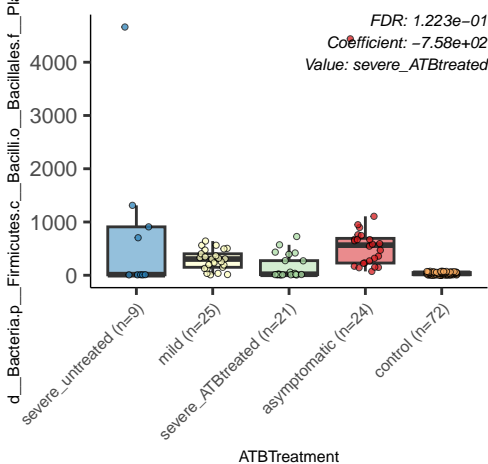


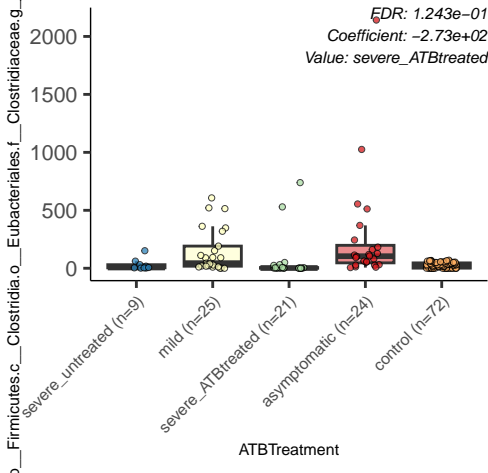


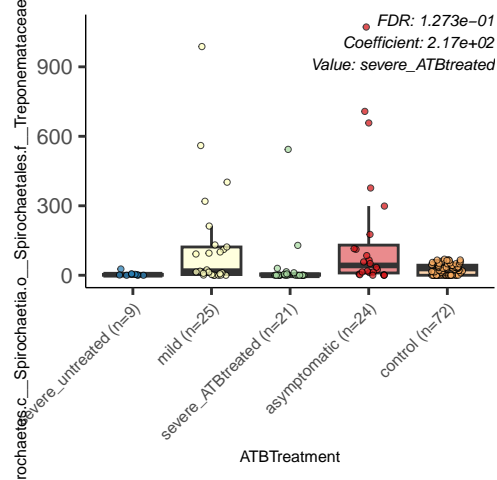


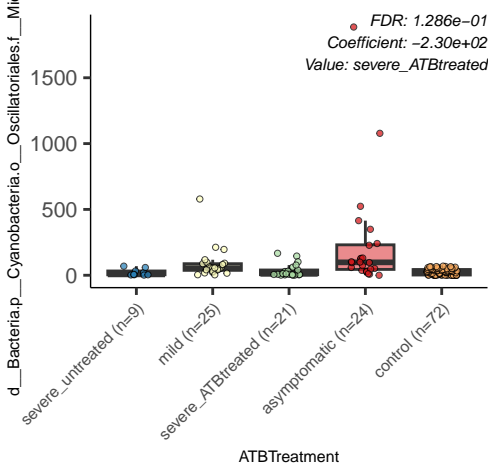
a.c._Epsilonproteobacteria.o__Campylobacteriales.f__Campylobact

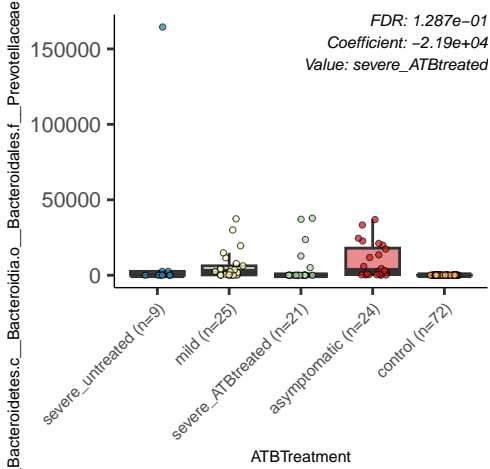


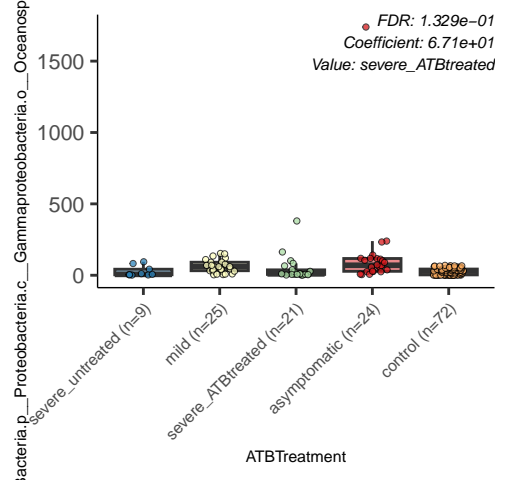


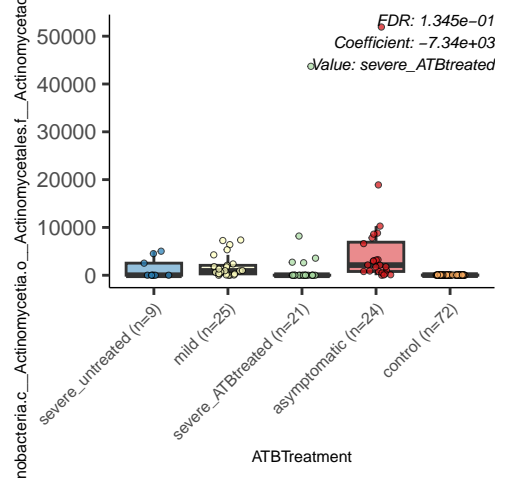


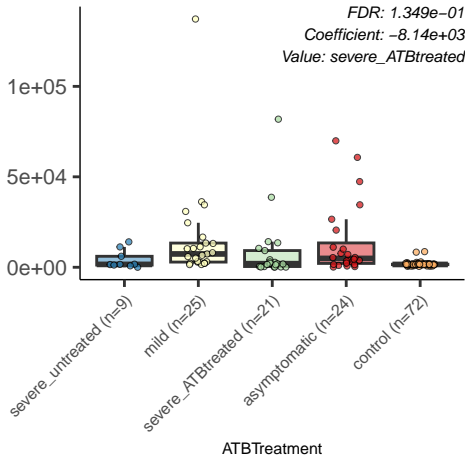


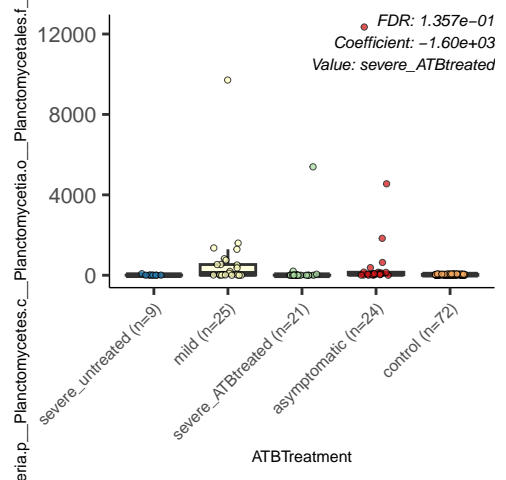


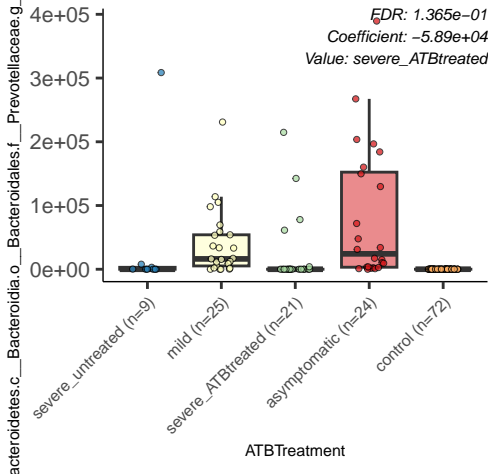






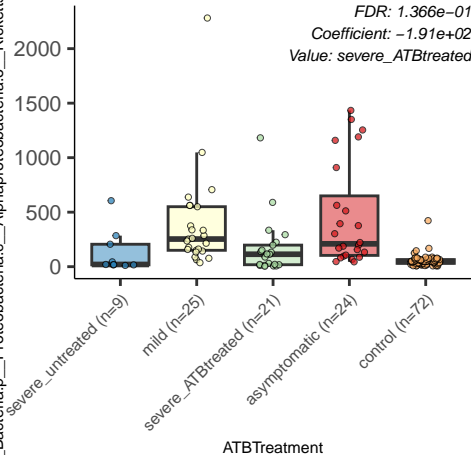




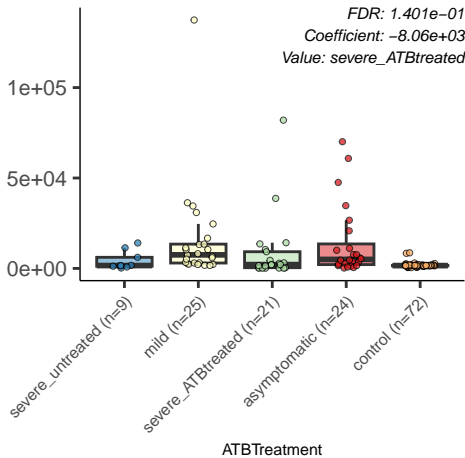


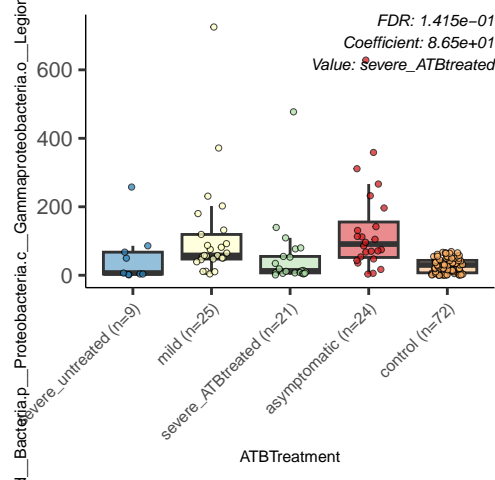
l_Bacteria.p__Proteobacteria.c__Alphaproteobacteria.o__Ricketts

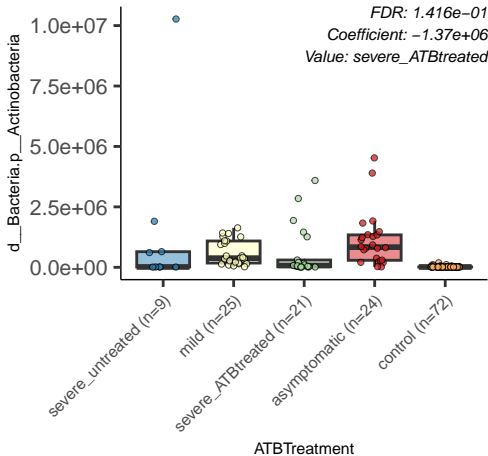
FDR: $1.366e-01$
Coefficient: $-1.91e+02$
Value: severe_ATBtreated

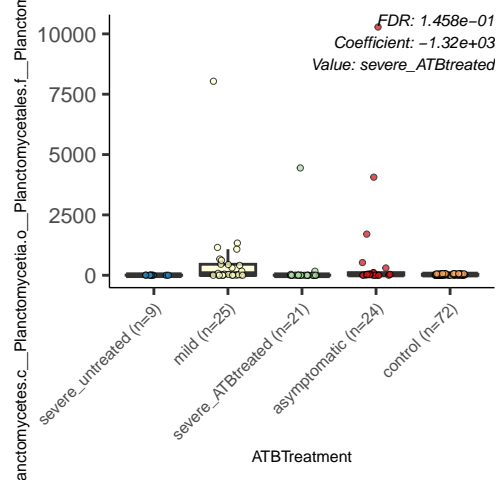


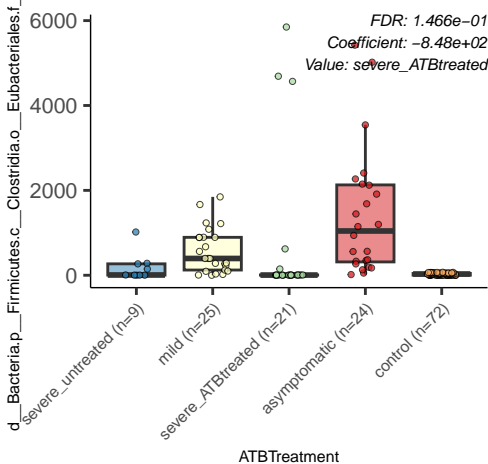
.p__Proteobacteria.c__Gammaproteobacteria.o__Oceanospirillales

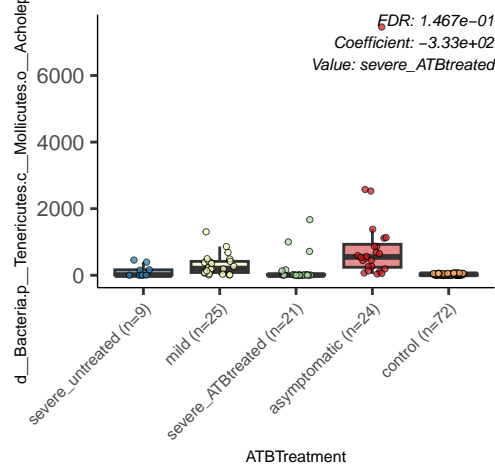












d_Bacteria.p__Tenericutes.c__Mollicutes.o__Acholeplasmatales.

FDR: 1.467×10^{-1}
Coefficient: -3.33×10^2
Value: severe_ATBtreated

6000

4000

2000

0

severe_unreated (n=9)

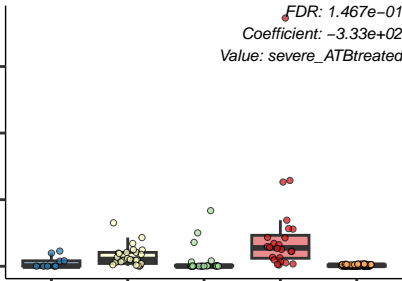
mild (n=25)

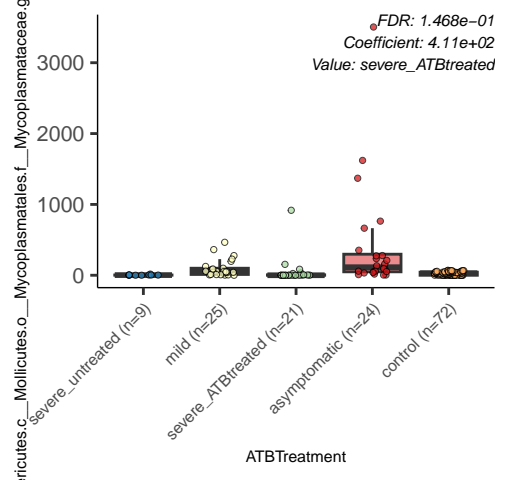
severe_ATBtreated (n=21)

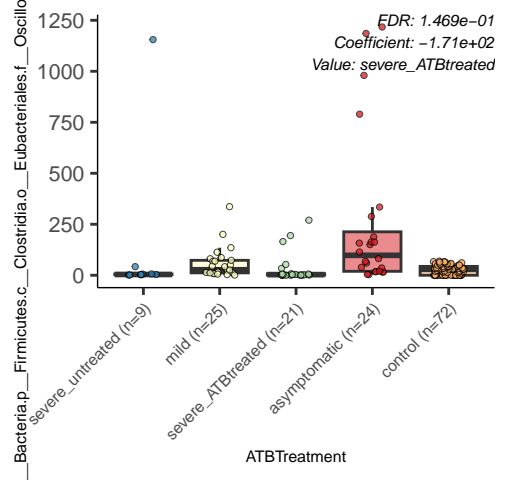
asymptomatic (n=24)

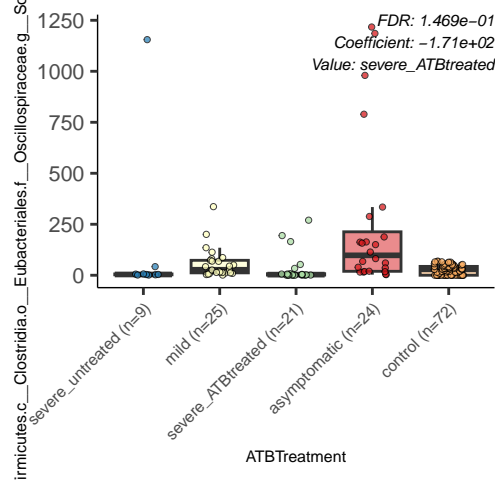
control (n=72)

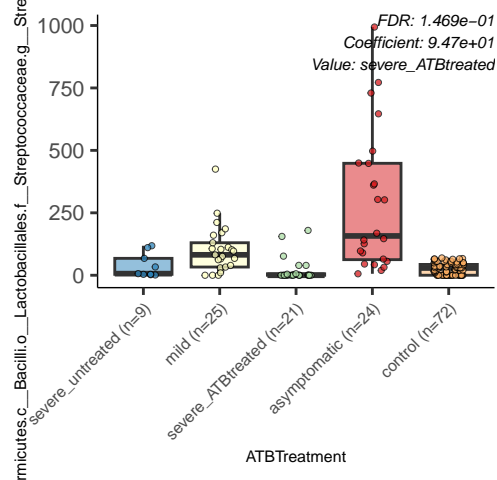
ATBTreatment

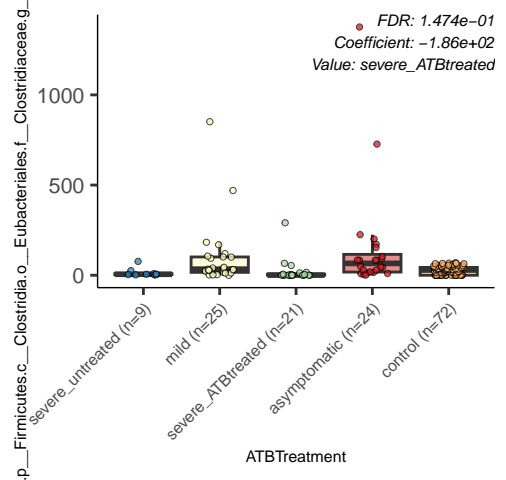


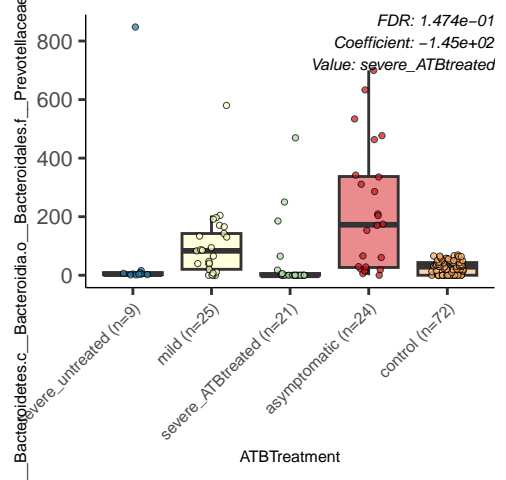


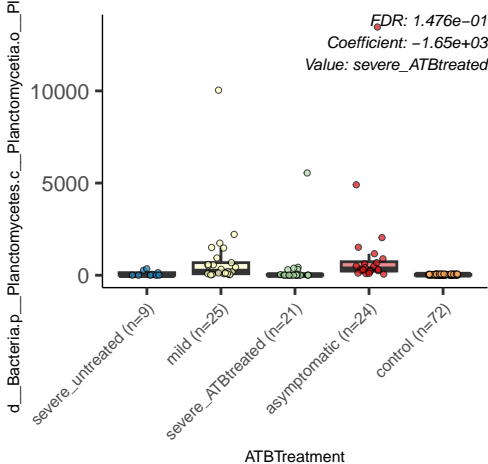


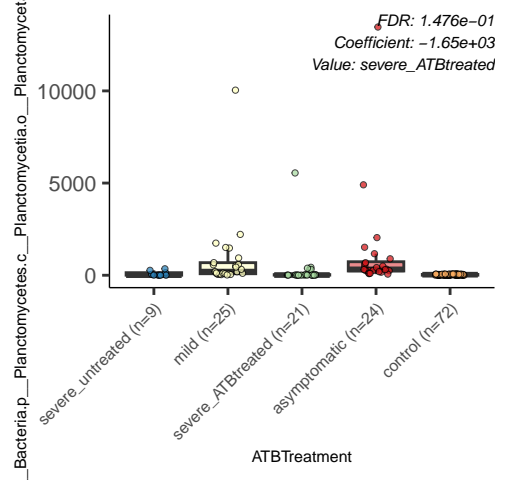


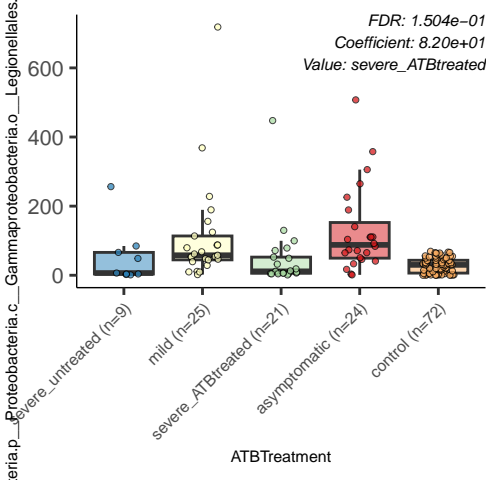


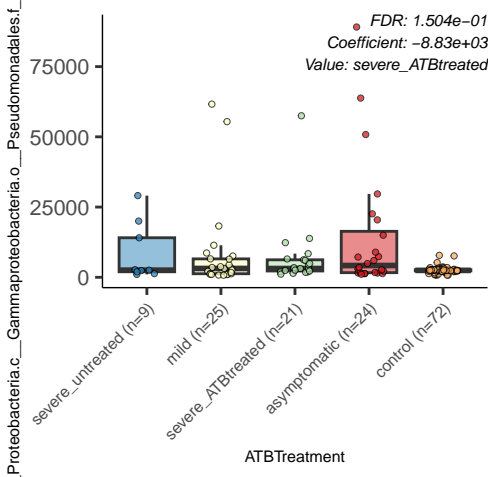


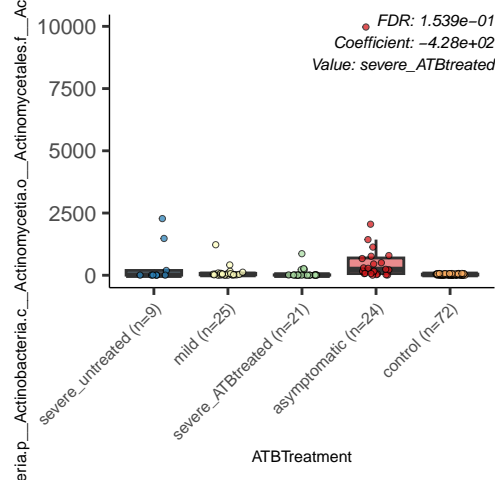


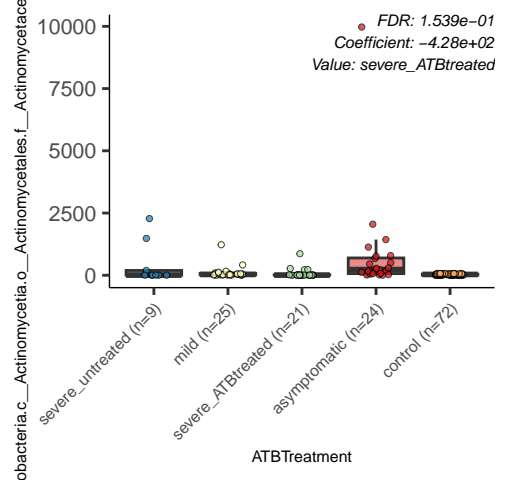


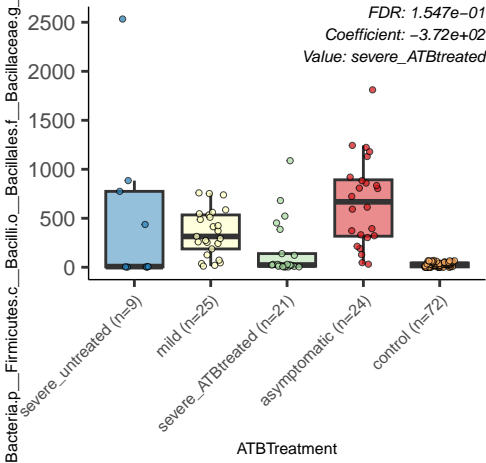


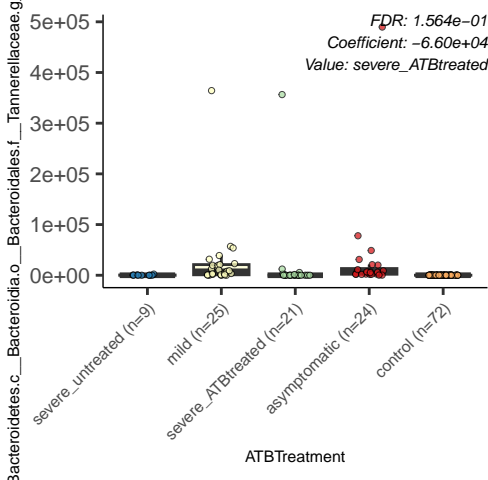


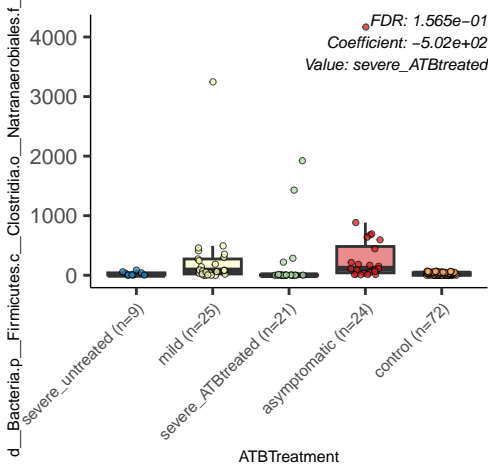


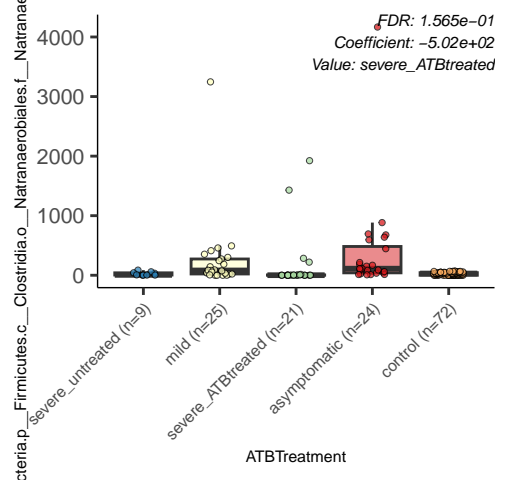


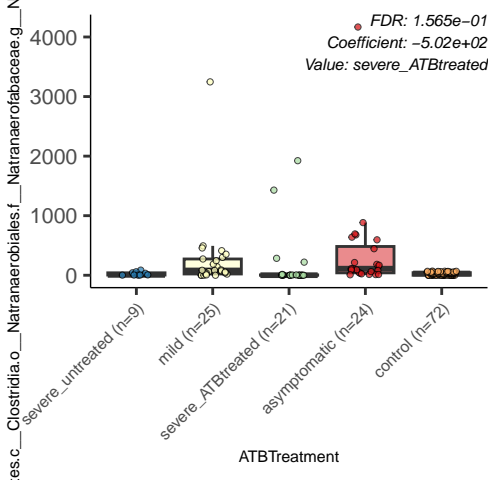




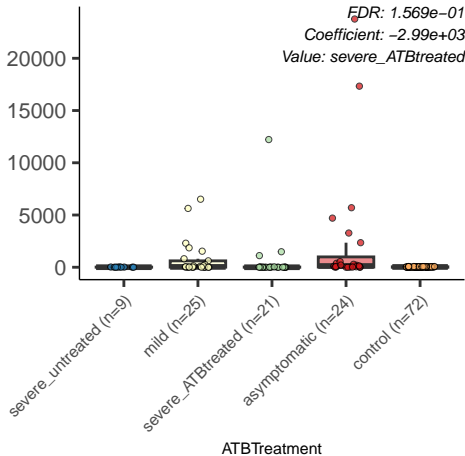


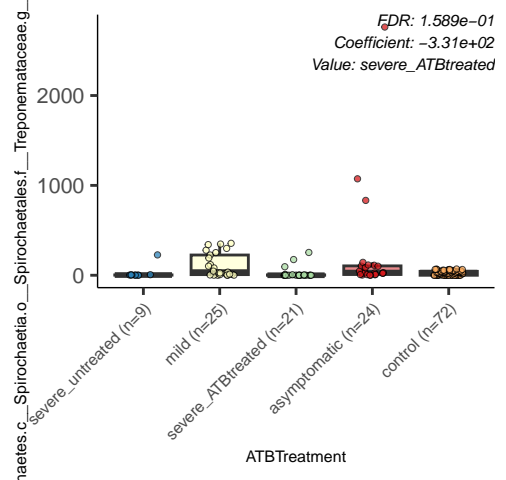


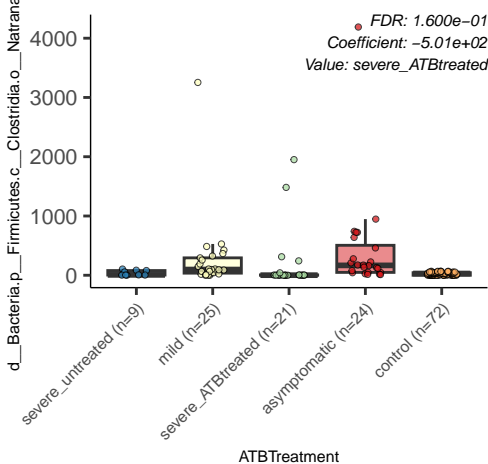


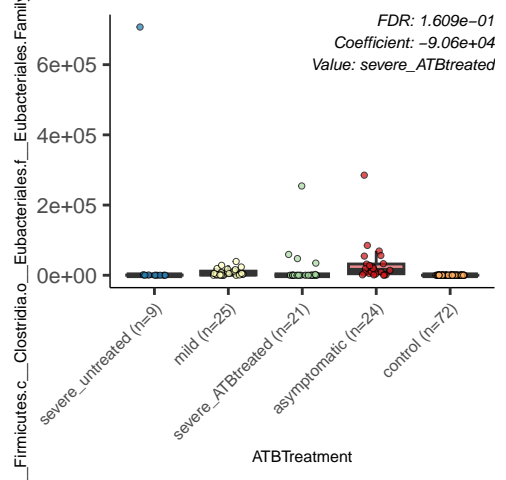


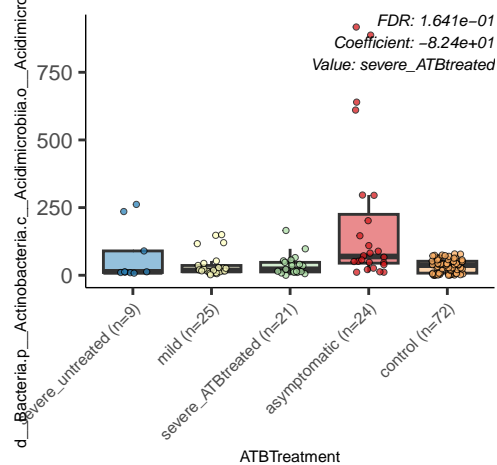
roidetes.c_Flavobacteriia.o_Flavobacteriales.f_Flavobacteriaceae

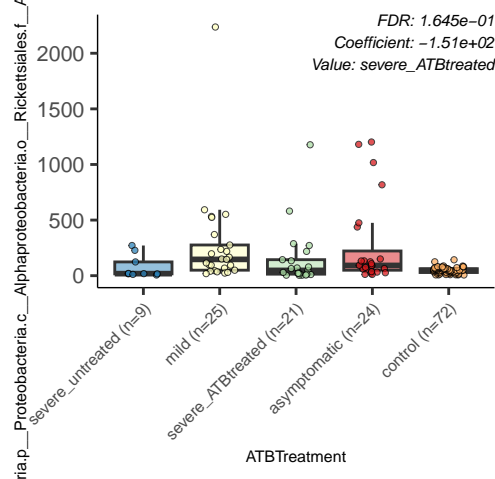




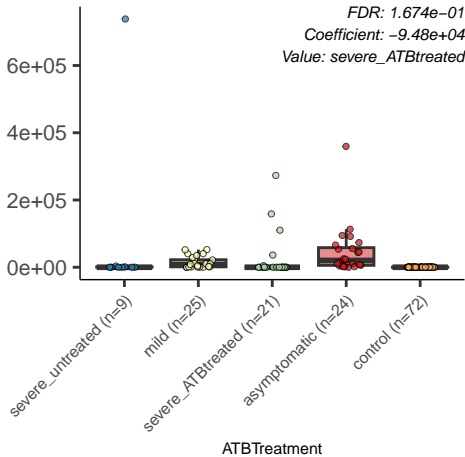


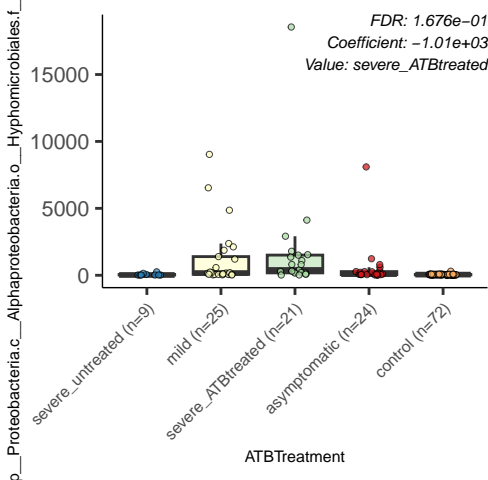






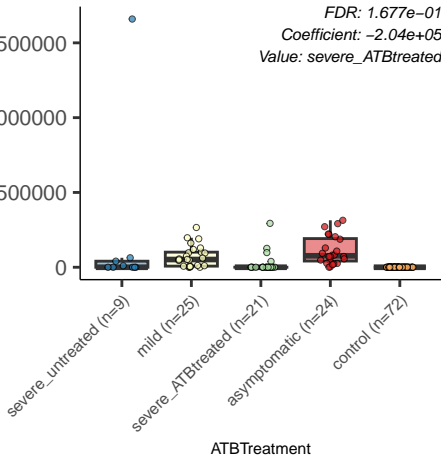
acteria.p_Firmicutes.c_Clostridia.o_Eubacteriales.f_Eubacteri



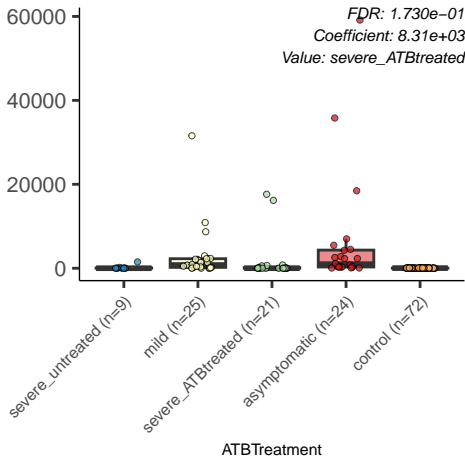


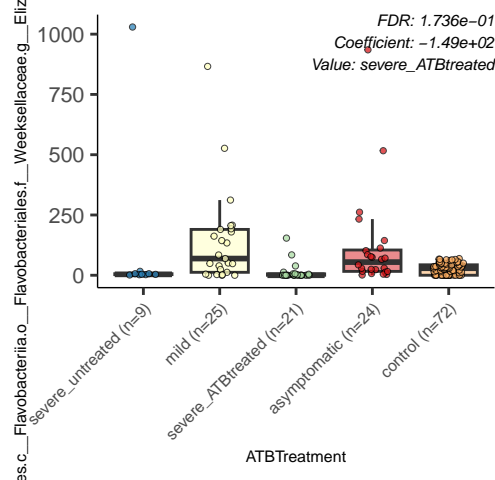
p__Firmicutes.c__Negativicutes.o__Veillonellales.f__Veillonellaceae

FDR: $1.677\text{e-}01$
Coefficient: $-2.04\text{e+}05$
Value: severe_ATBtreated

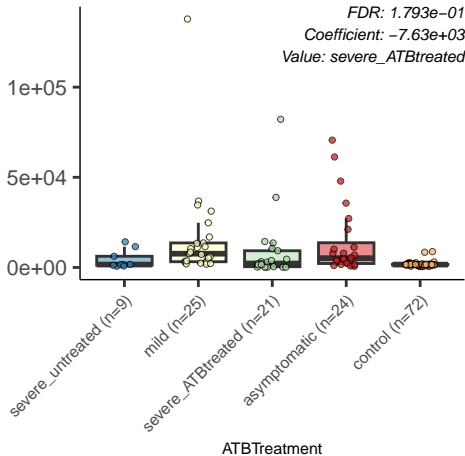


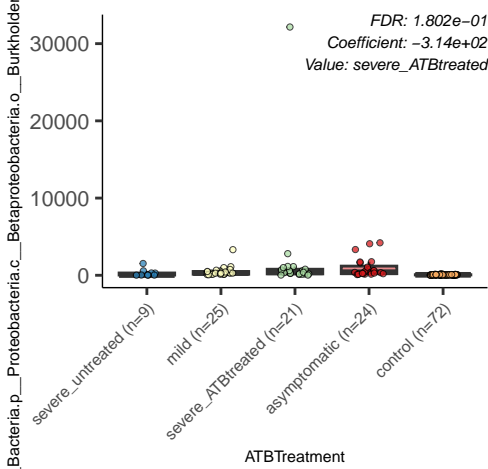
bidetes.c__Bacteroidia.o__Bacteroidales.f__Porphyromonadaceae.g

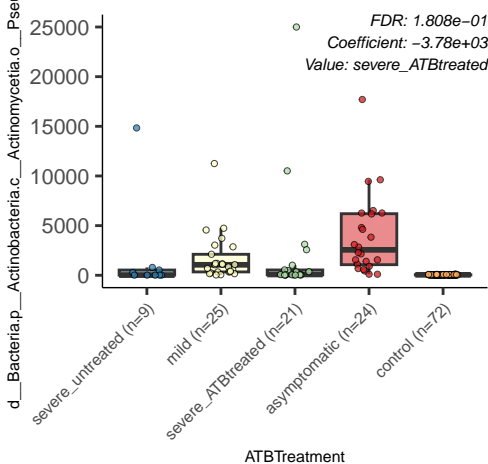


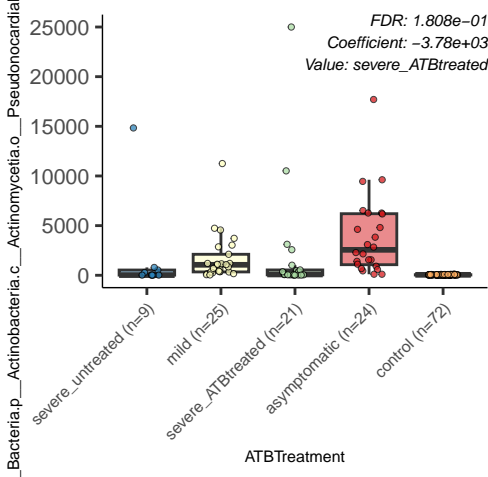


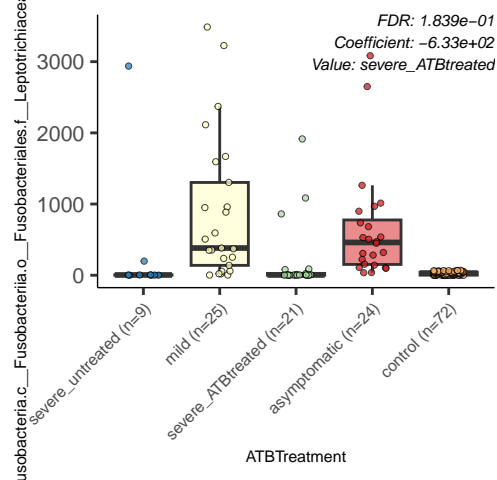
Bacteria.p__Proteobacteria.c__Gammaproteobacteria.o__Oceanos

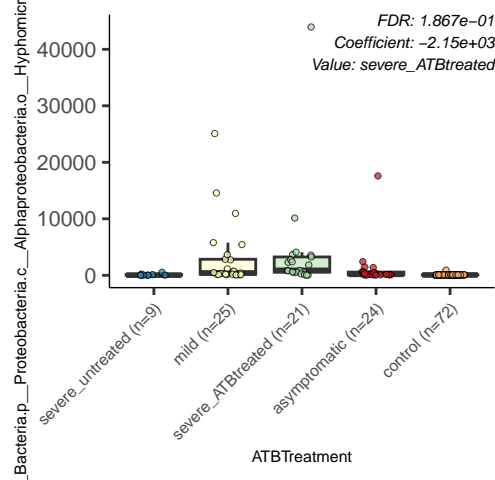












__Bacteroidetes.c__Bacteroidia.o__Bacteroidales.f__Prevotellaceae

FDR: $1.901e-01$
Coefficient: $-2.90e+04$
Value: severe_ATBtreated

150000

100000

50000

0

severe_unreated (n=9)

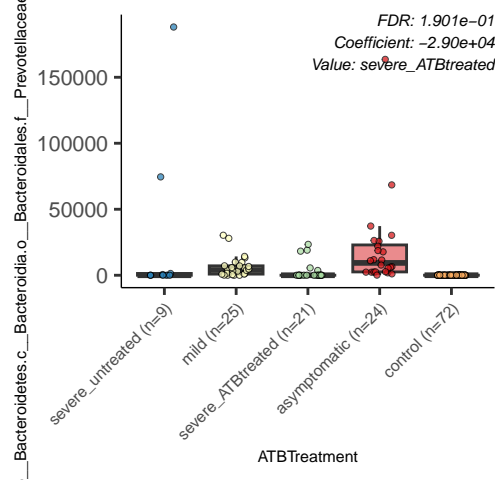
mild (n=25)

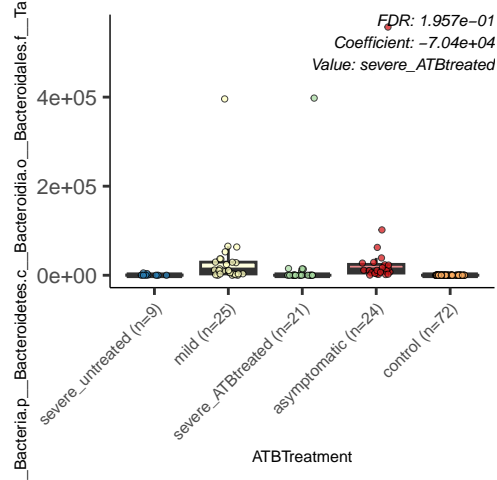
severe_ATBtreated (n=21)

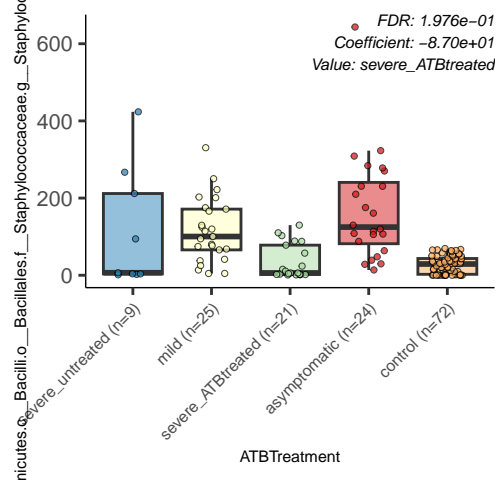
asymptomatic (n=24)

control (n=72)

ATBTreatment

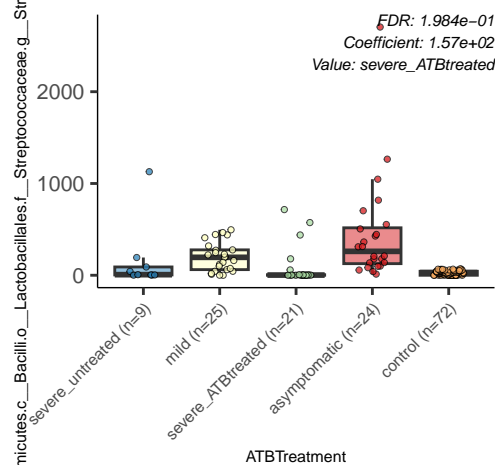


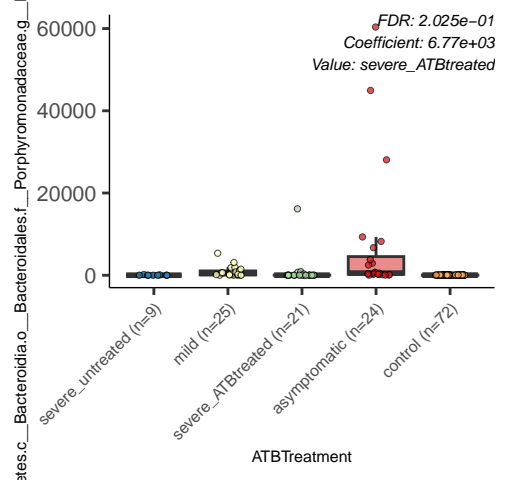


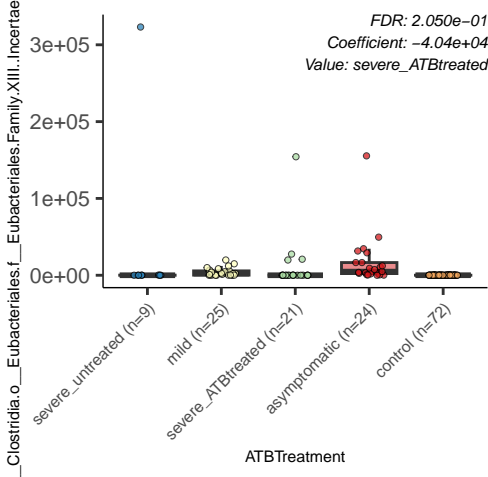


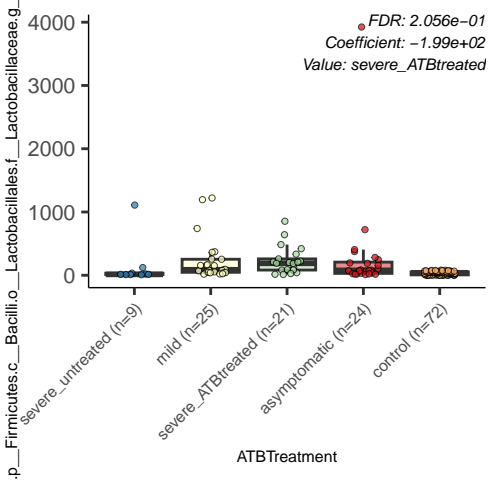
Firmicutes.c__Bacilli.o__Lactobacillales.f__Streptococcaceae.g__Str

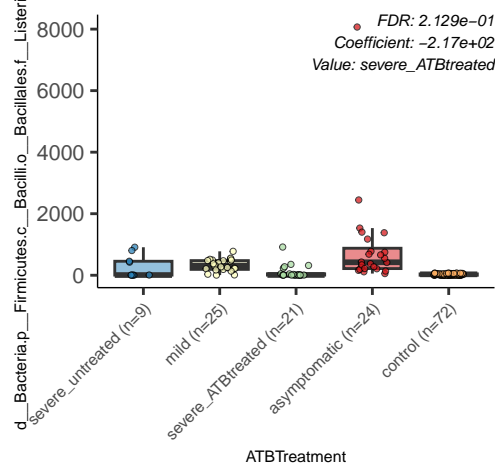
FDR: 1.984e-01
Coefficient: 1.57e+02
Value: severe_ATBtreated

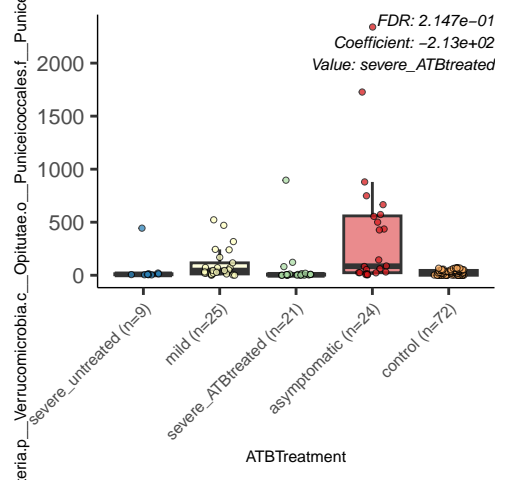


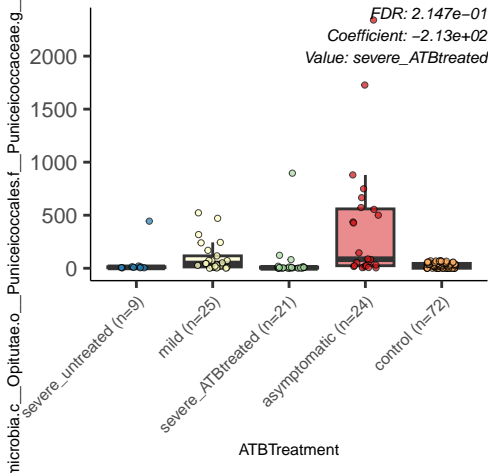


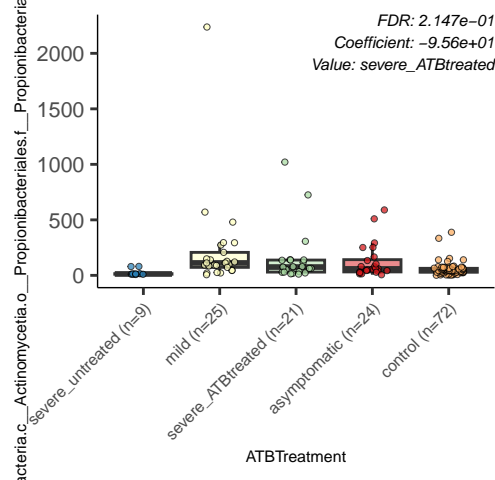


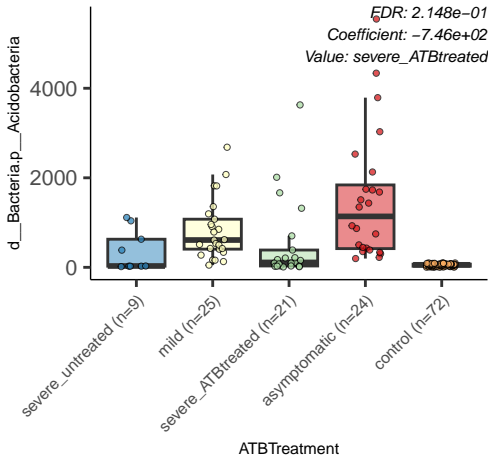


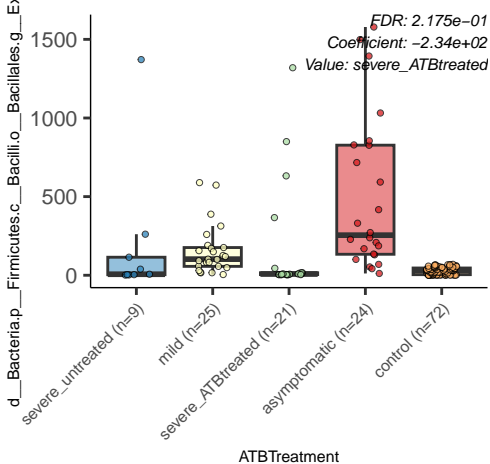


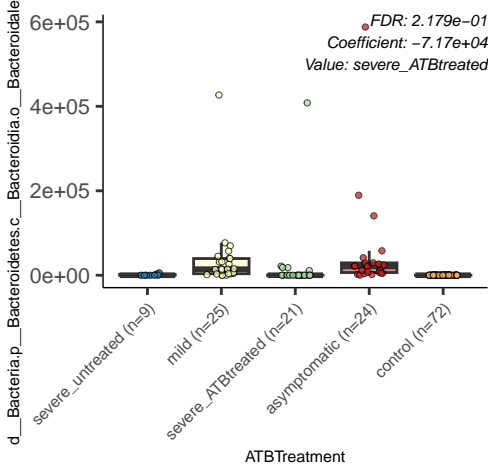


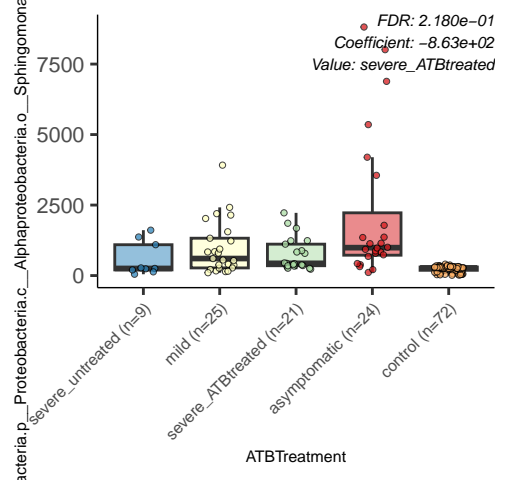












acteria.p_Bacteroidetes.c_Flavobacteriia.o_Flavobacteriales.f

FDR: 2.183e-01
Coefficient: -1.71e+03
Value: severe_ATBtreated

10000

5000

0

severe_untreated (n=9)

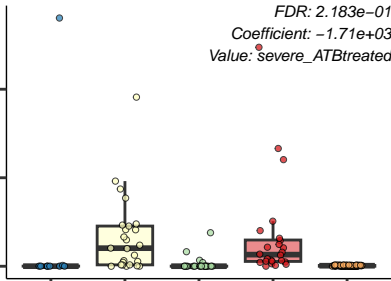
mild (n=25)

severe_ATBtreated (n=21)

asymptomatic (n=24)

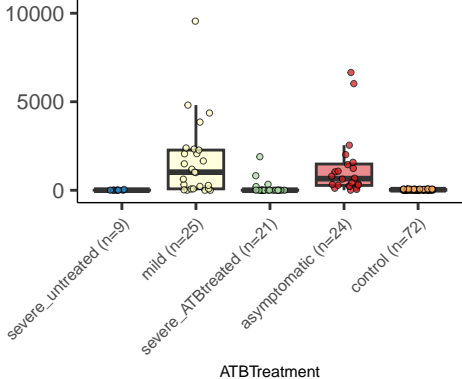
control (n=72)

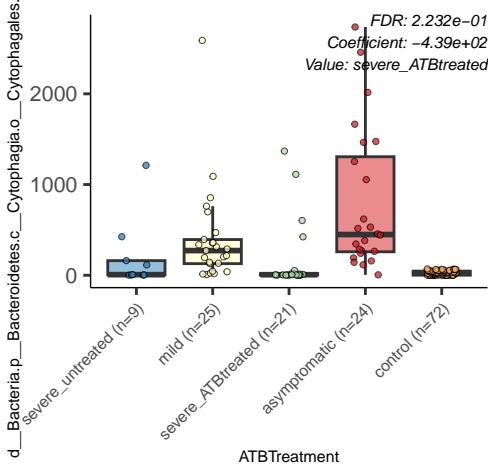
ATBTreatment

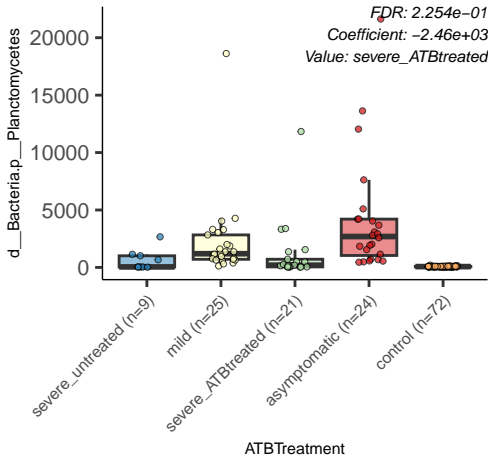


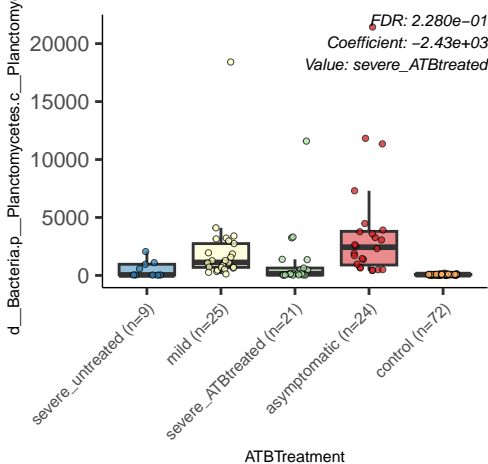
Bacteroidetes.c_Flavobacteriia.o_Flavobacteriales.f_Weeksellac

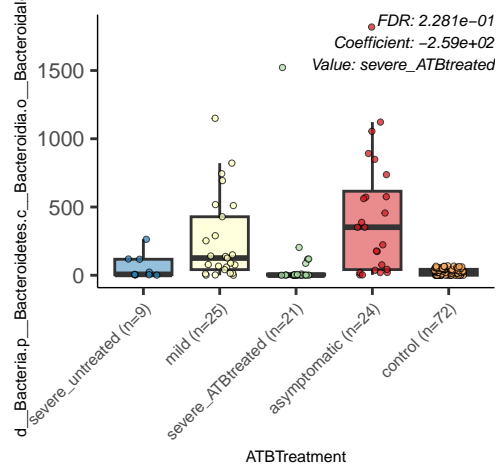
FDR: $2.183e-01$
Coefficient: $-1.71e+03$
Value: severe_ATBtreated

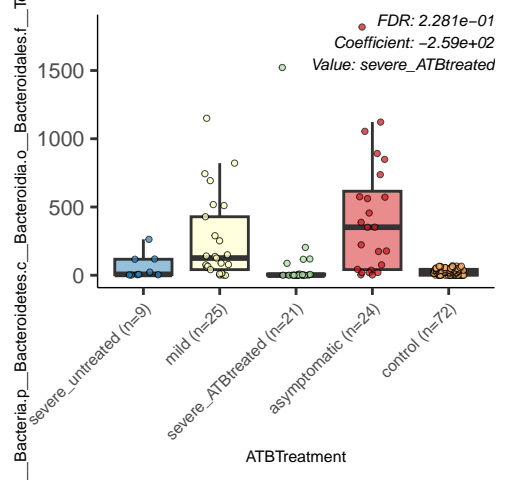




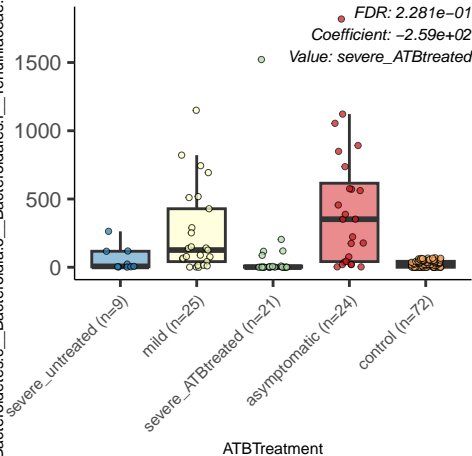


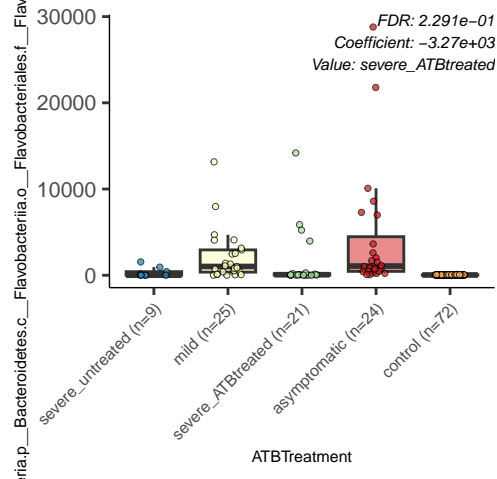


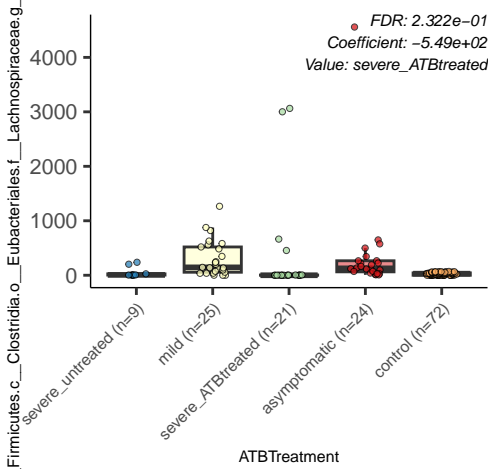


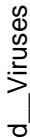


Bacteroidetes.c_Bacteroidia.o_Bacteroidales.f_Tenuifilaceae.g









FDR: 2.322e-01

Coefficient: $-1.94e+06$

Value: severe_ATBtreated

