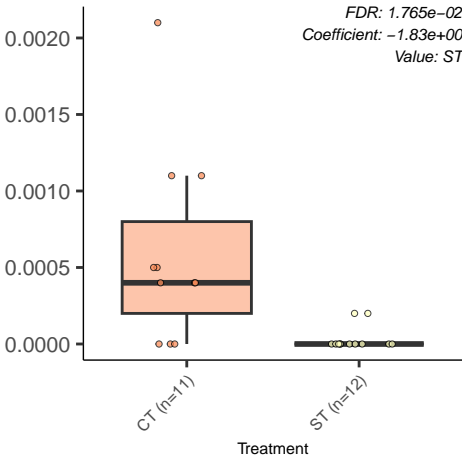
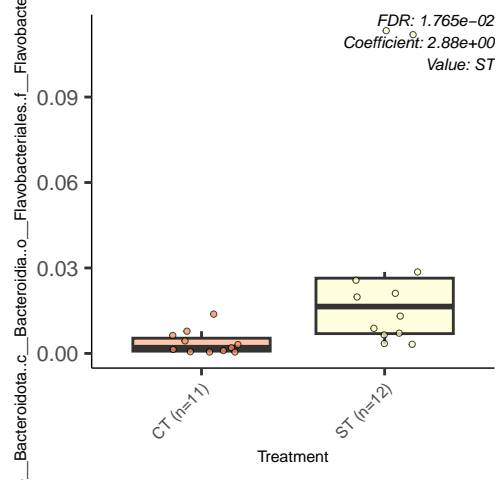


acteria..p__Acidobacteriota..c__Blastocatellia..o__UBA7656..f__UBA





proteobacteria.c__Gammaproteobacteria..o__Burkholderiales..f__Bu

FDR: 1.765e-02
Coefficient: 2.42e+00
Value: ST

0.010

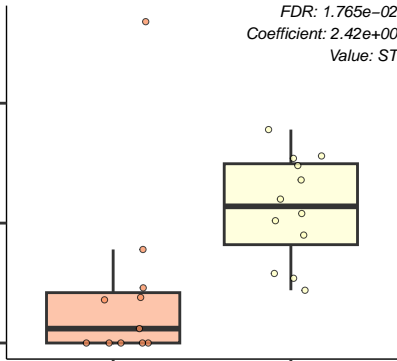
0.005

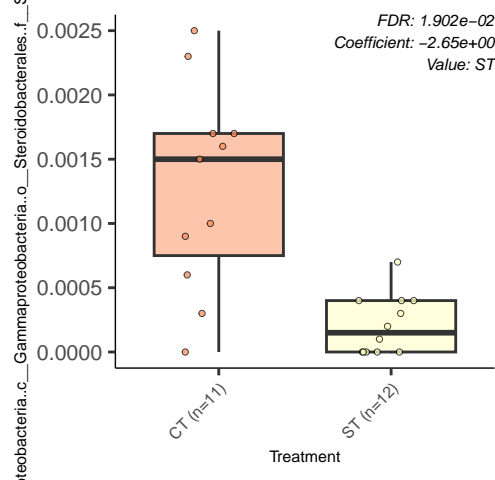
0.000

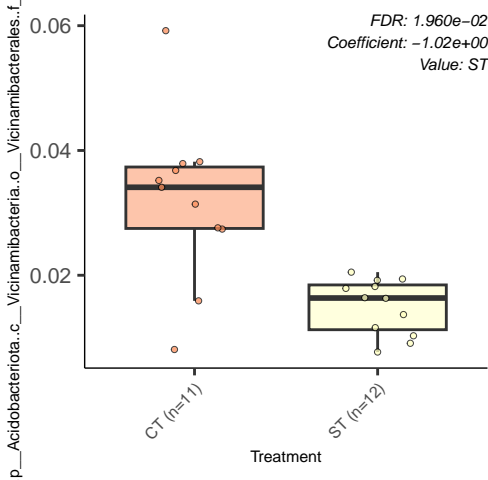
CT (n=11)

ST (n=12)

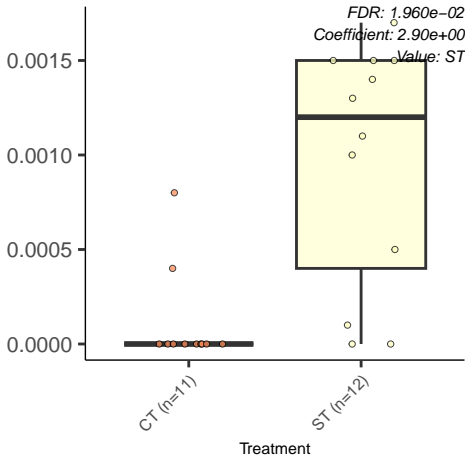
Treatment

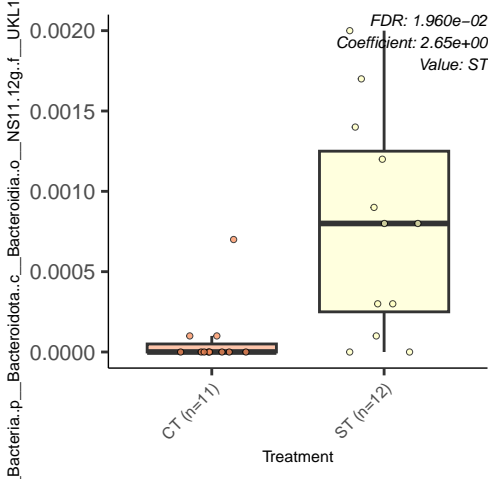


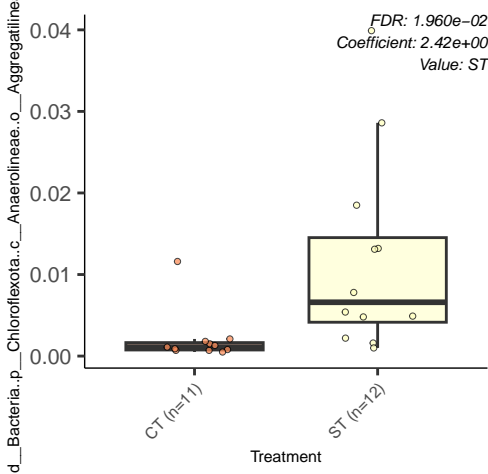


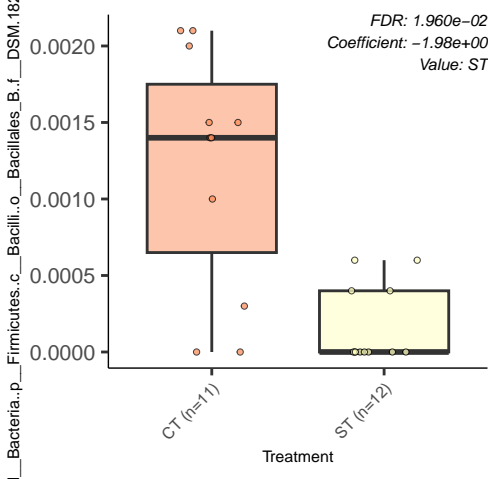


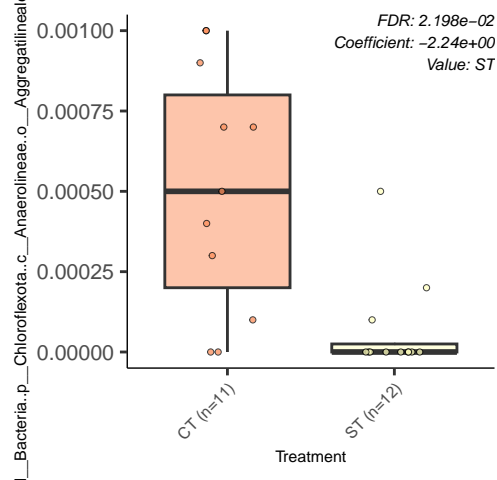
ia..p__Actinobacteriota..c__Actinomycetia..o__Propionibacteriales..f

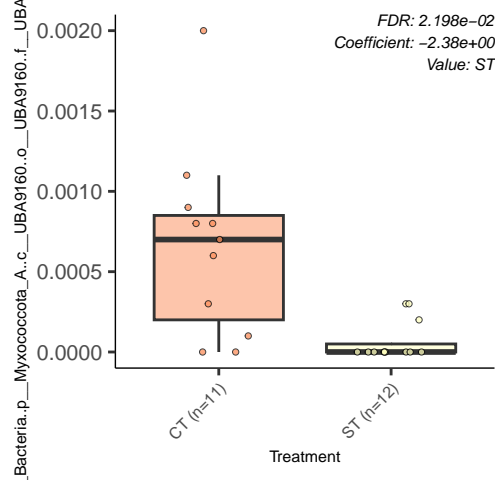


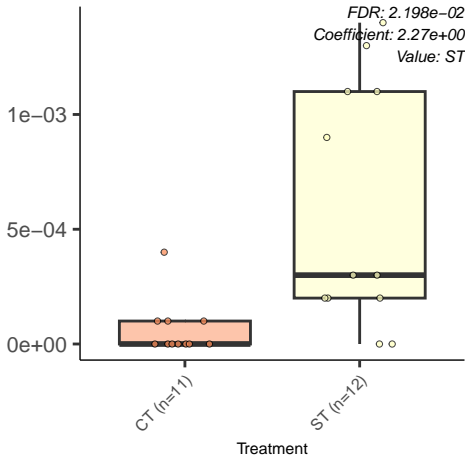


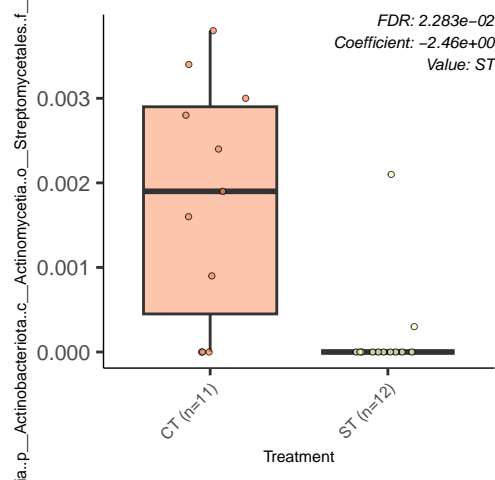


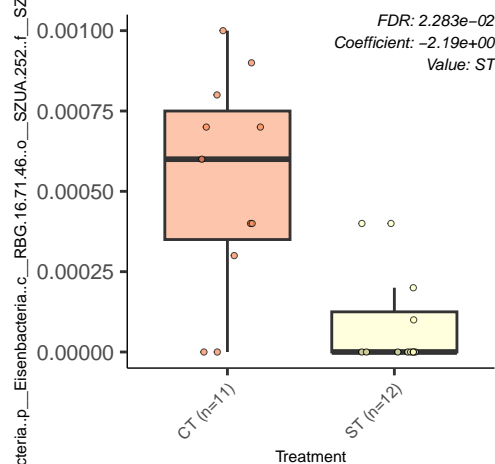


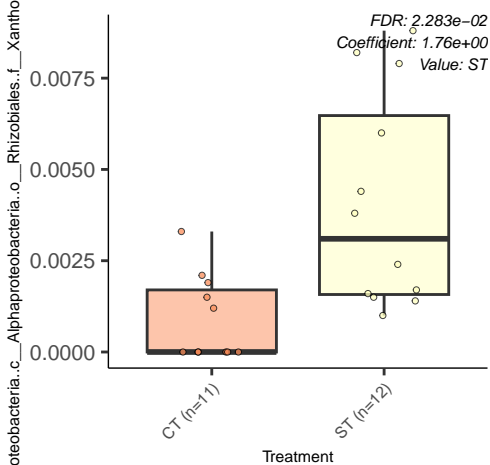


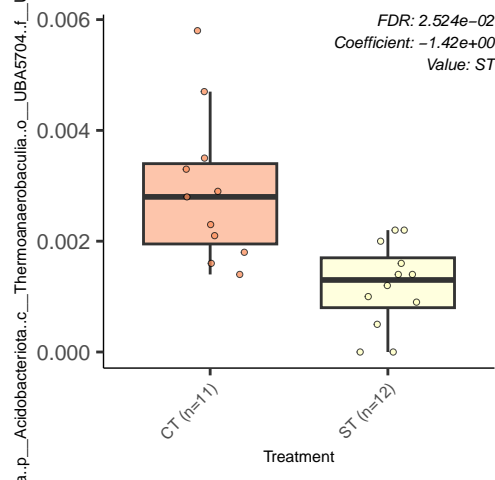


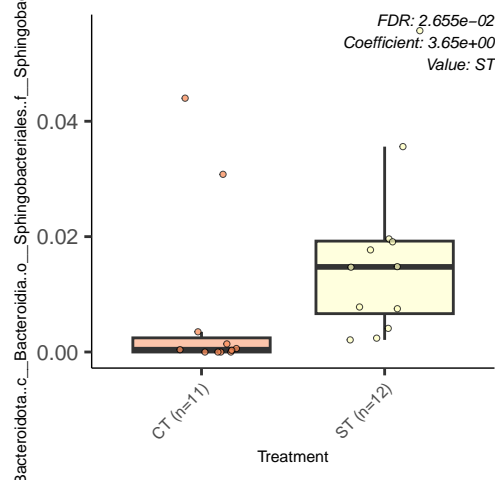


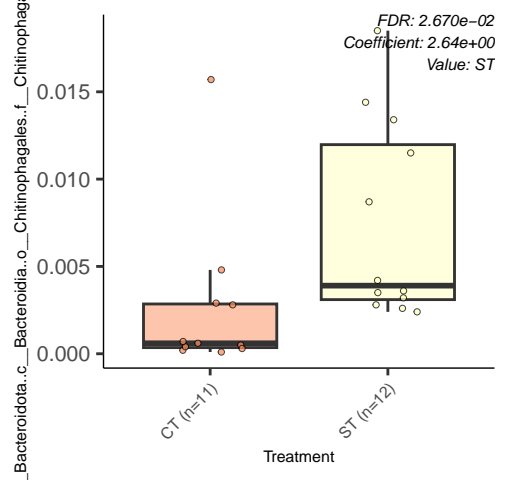


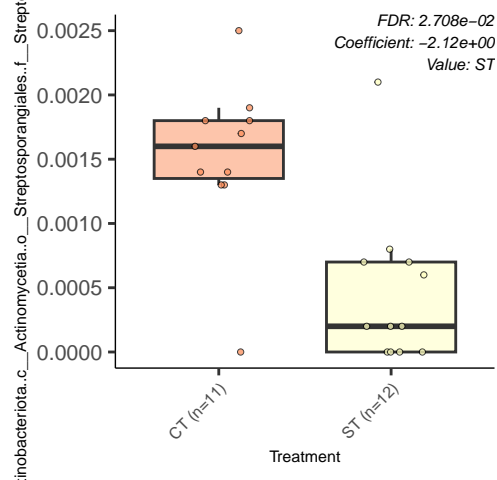


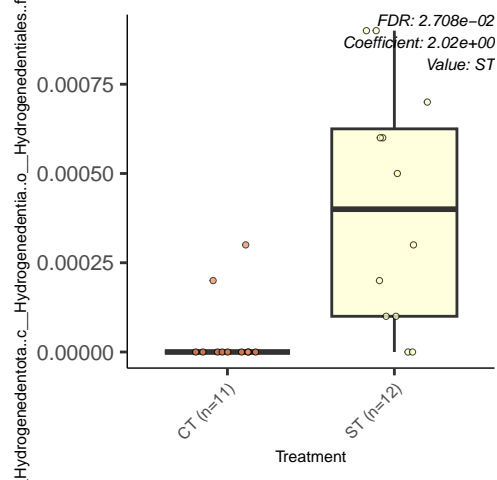


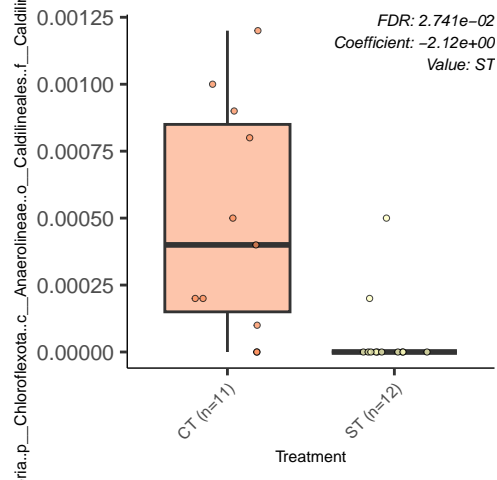


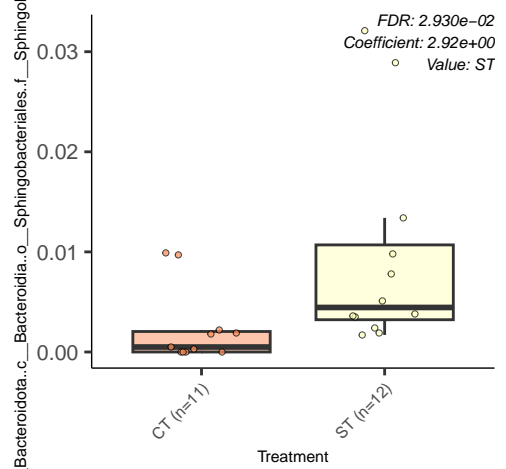


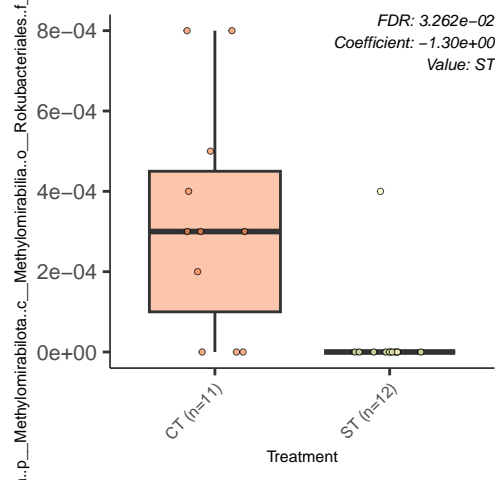


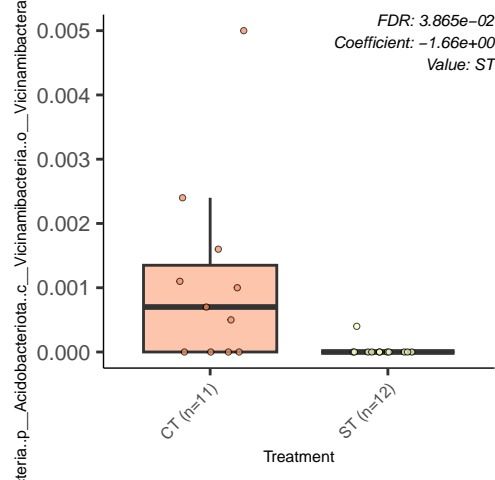






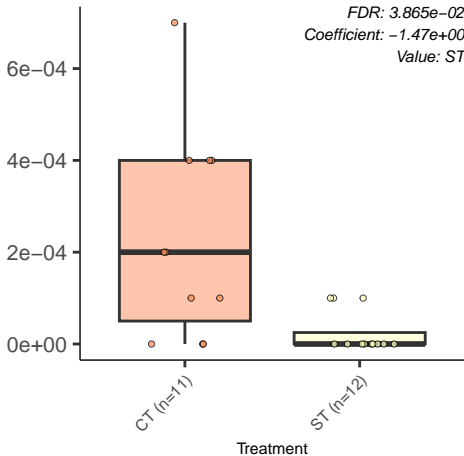


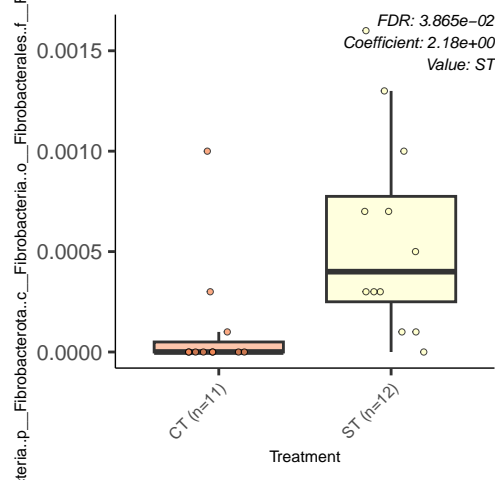


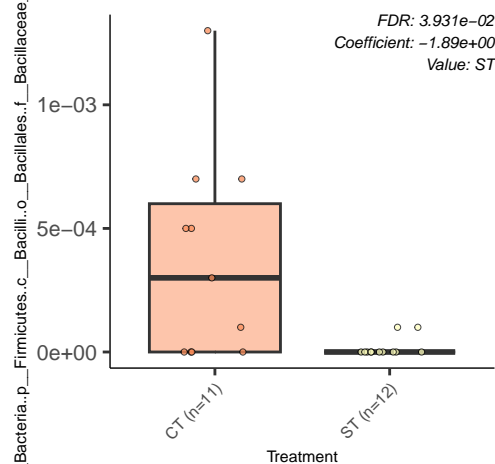


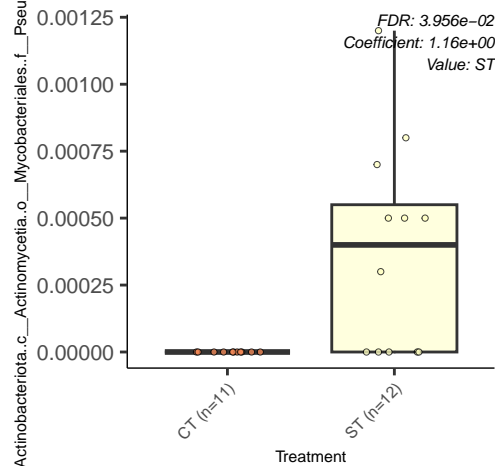
ctinobacteriota..c_Thermoleophilia..o_Miltoncostaeales..f_Milton

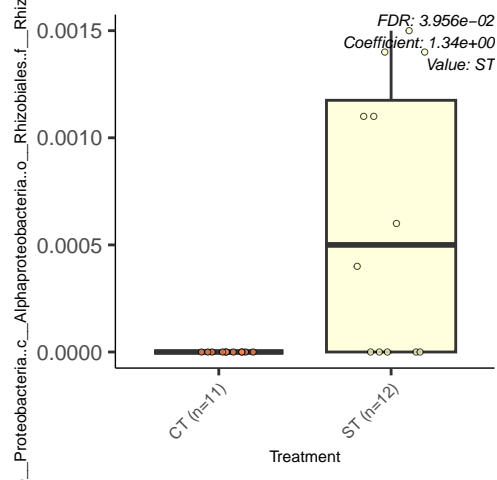
FDR: $3.865e-02$
Coefficient: $-1.47e+00$
Value: ST

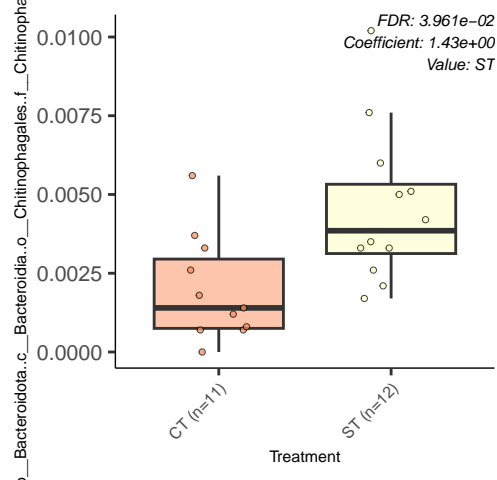


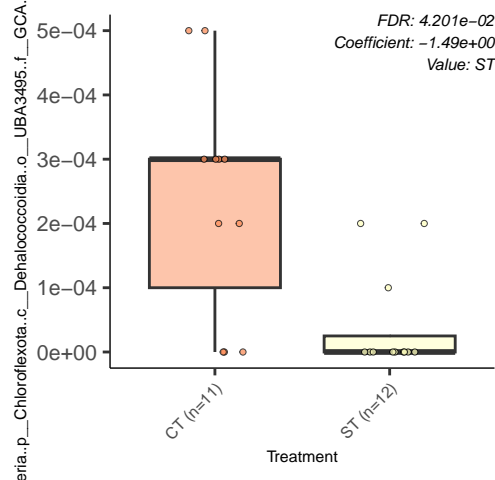


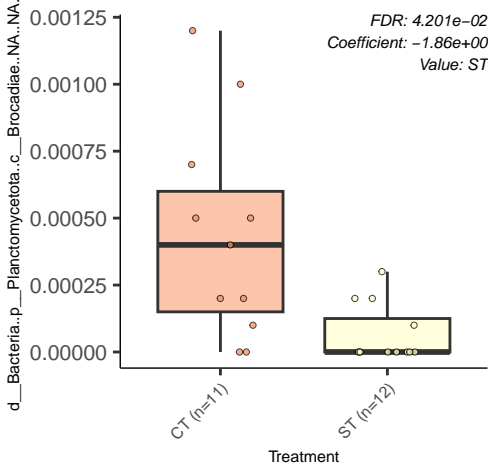


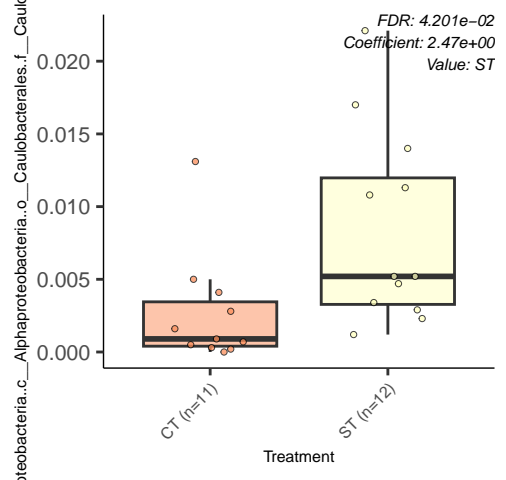




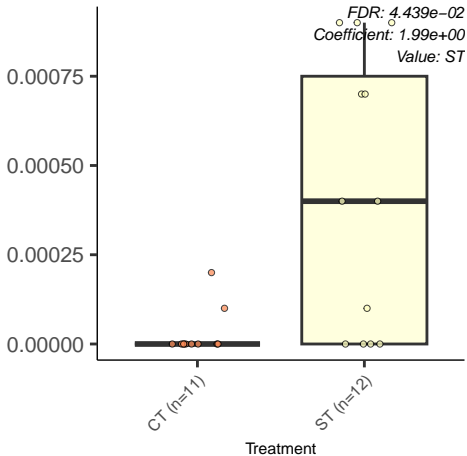


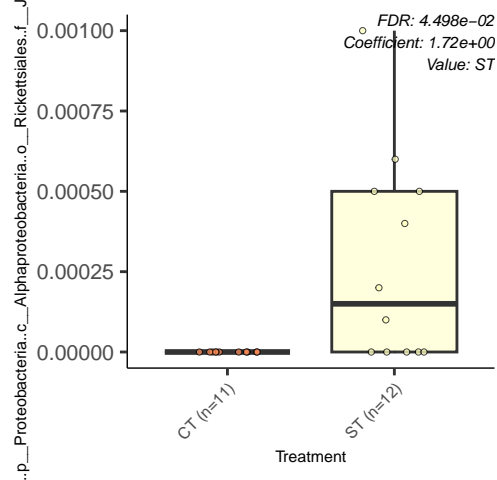






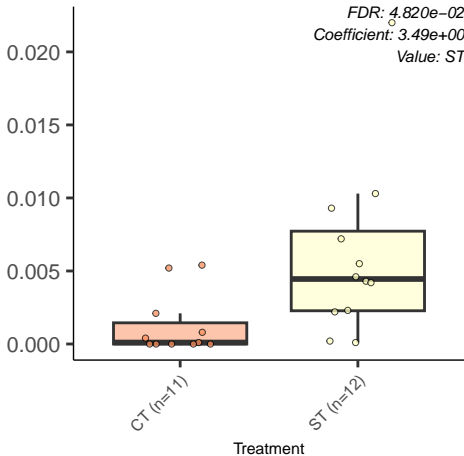
a..p__UBA10199..c__UBA10199..o__UBA10199..f__2.12.FULL.40.

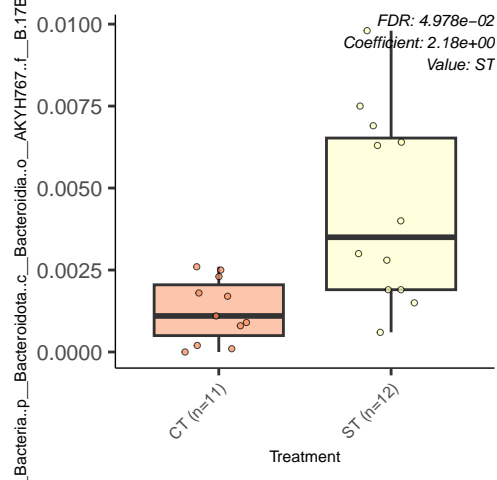


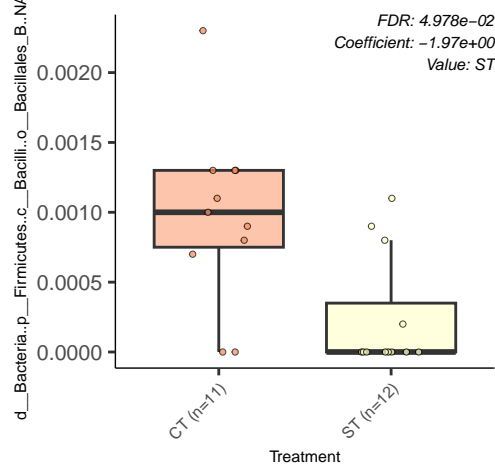


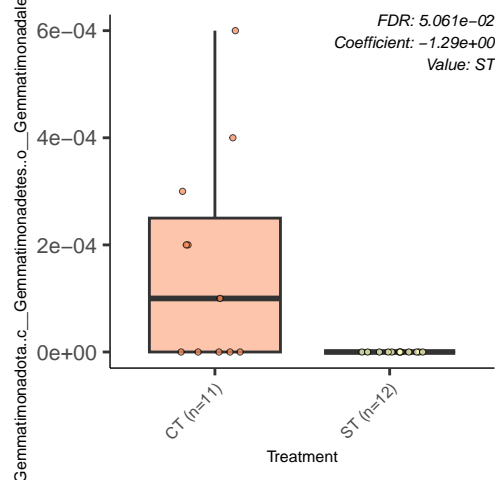
matimonadota.c__Fimbrimonadia..o__Fimbrimonadales.f__Fimbr

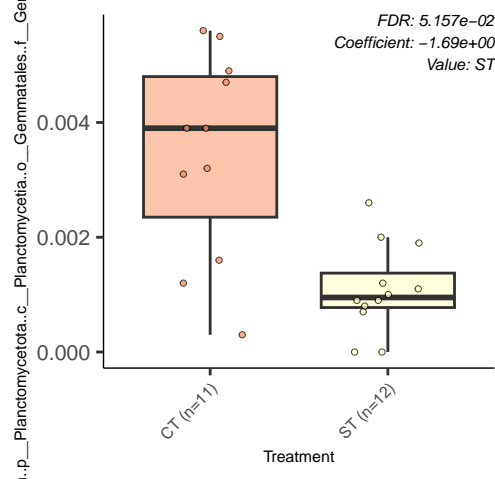
FDR: 4.820e-02
Coefficient: 3.49e+00
Value: ST

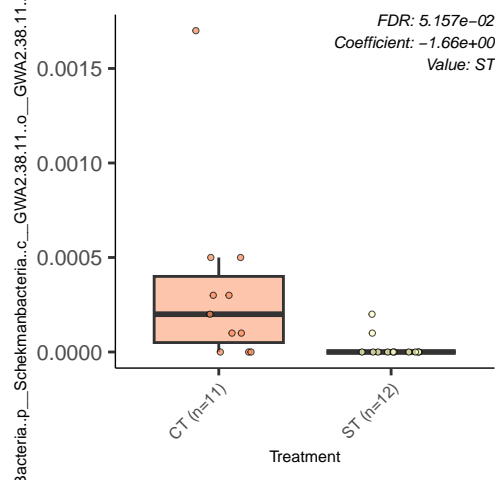












d__Bacteria.p__Bacteroidota.c__Bacteroidia.o__Chitinophaga

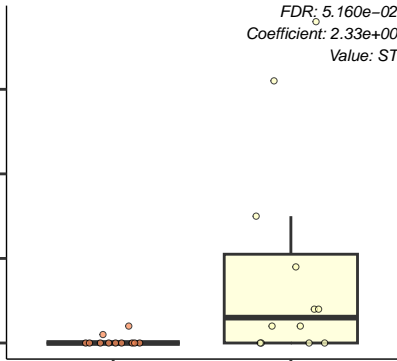
FDR: 5.160×10^{-2}
Coefficient: 2.33×10^0
Value: ST

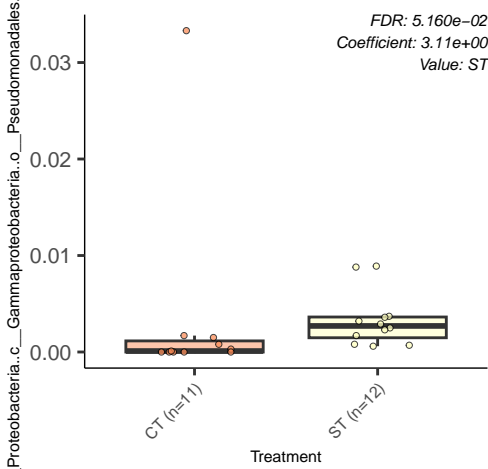
0.003
0.002
0.001
0.000

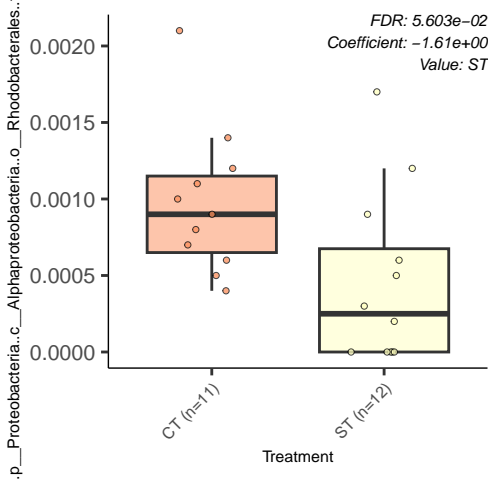
CT (n=11)

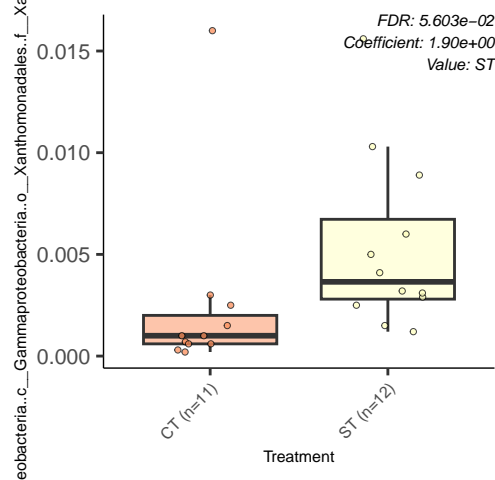
ST (n=12)

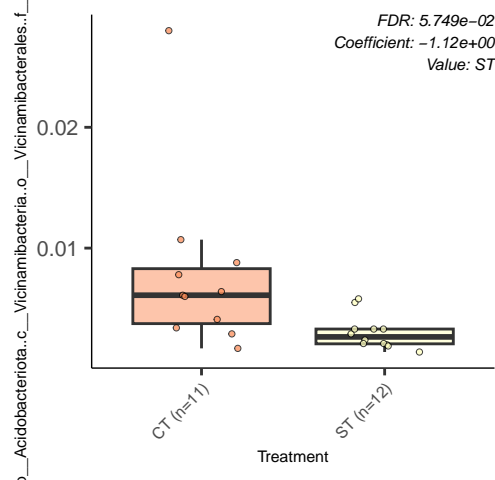
Treatment

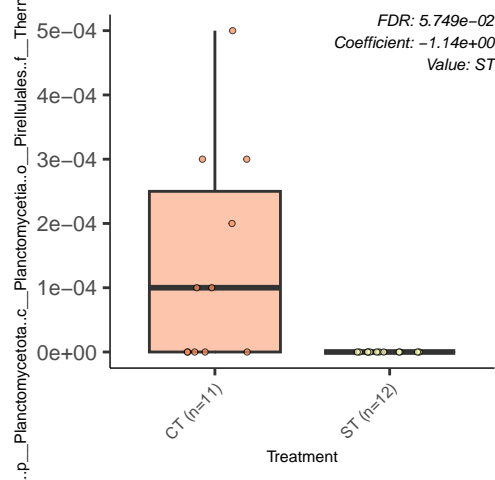


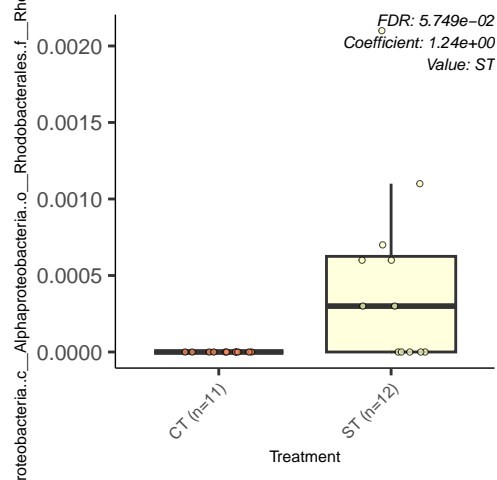


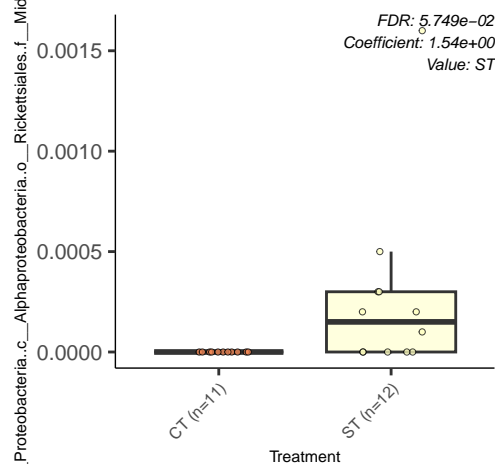


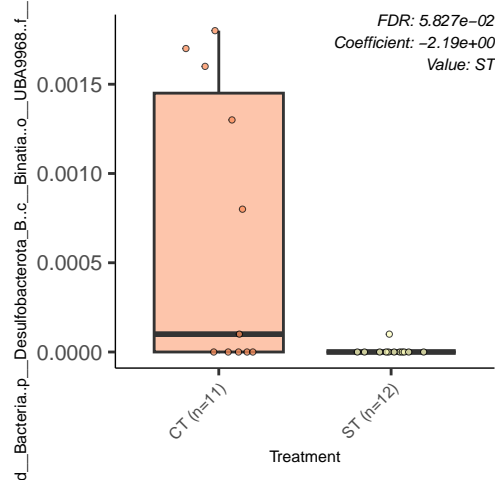


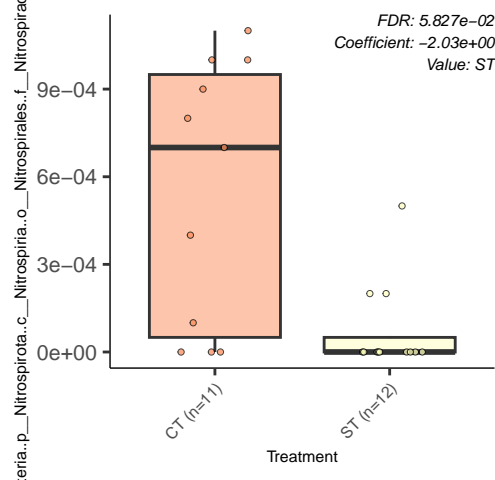


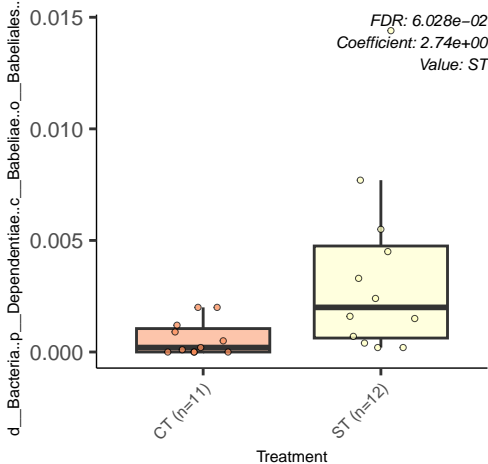


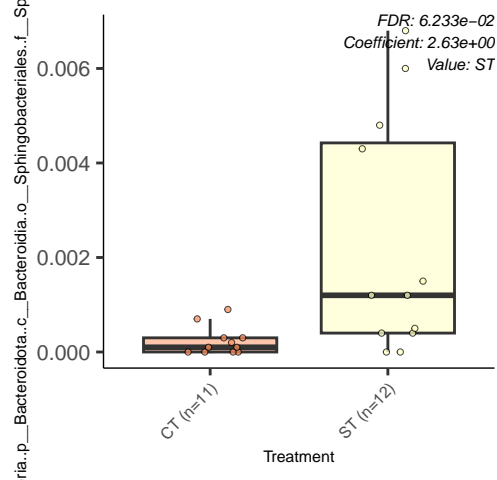


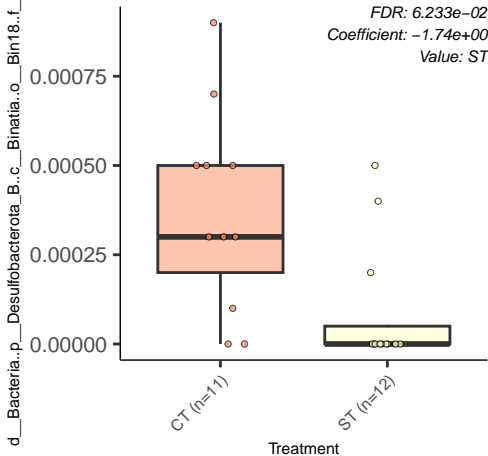


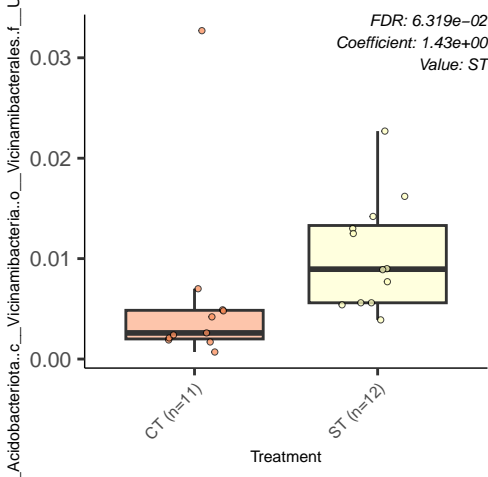


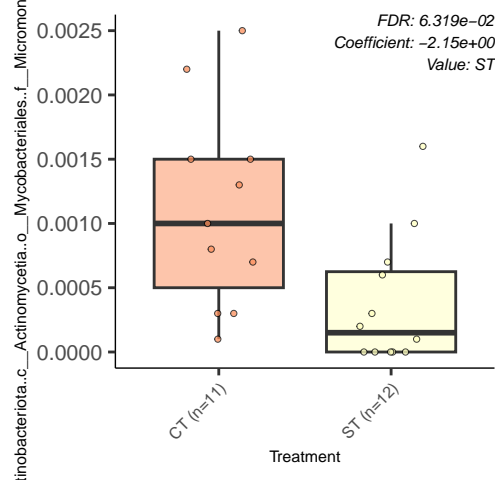


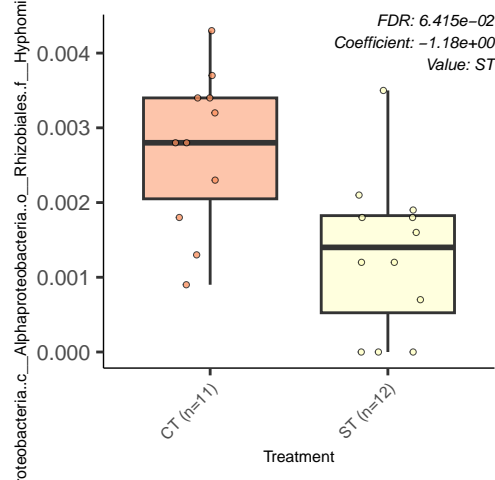


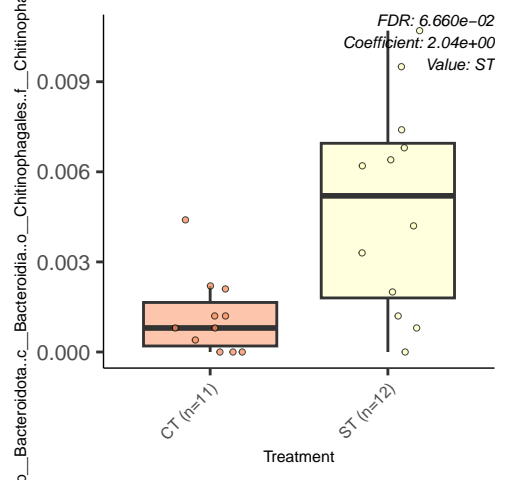


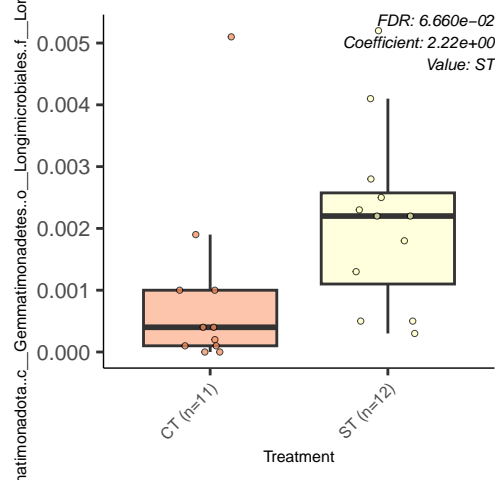


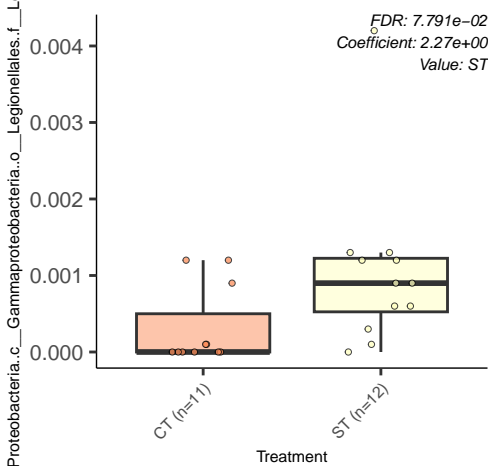


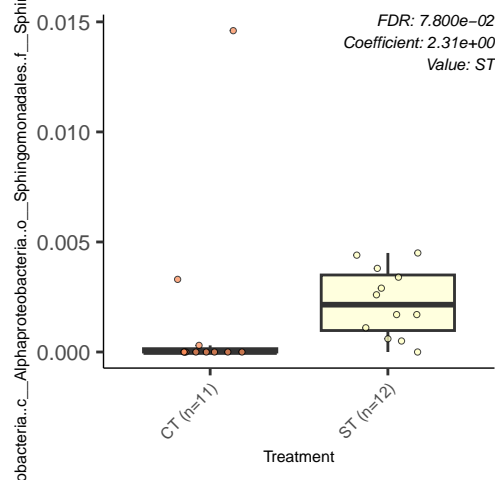






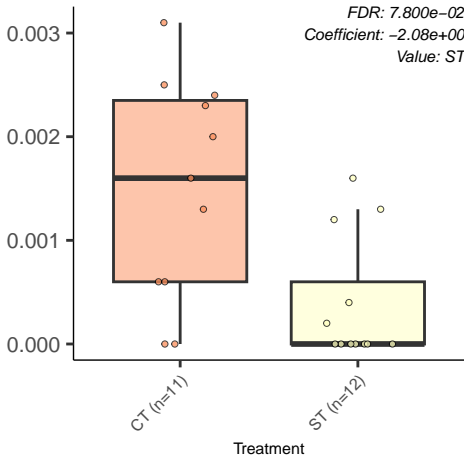






a.p__Verrucomicrobiota.c__Verrucomicrobiae.o__Pedosphaerales

FDR: $7.800e-02$
Coefficient: $-2.08e+00$
Value: ST



Bacteria..p__Chloroflexota..c__Anaerolineae..o__Aggregatilineales..f__

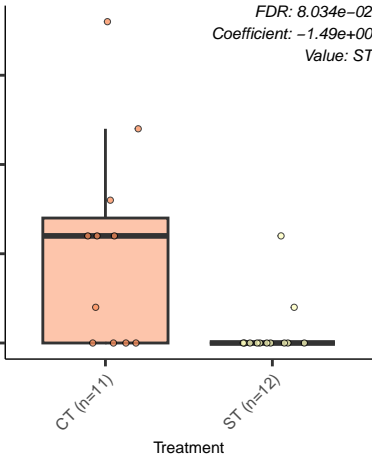
FDR: 8.034e-02
Coefficient: -1.49e+00
Value: ST

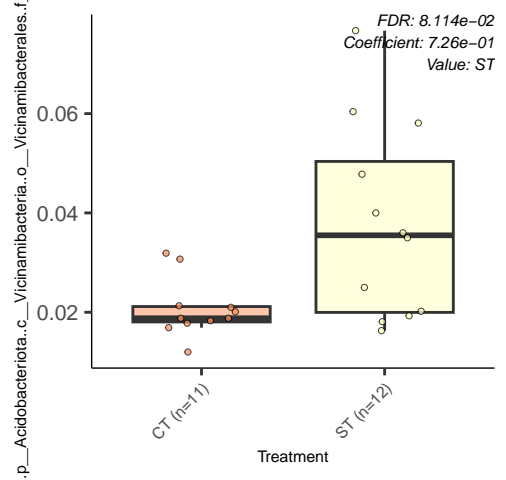
0.00075
0.00050
0.00025
0.00000

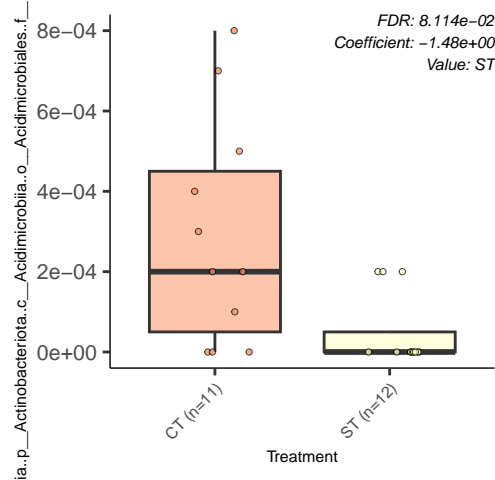
CT (n=11)

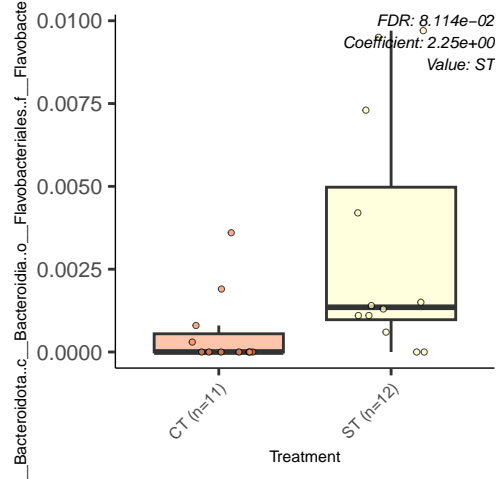
ST (n=12)

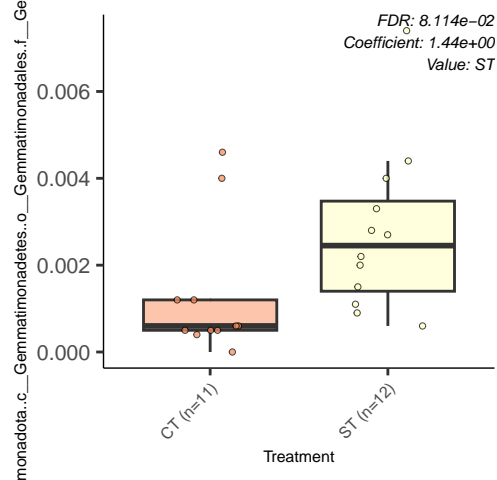
Treatment

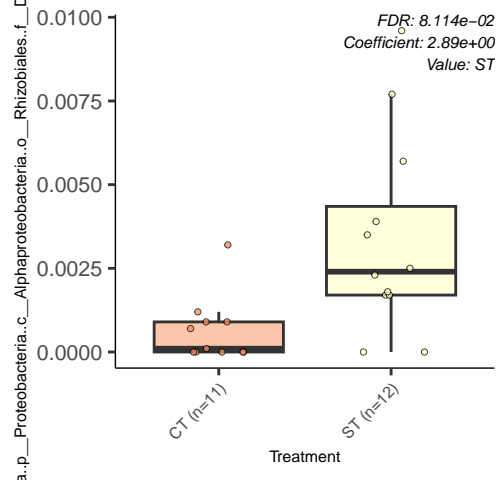


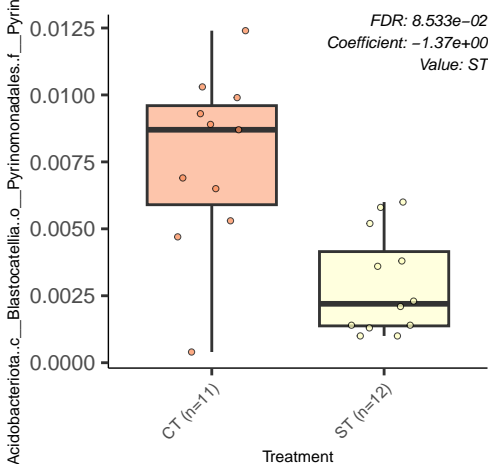


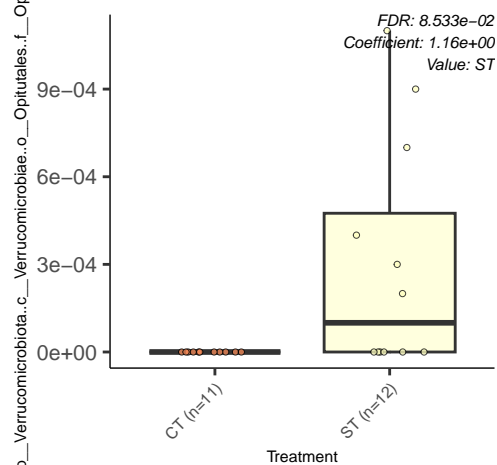


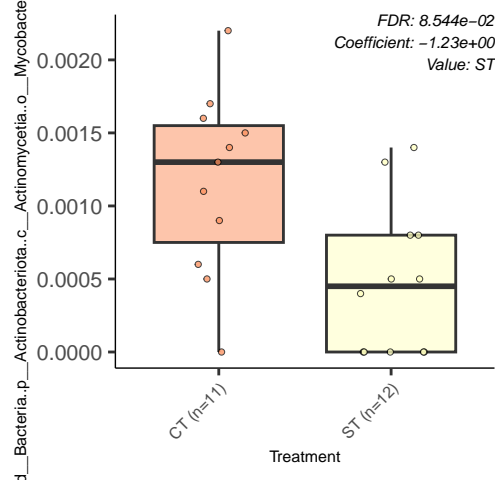


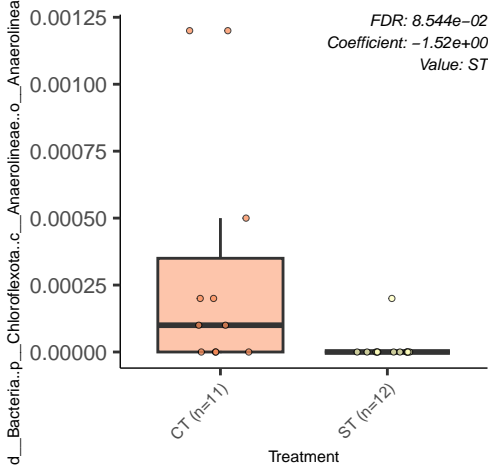


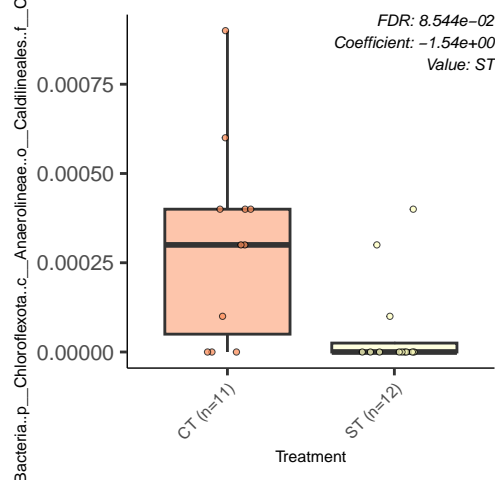


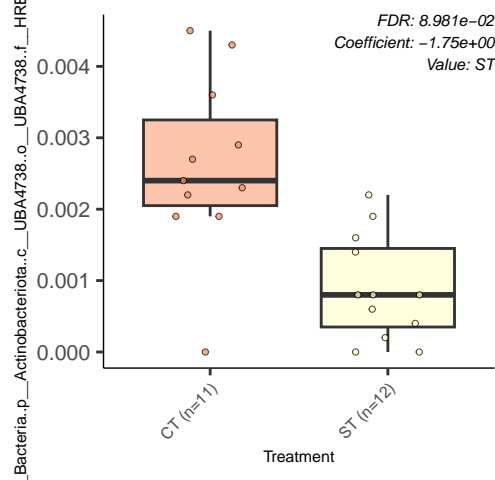


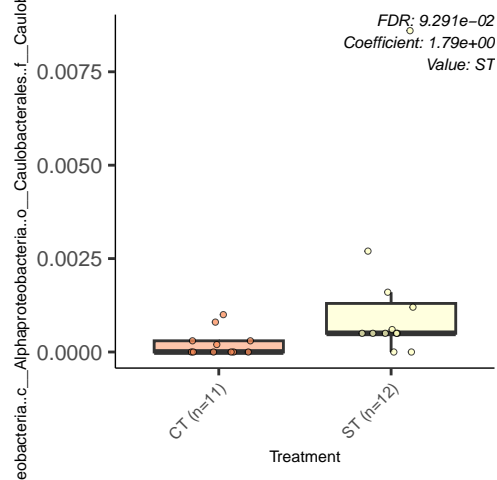




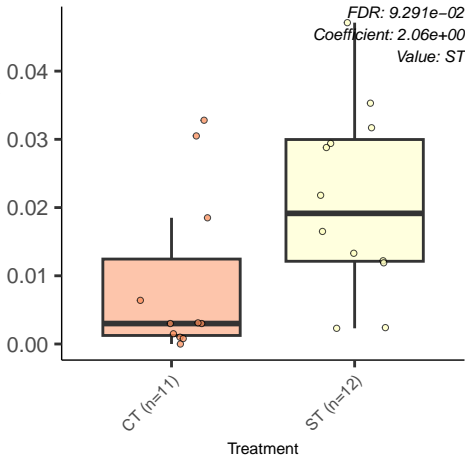








__Proteobacteria.c__Alphaproteobacteria.o__Sphingomonadales.



ia.p__Proteobacteria.c__Gammaproteobacteria.o__Legionellales

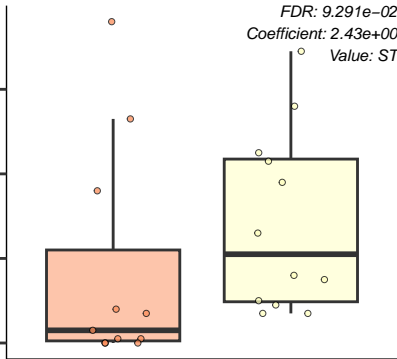
FDR: 9.291e-02
Coefficient: 2.43e+00
Value: ST

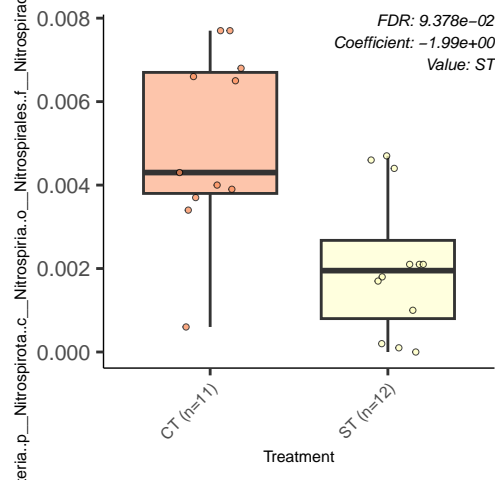
0.006
0.004
0.002
0.000

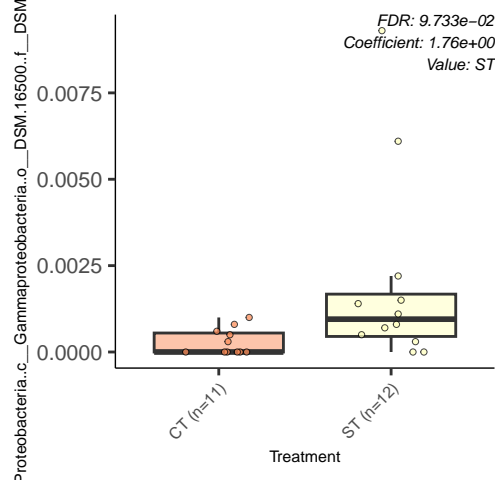
CT (n=11)

ST (n=12)

Treatment







Acidobacteriota..c__Vicinamibacteria..o__Vicinamibacterales..f__2.

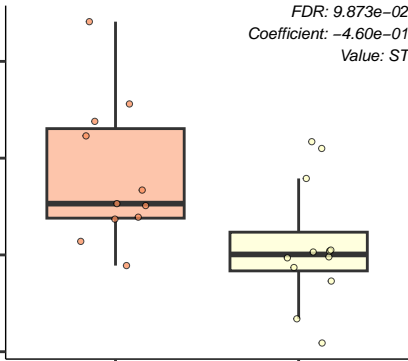
FDR: $9.873e-02$
Coefficient: $-4.60e-01$
Value: ST

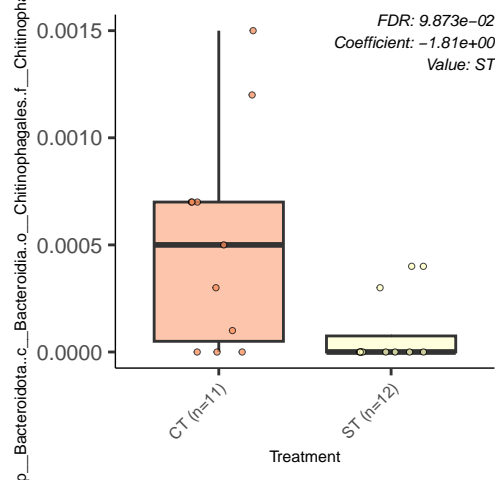
CT (n=11)

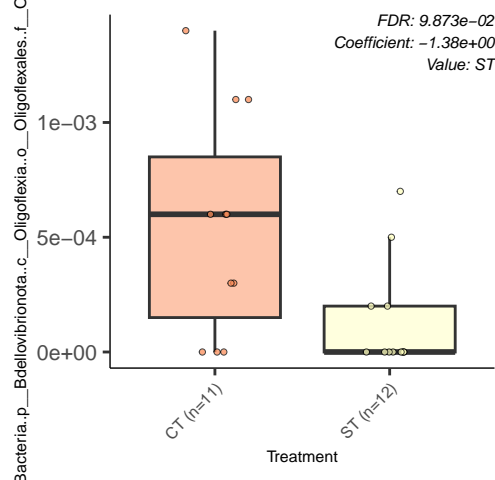
ST (n=12)

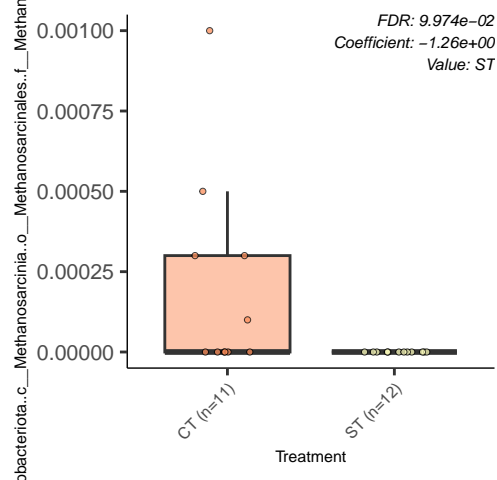
Treatment

0.01
0.02
0.03
0.04

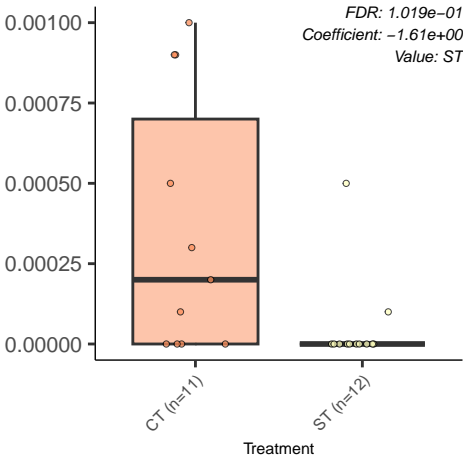


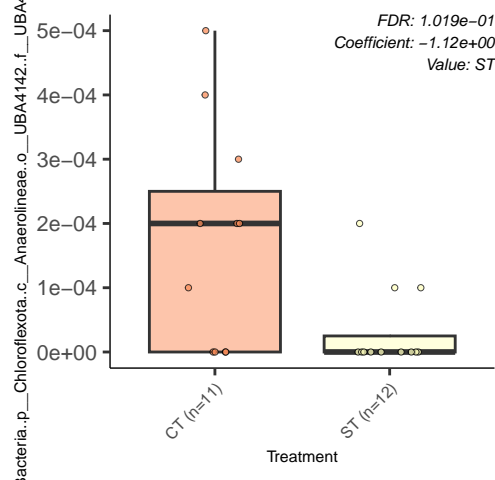


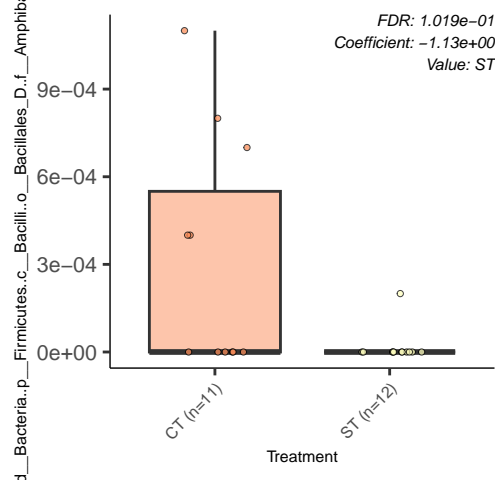


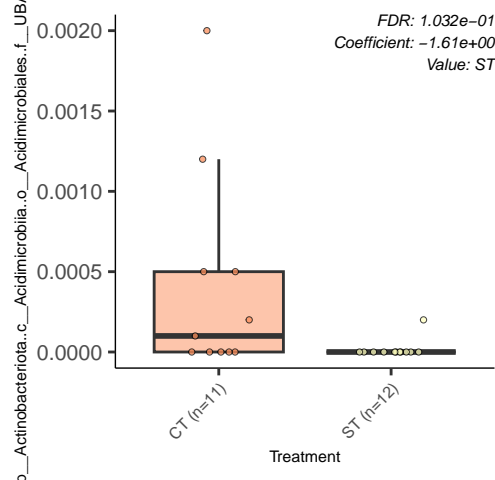


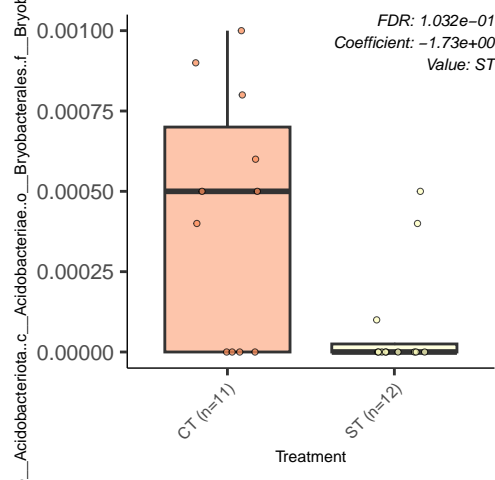
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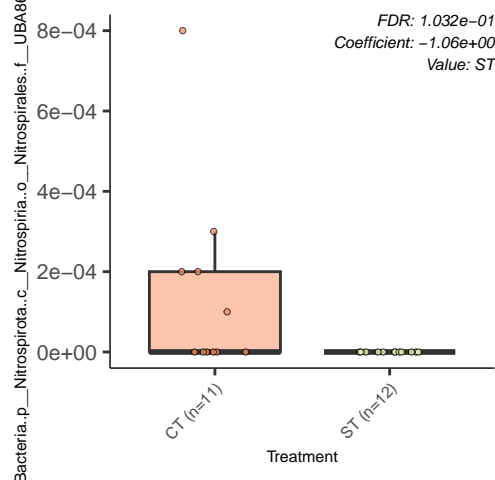


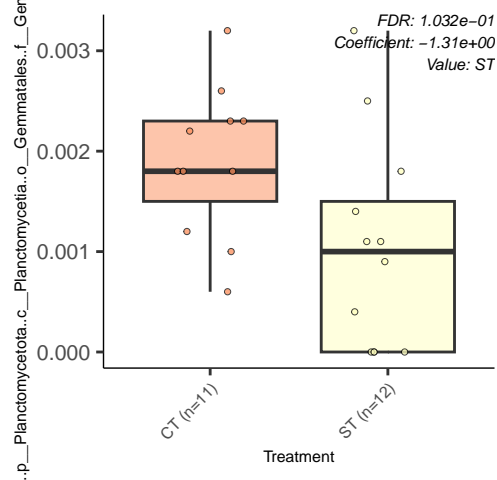


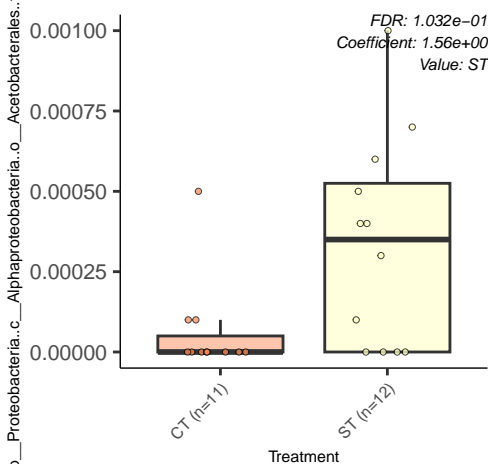


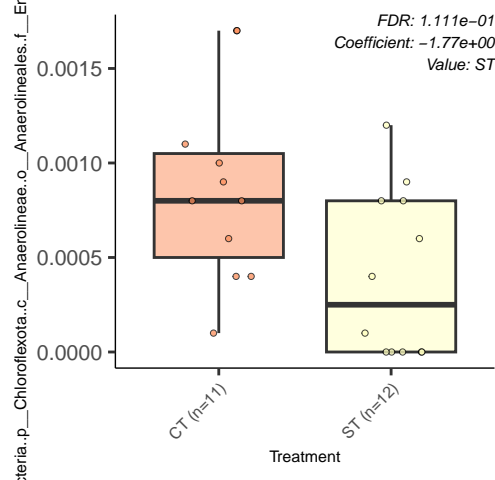


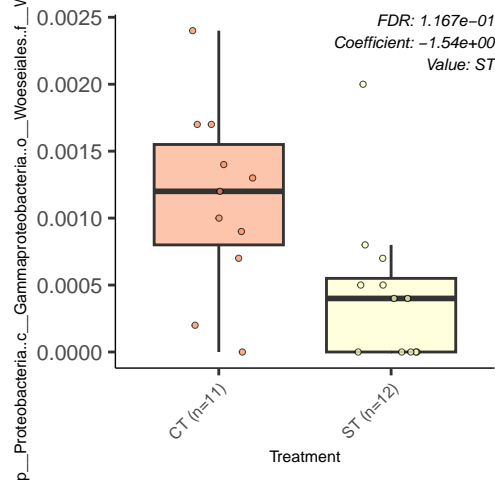


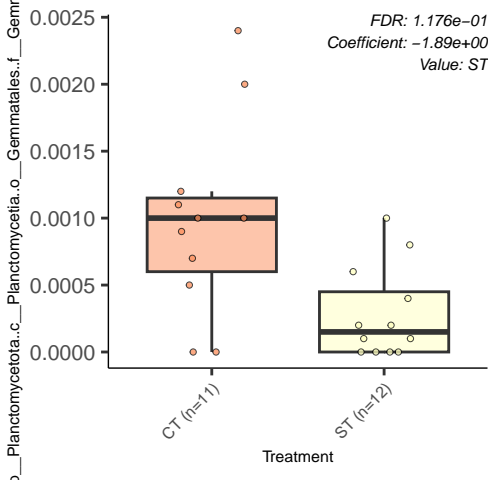


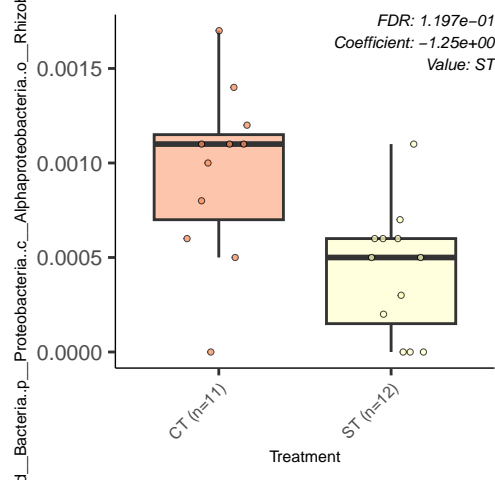


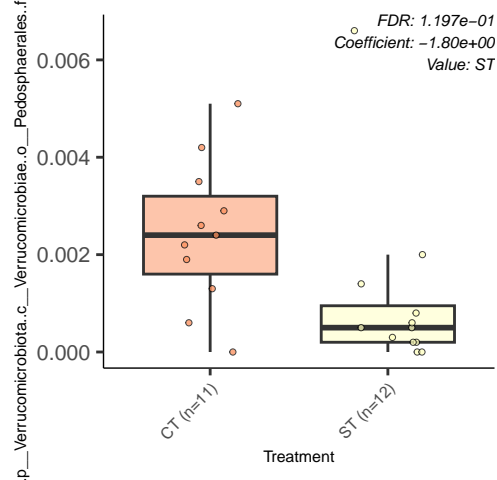


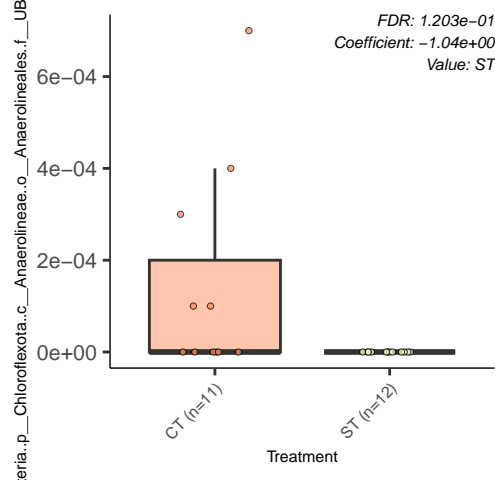


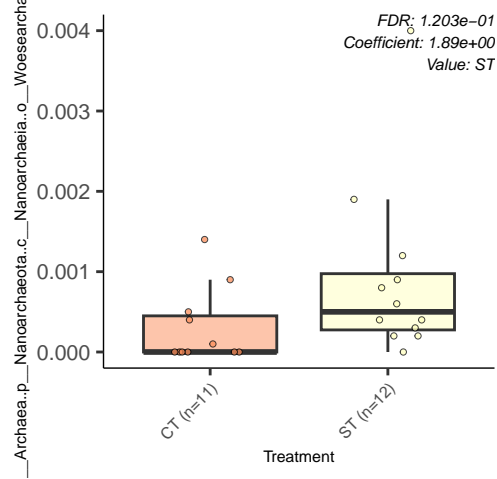


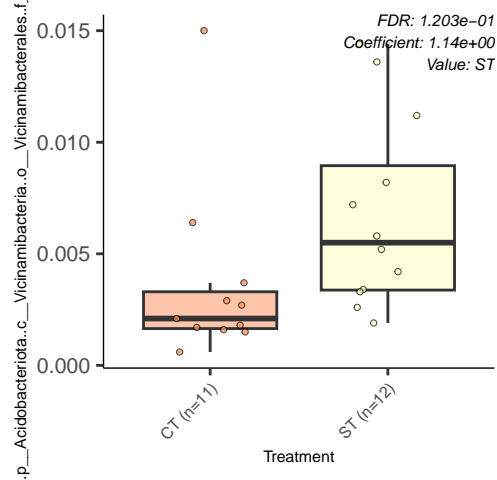


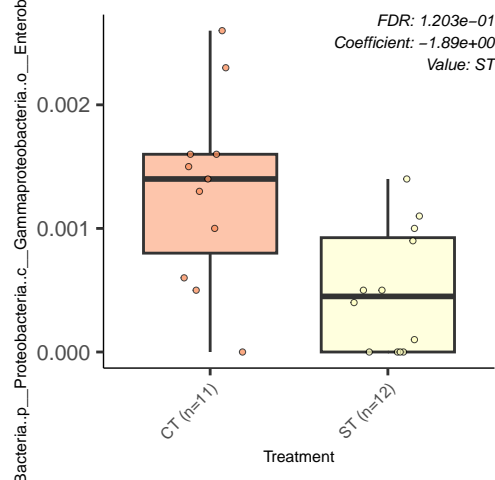


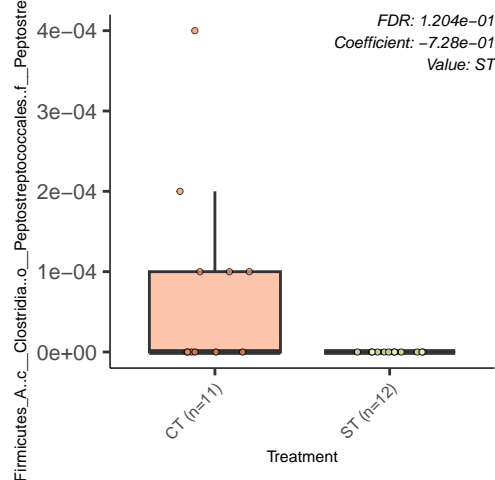


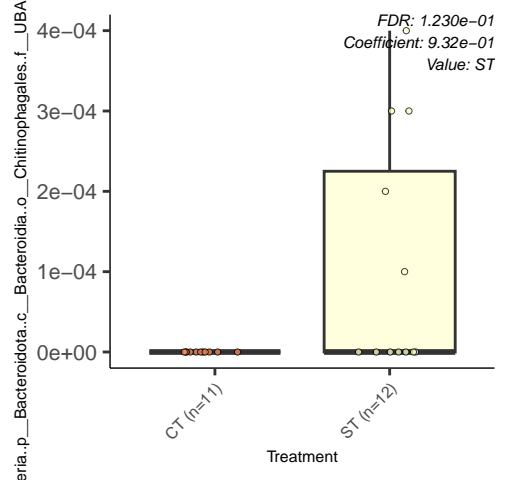


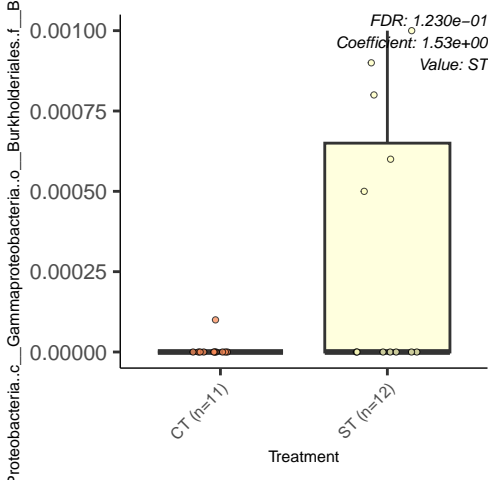


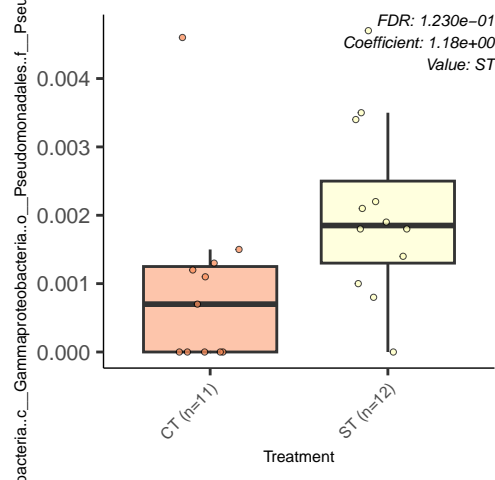


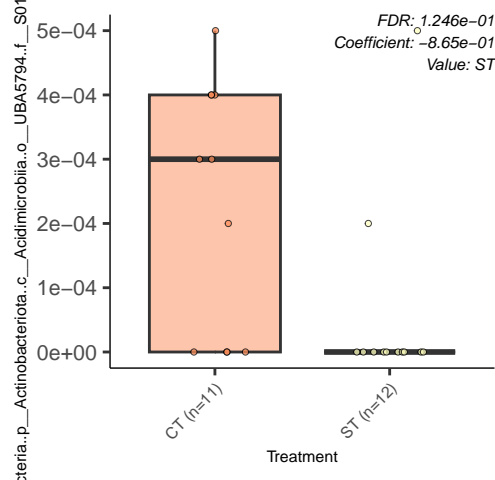


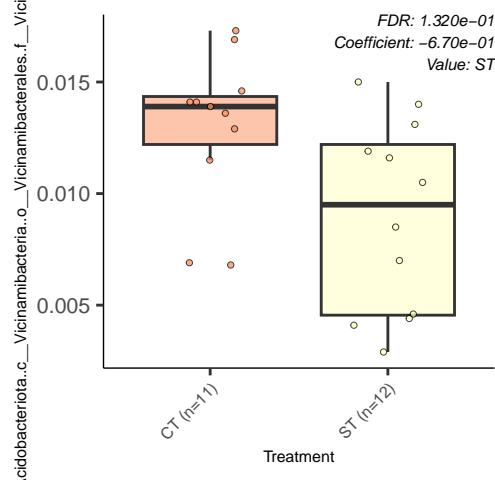


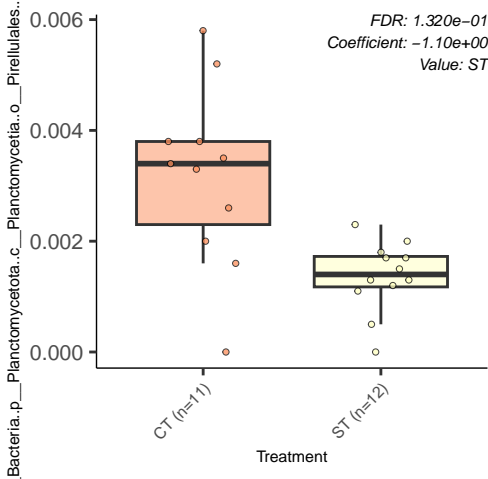


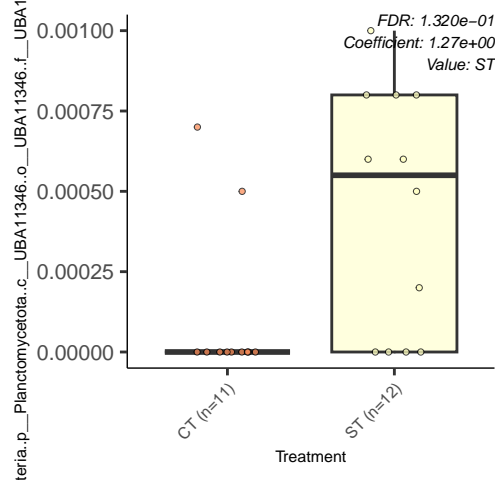






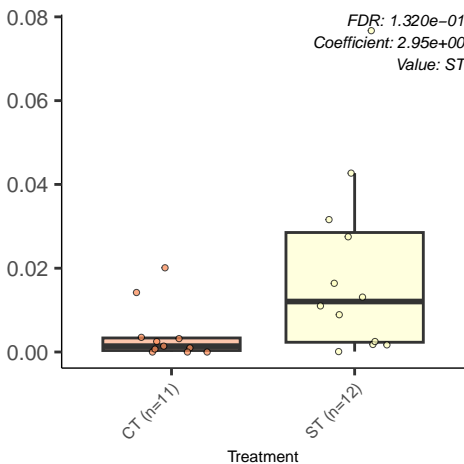


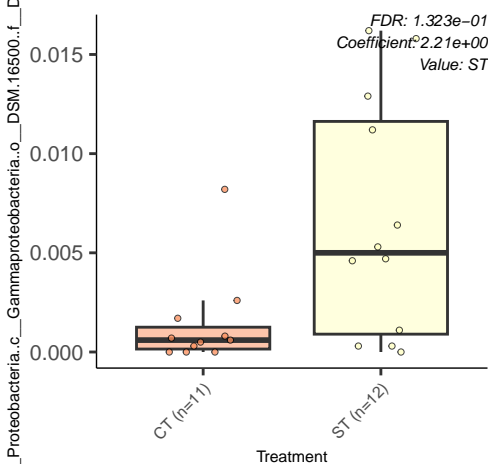


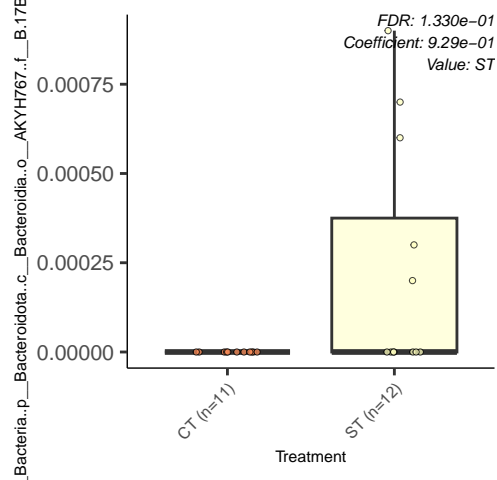


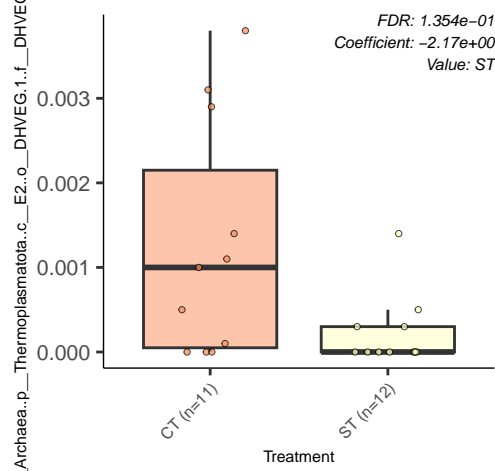
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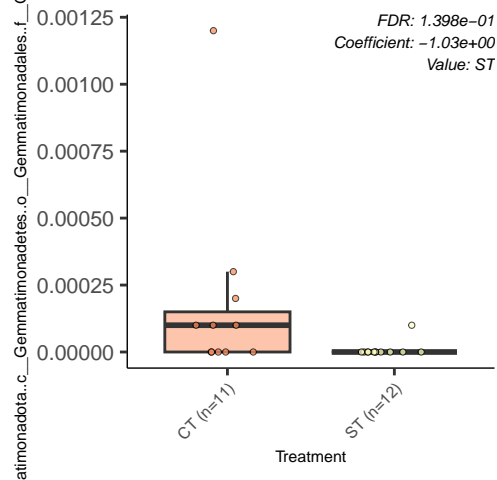
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Coefficient: 2.95e+00
Value: ST

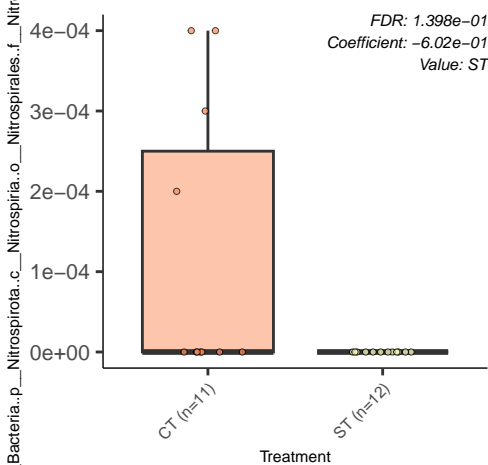


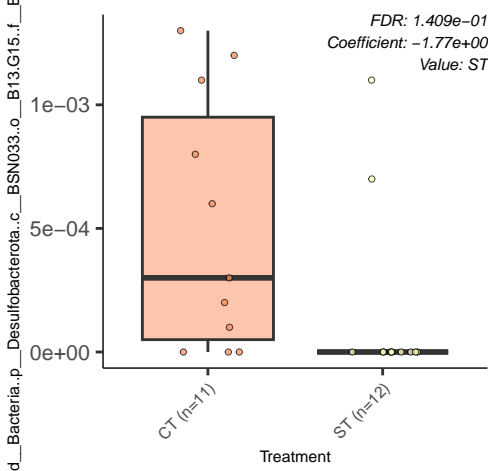


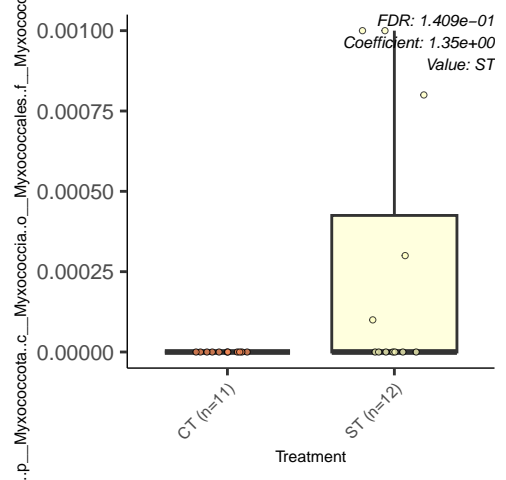












terial.p_Actinobacteriota.c_Acidimicrobiia.o_UBA5794.f_ZC4

FDR: 1.414e-01
Coefficient: -9.64e-01
Value: ST

0.006

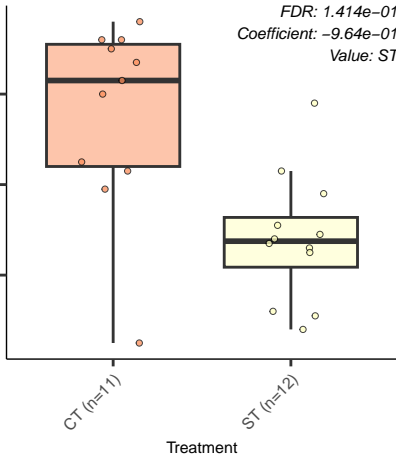
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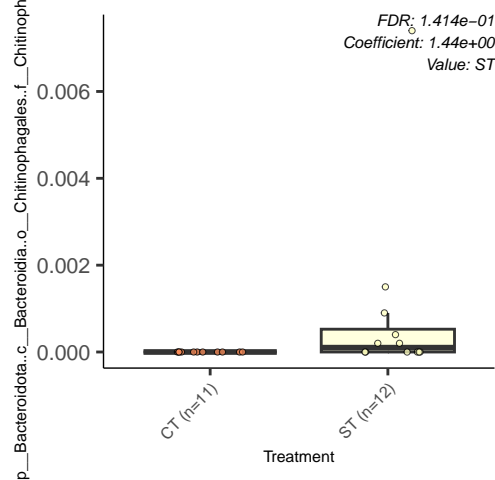
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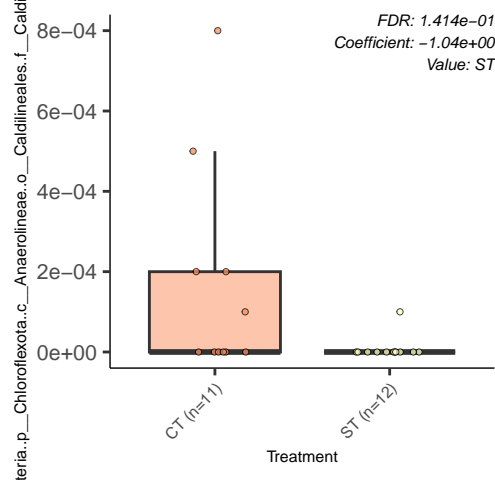
CT (n=11)

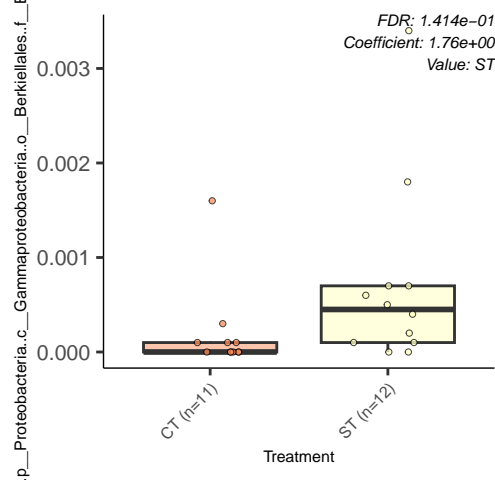
ST (n=12)

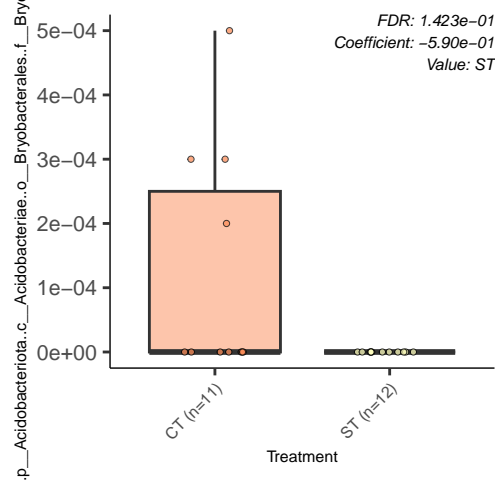
Treatment

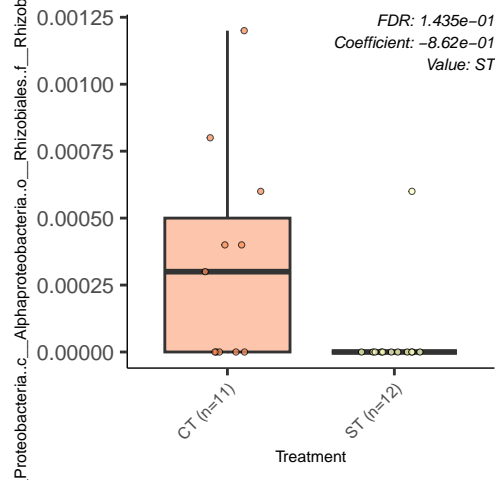


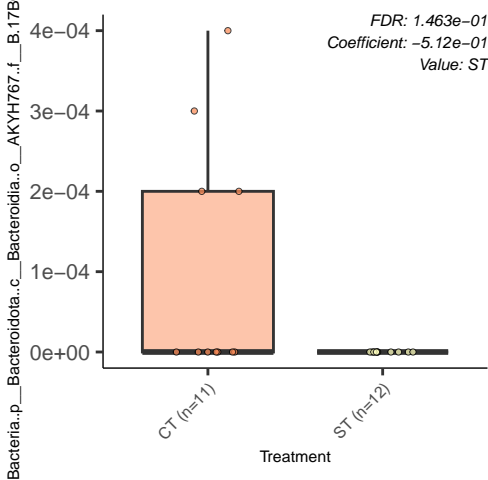


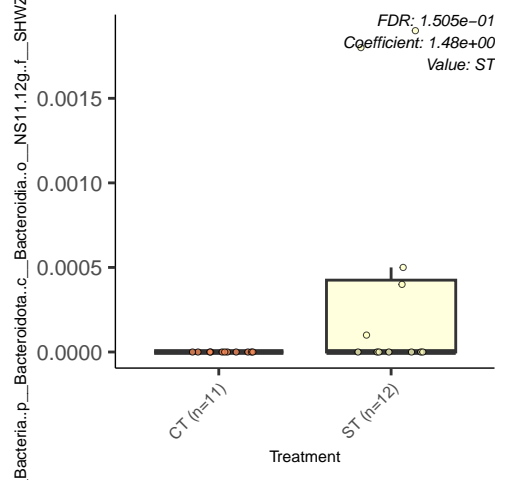


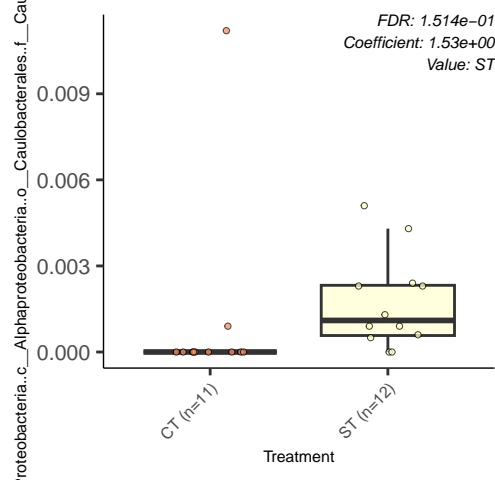


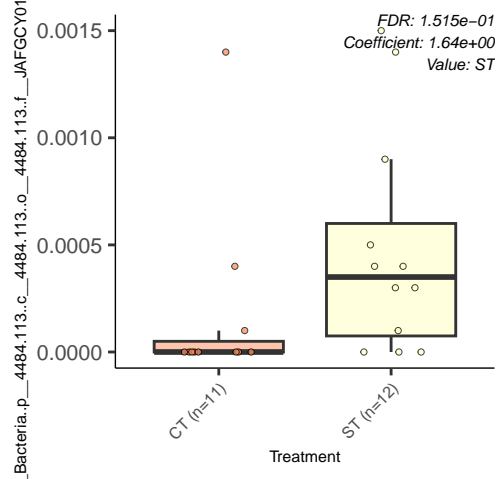


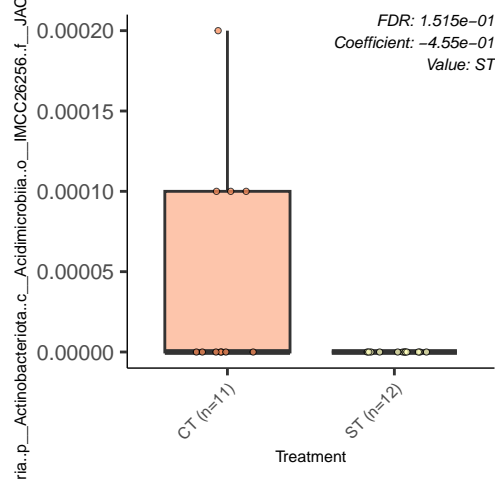


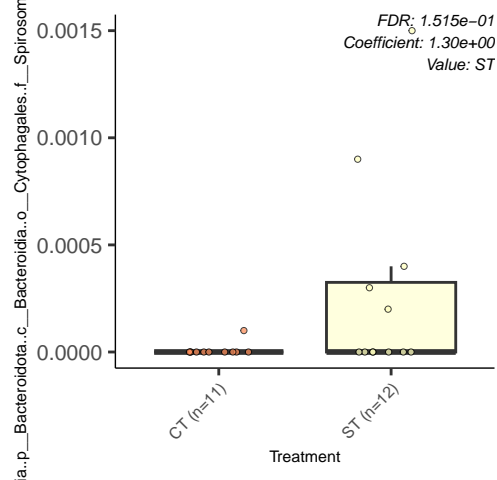


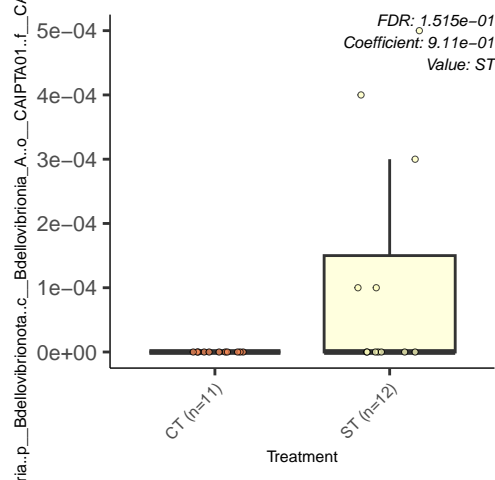


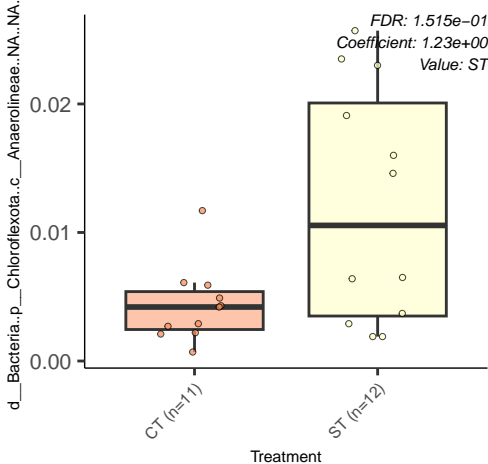


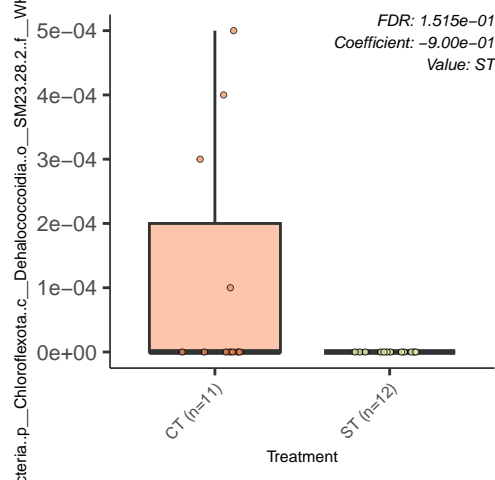


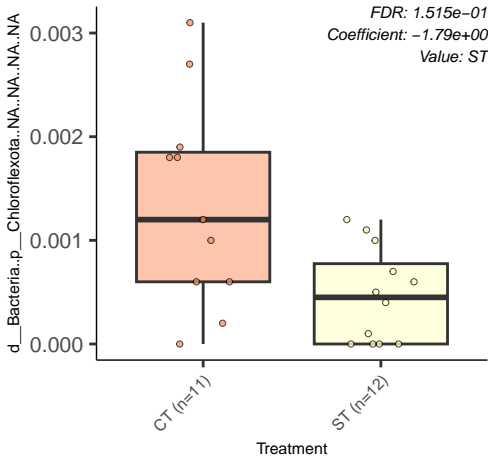


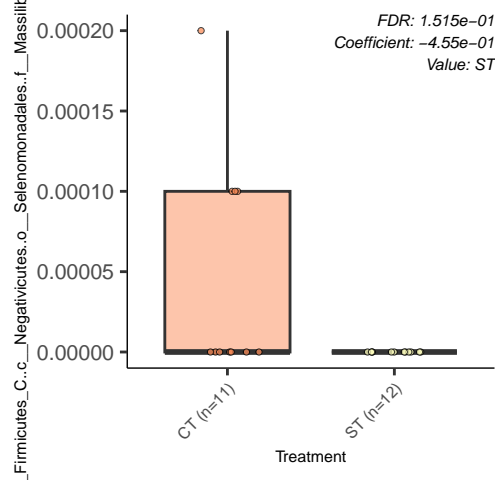


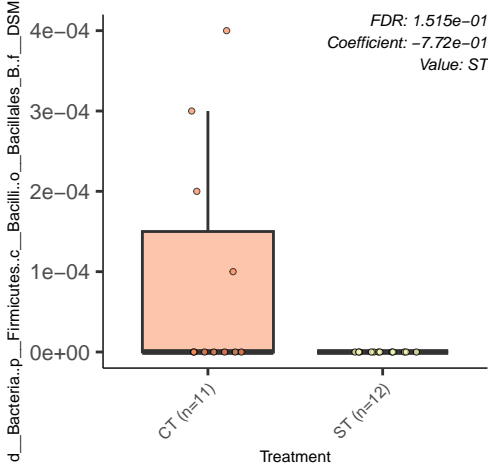


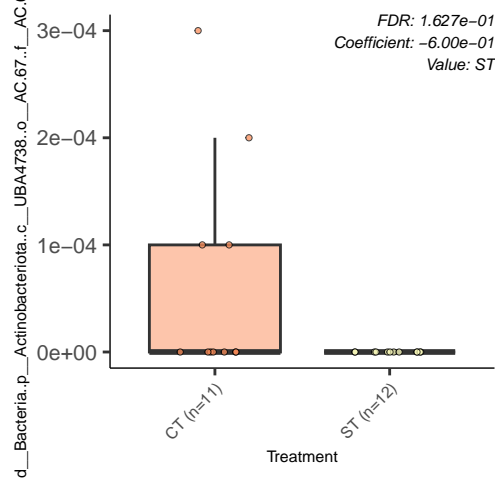






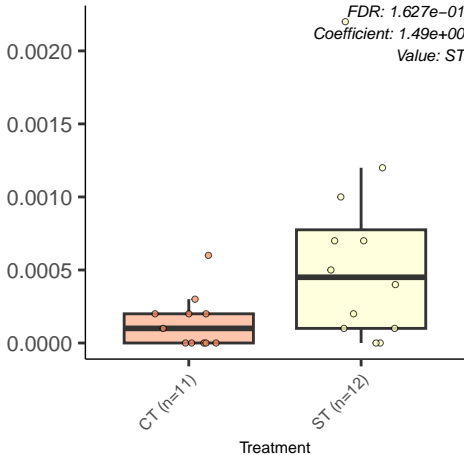


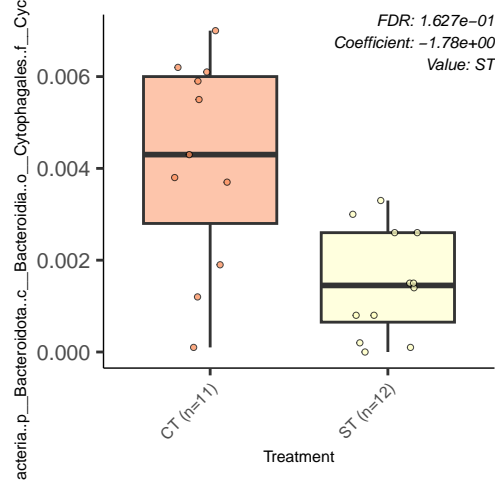


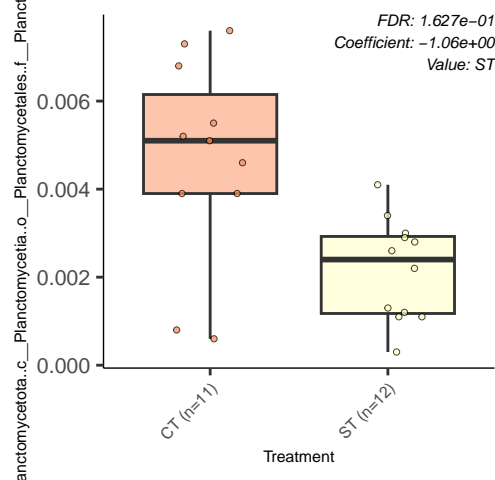


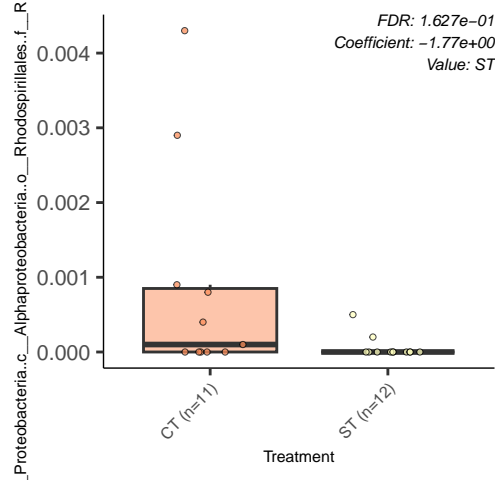
Bacteria.p__Bacteroidota..c__Bacteroidia.o__CAILMK01..f__CAILMK01

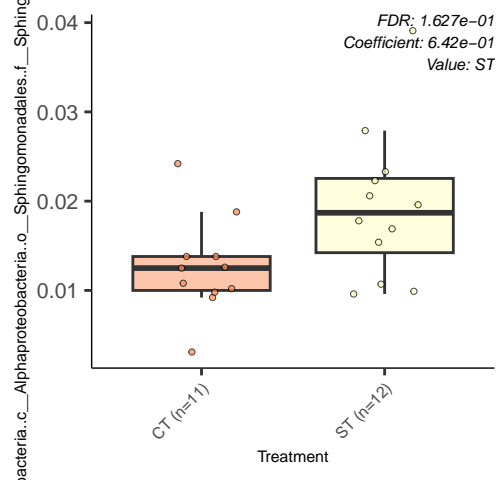
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Coefficient: 1.49e+00
Value: ST



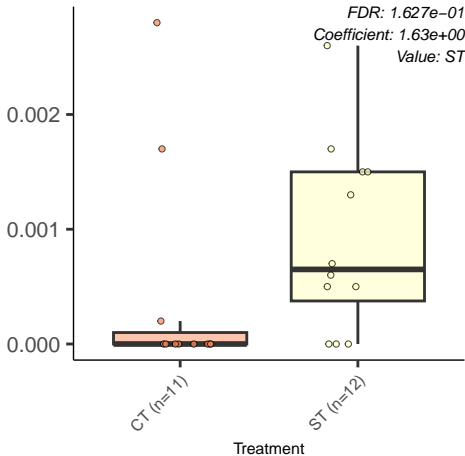


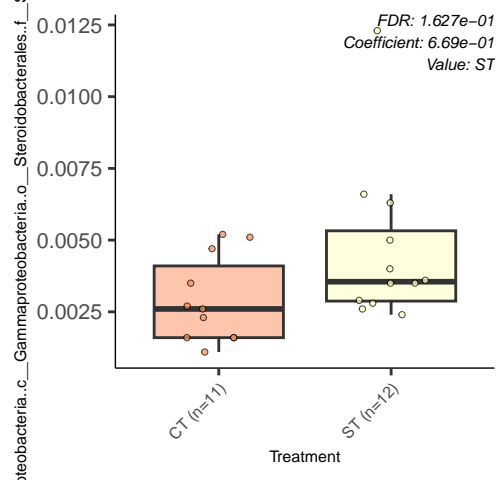


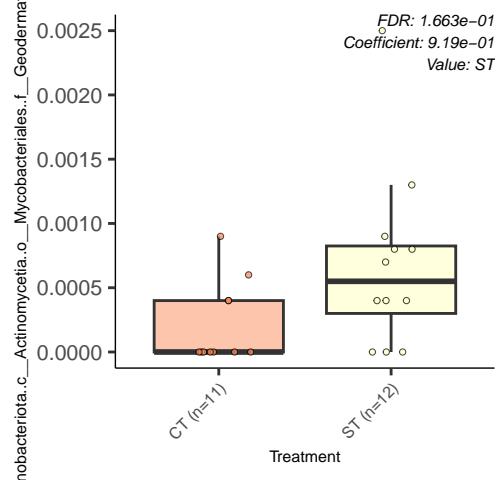


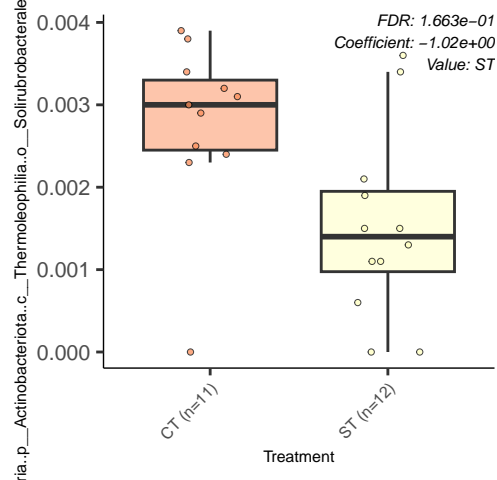


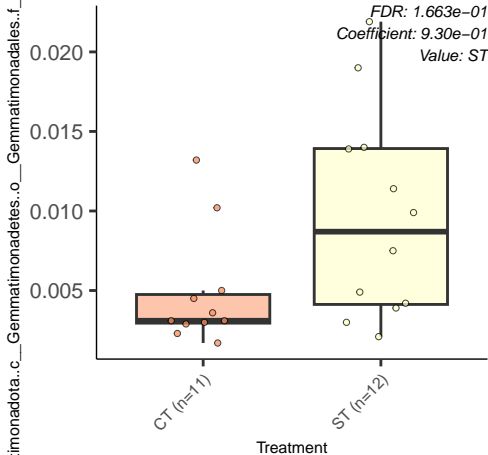
proteobacteria.c__Gammaproteobacteria.o__Burkholderiales.f__Bu

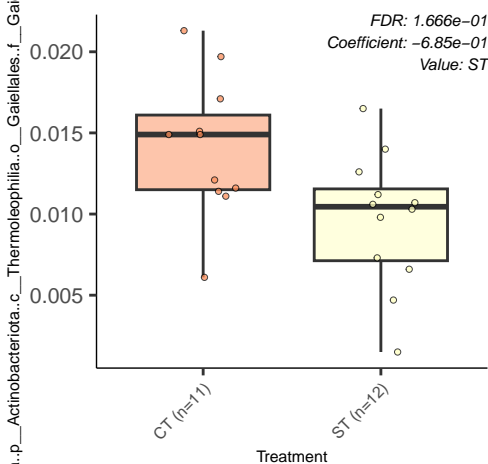


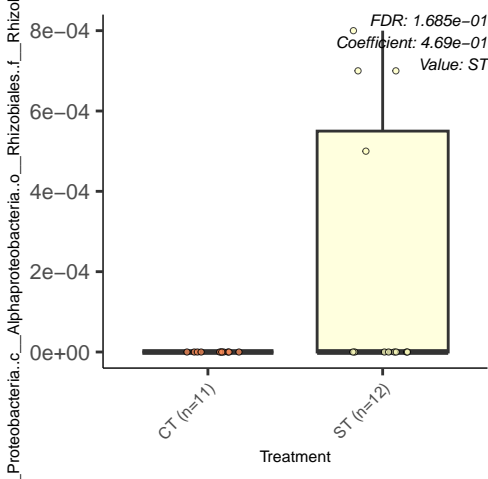


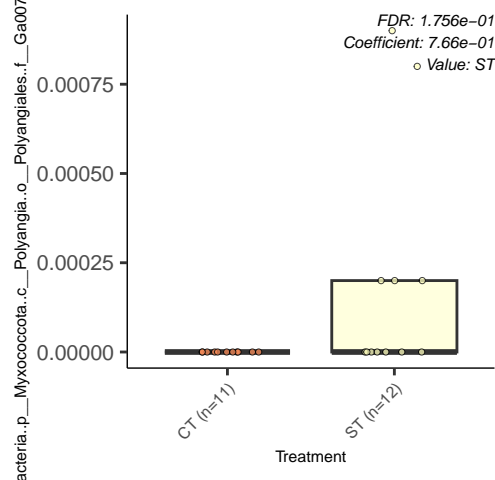


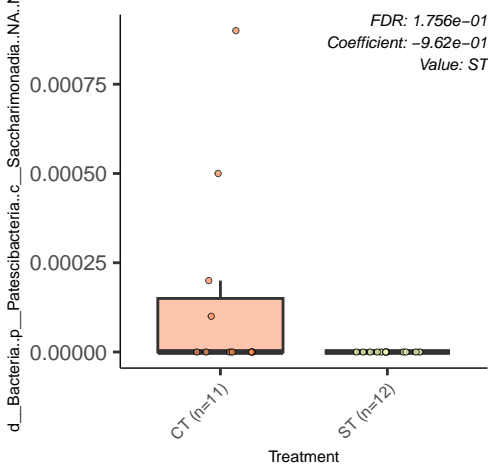


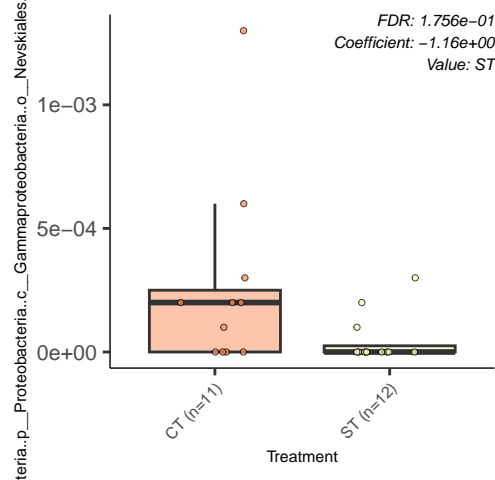






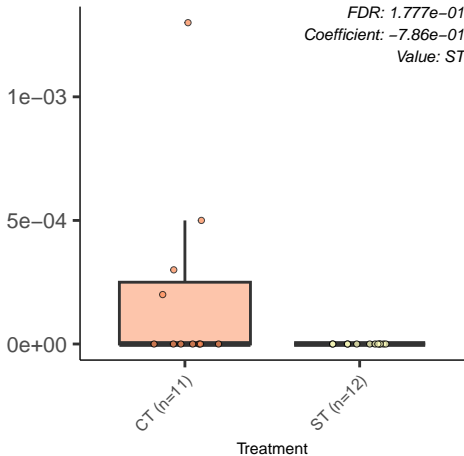


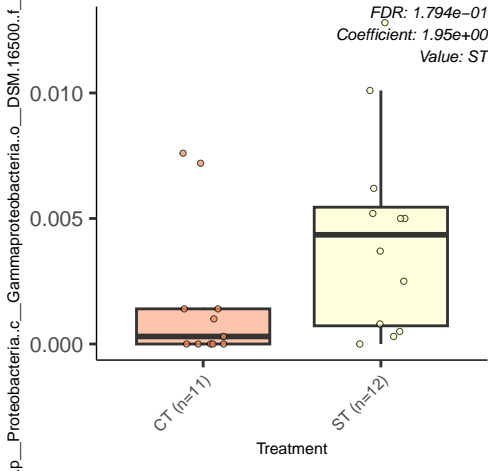


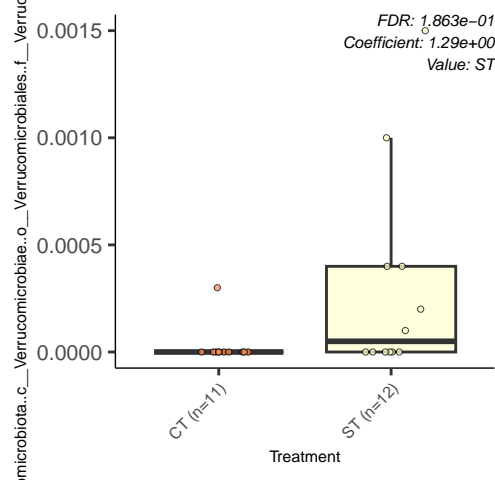


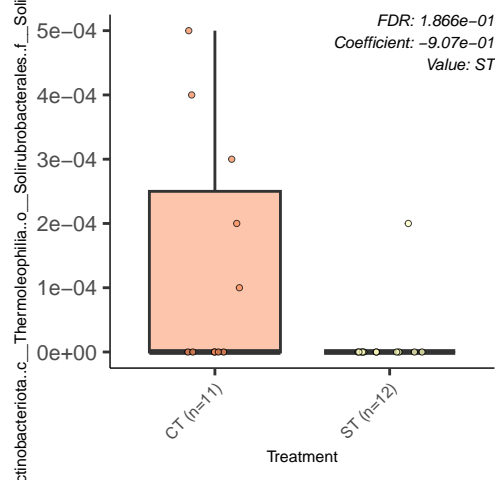
a.p__Planctomycetota..c__Planctomycetia..o__Pirellulales..f__Lacip

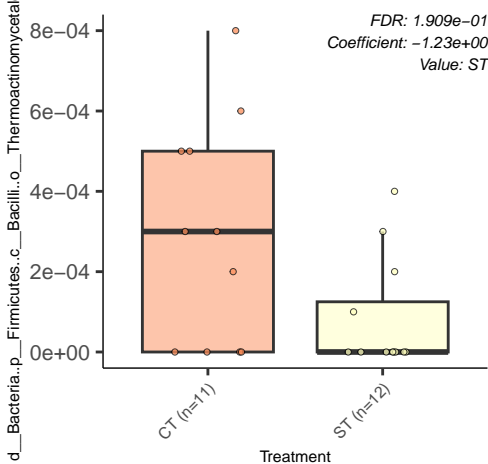
FDR: $1.777\text{e-}01$
Coefficient: $-7.86\text{e-}01$
Value: ST

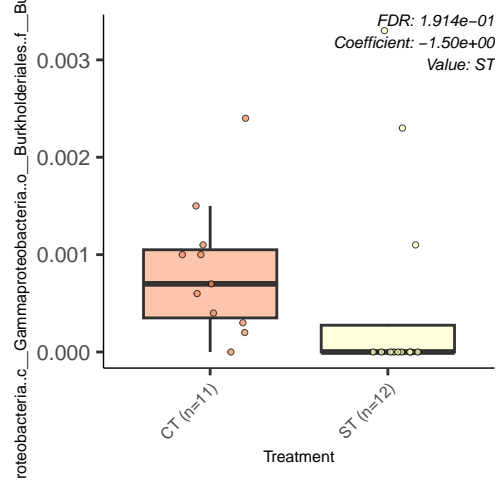


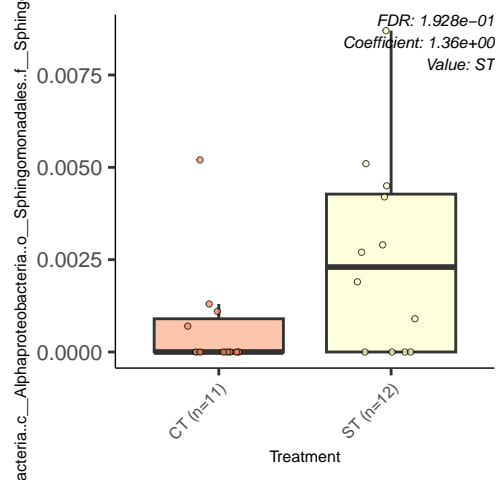


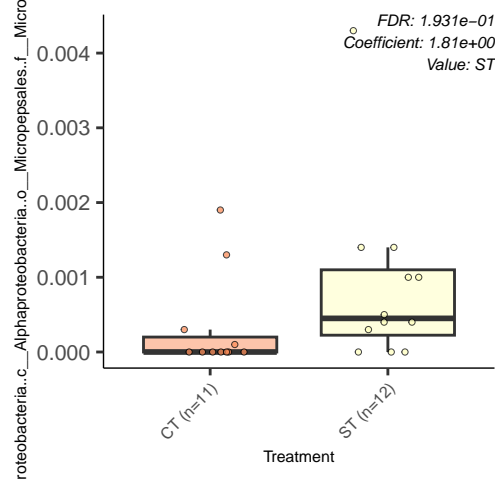


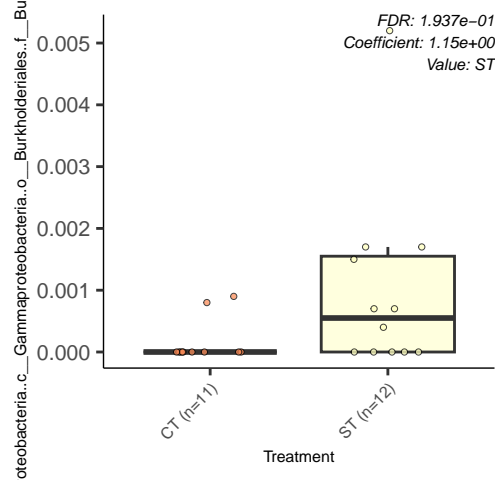


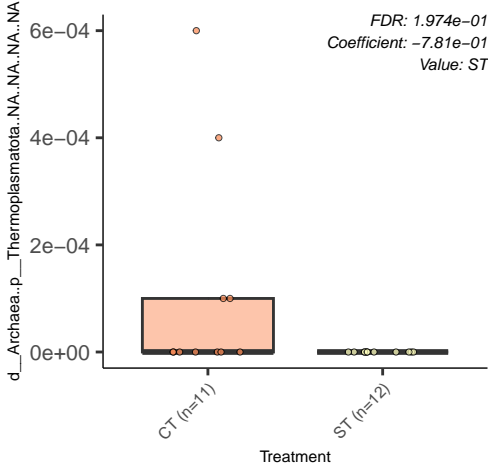


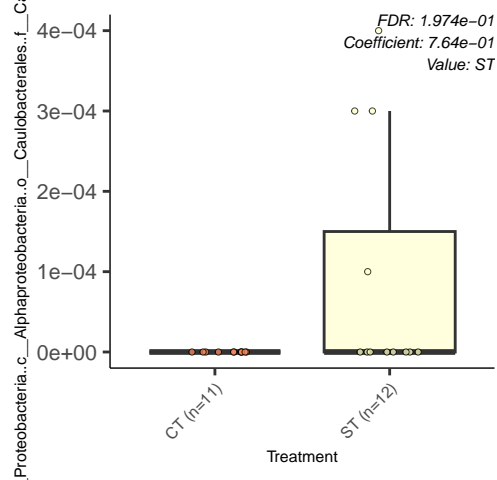


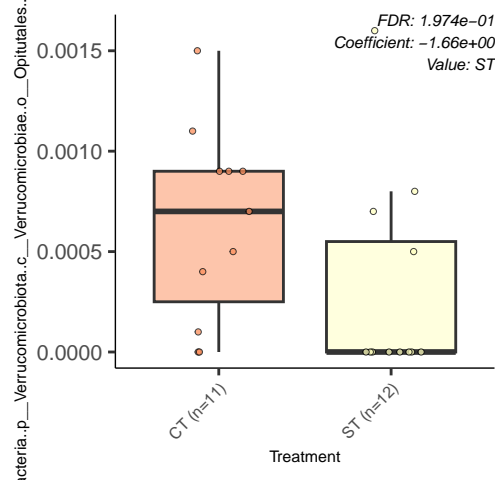


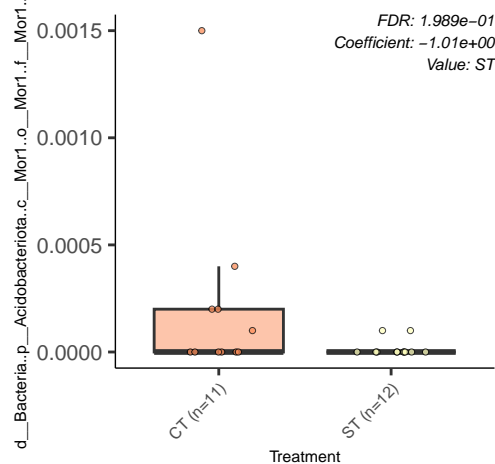


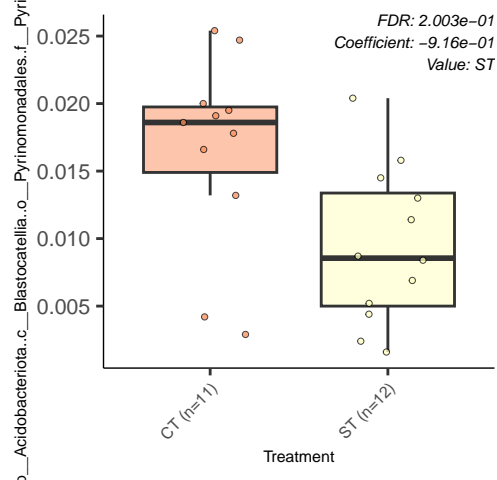




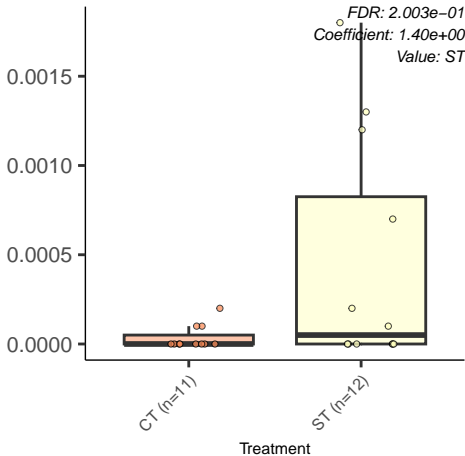


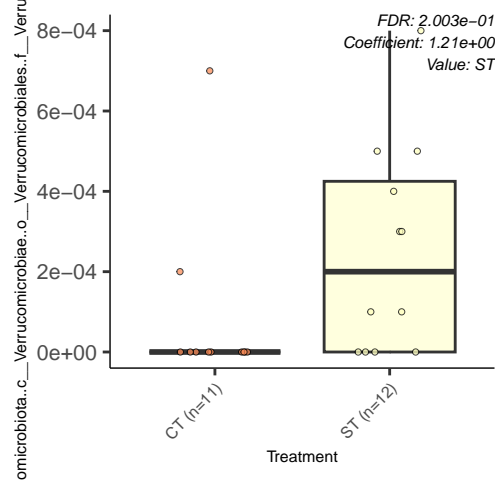


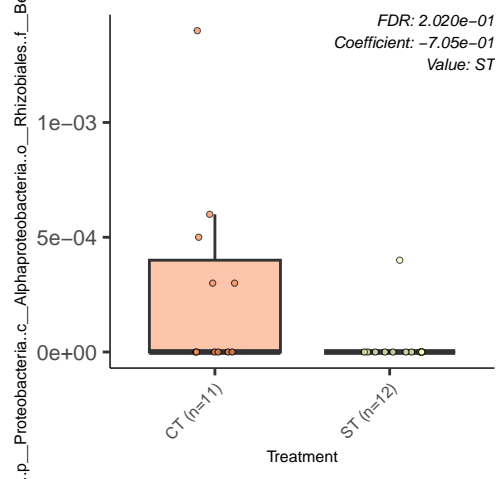


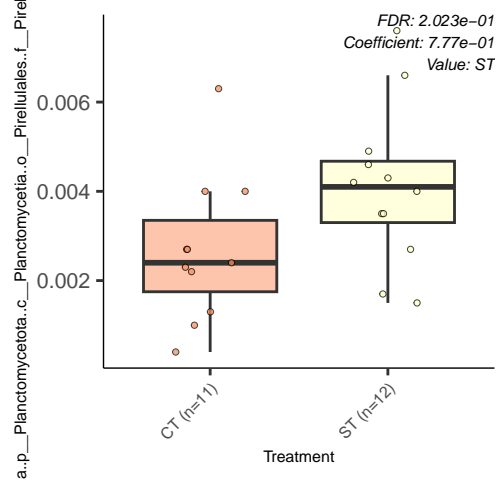


acteria..p__Dependentiae..c__Babeliae..o__Babeliales..f__UBA124









.p__Patescibacteria..c__UBA1384..o__2.12.FULL.50.11..f__2.12.F

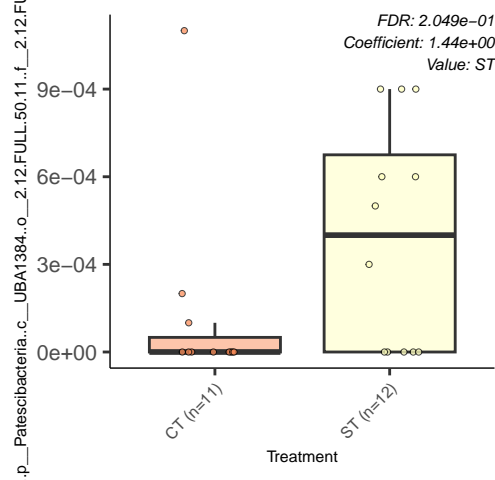
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Coefficient: 1.44e+00
Value: ST

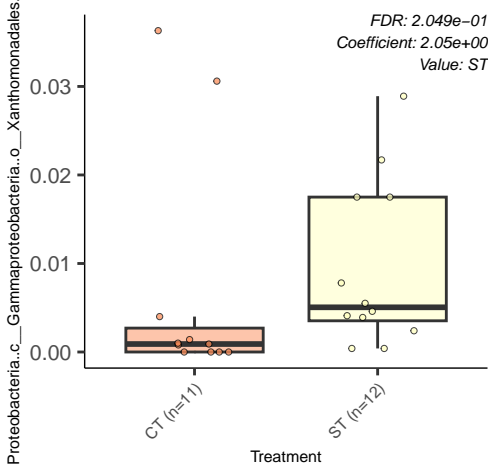
9e-04
6e-04
3e-04
0e+00

CT (n=11)

ST (n=12)

Treatment





d_Bacteria..NA..NA..NA..NA..NA

0.06

0.04

0.02

CT (n=11)

ST (n=12)

Treatment

FDR: 2.124e-01
Coefficient: 4.25e-01
Value: ST

