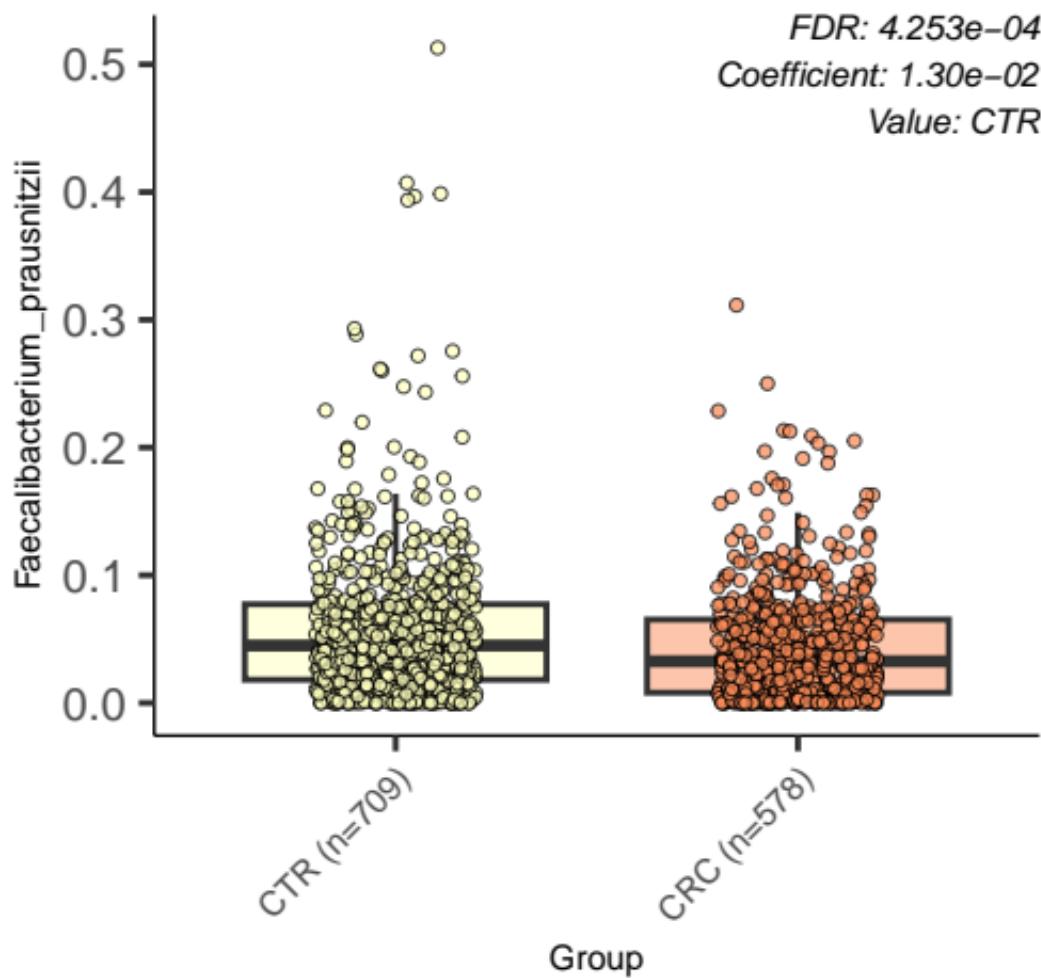
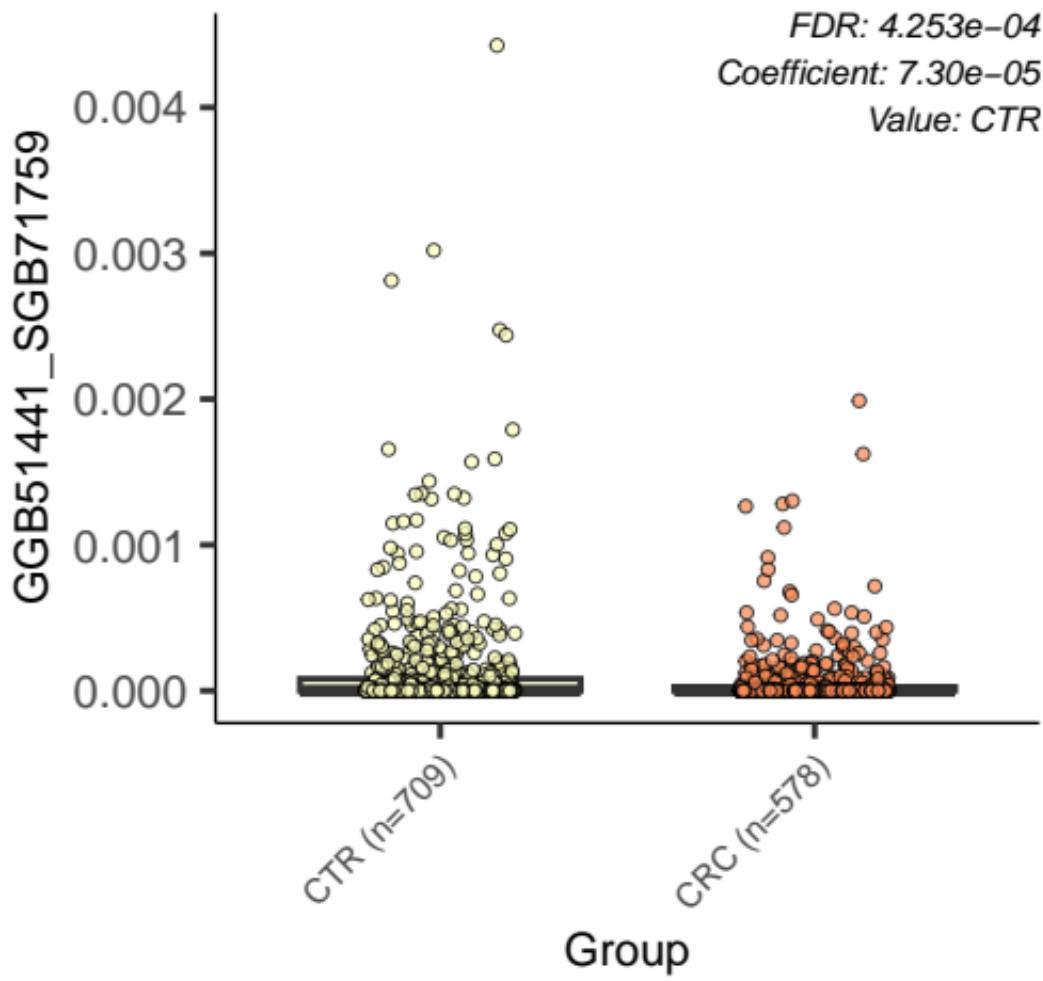
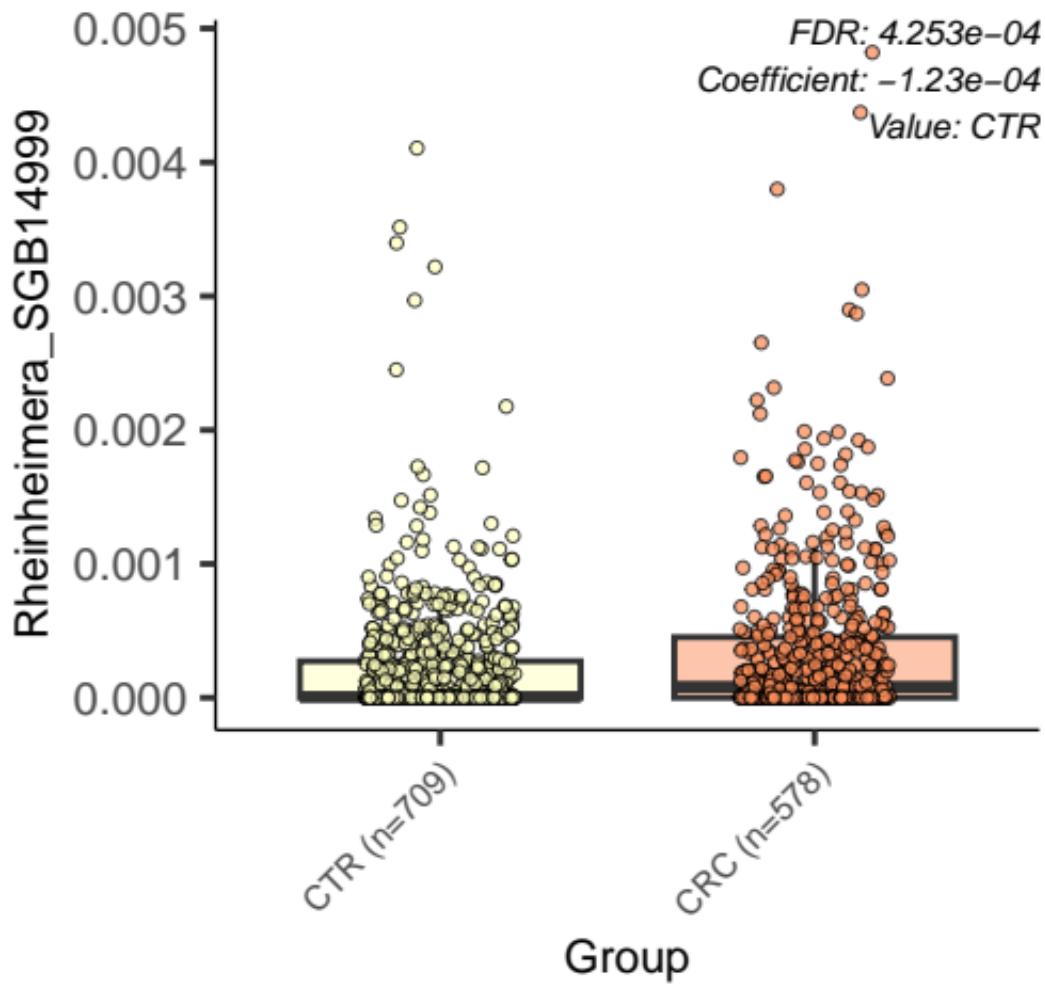
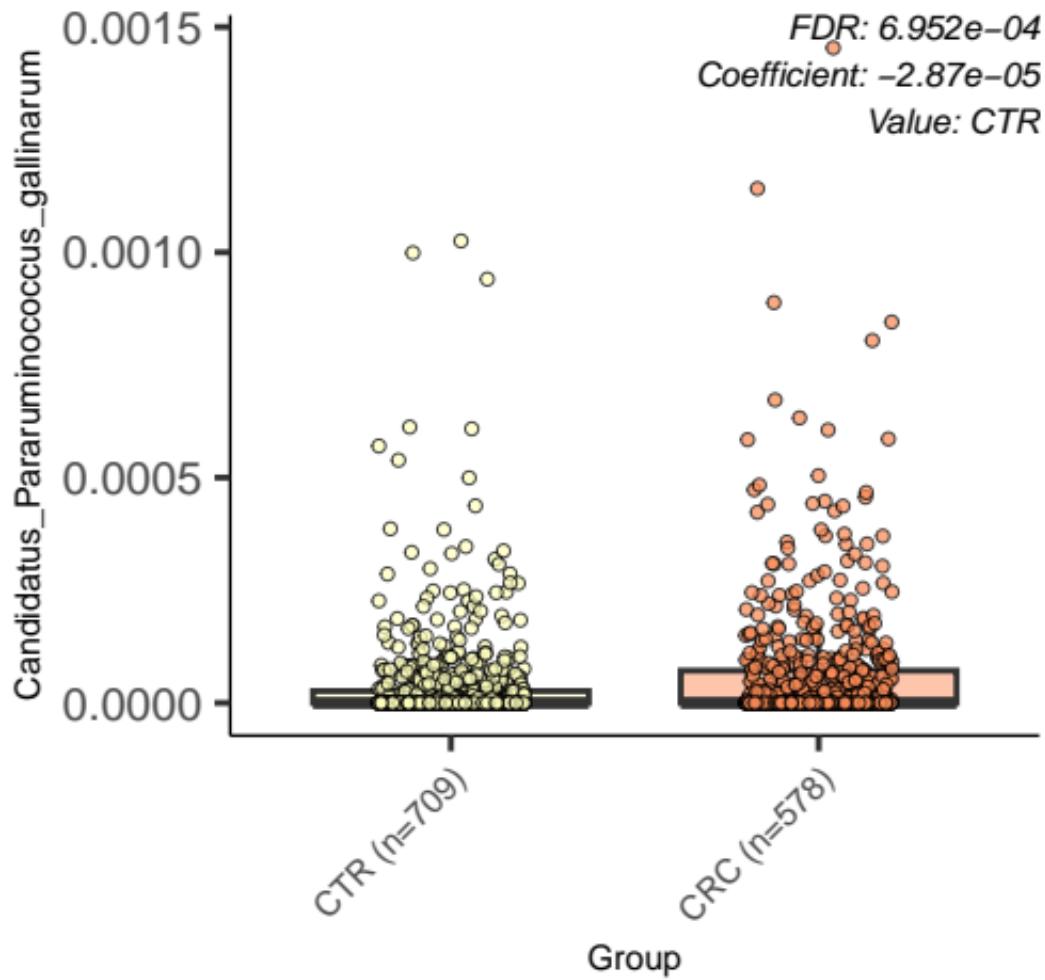


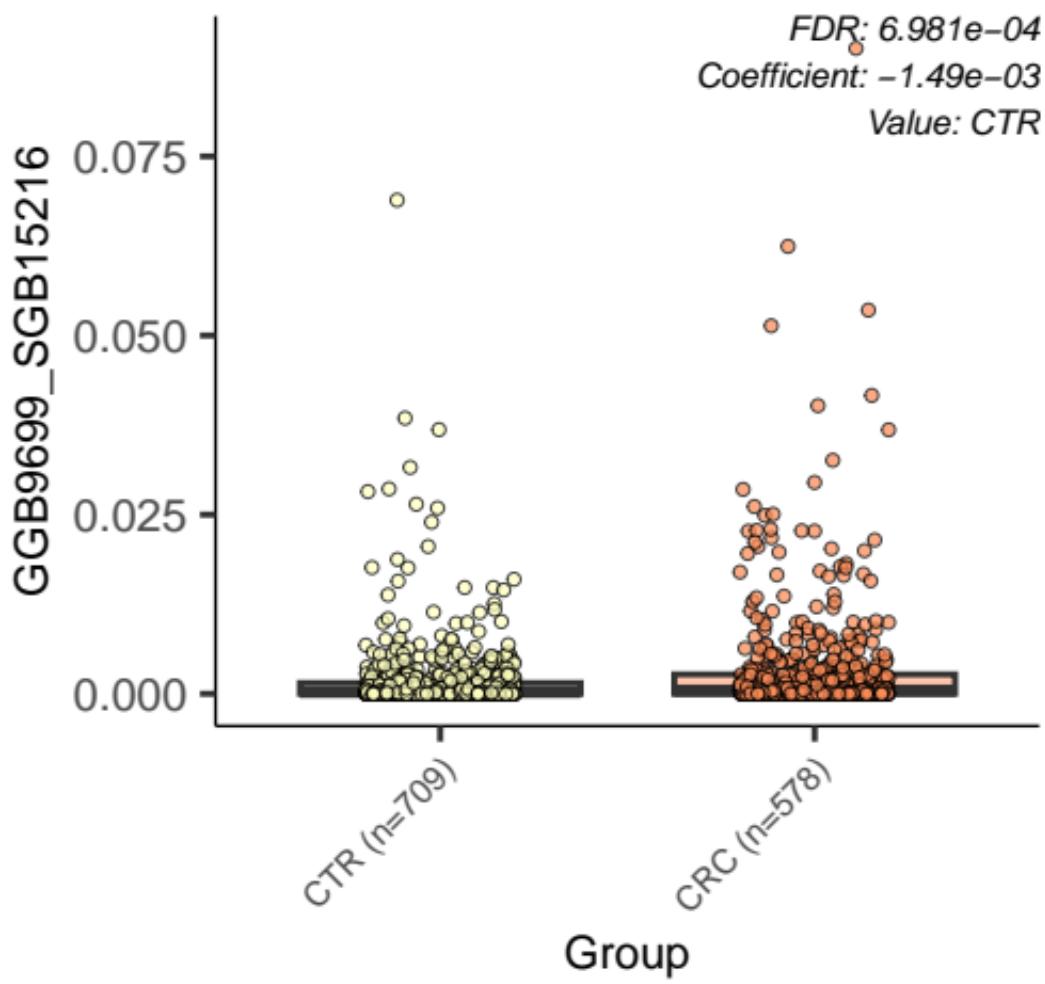
FDR: 4.253e-04
Coefficient: 1.30e-02
Value: CTR

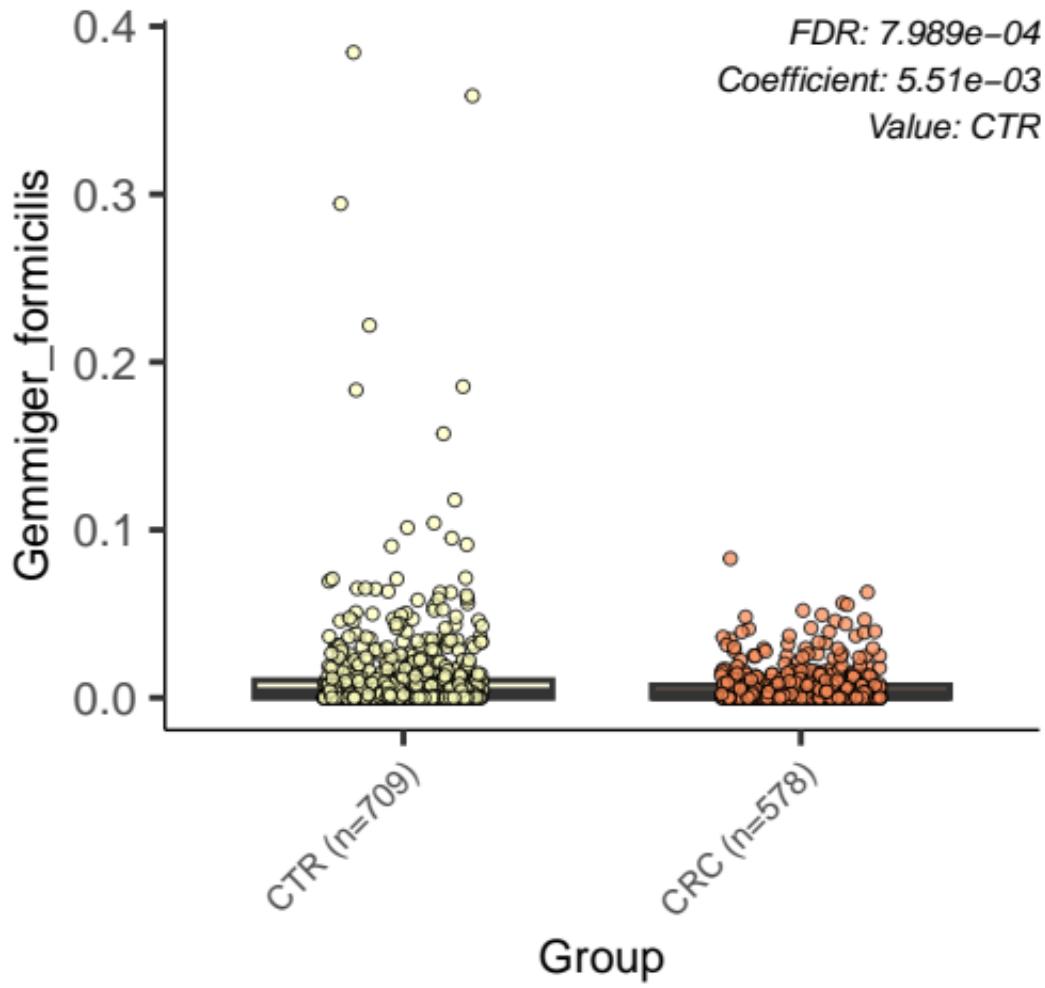


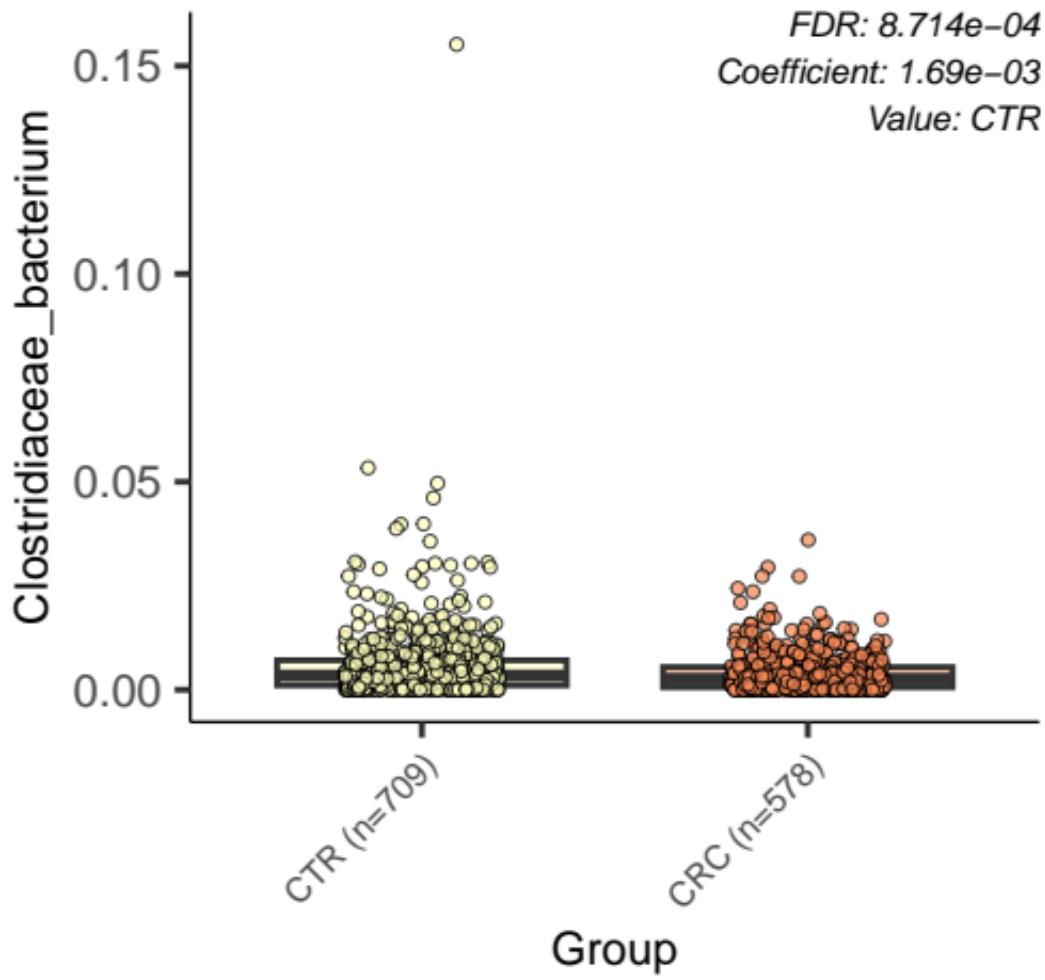


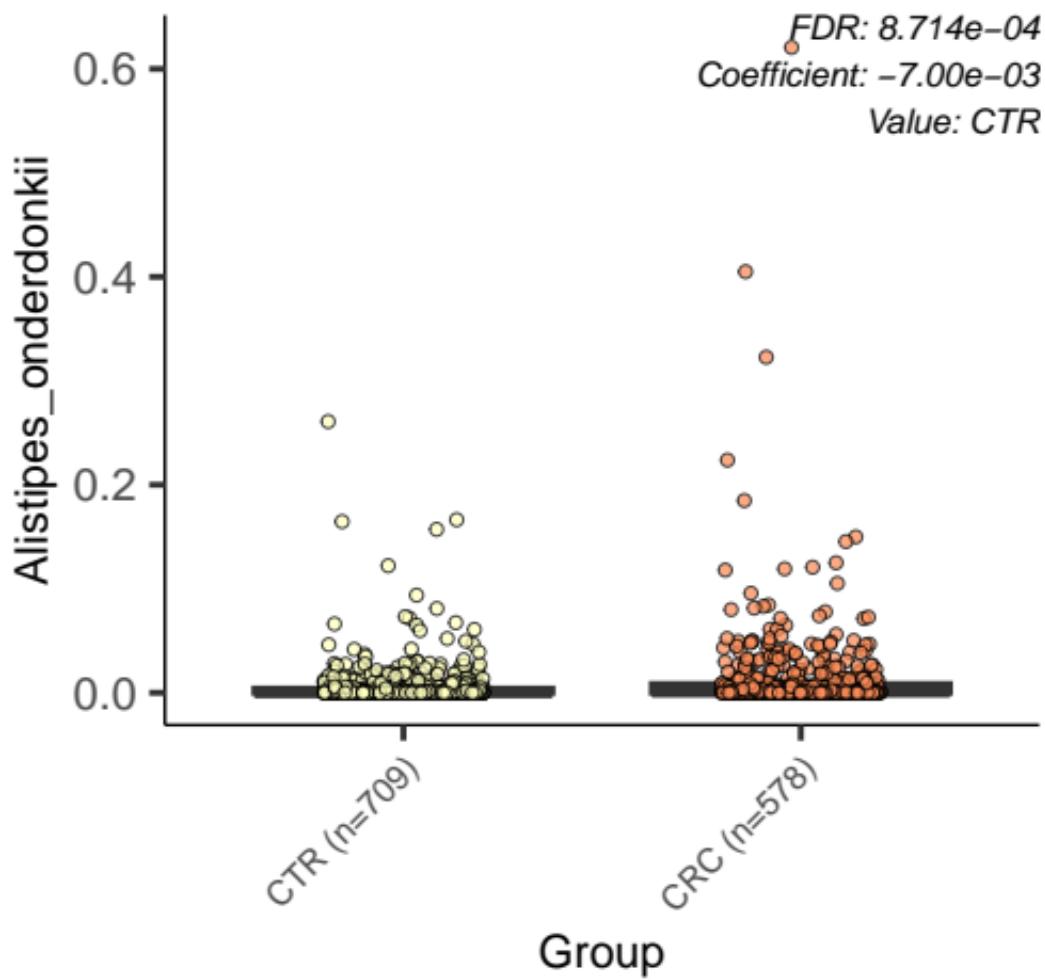


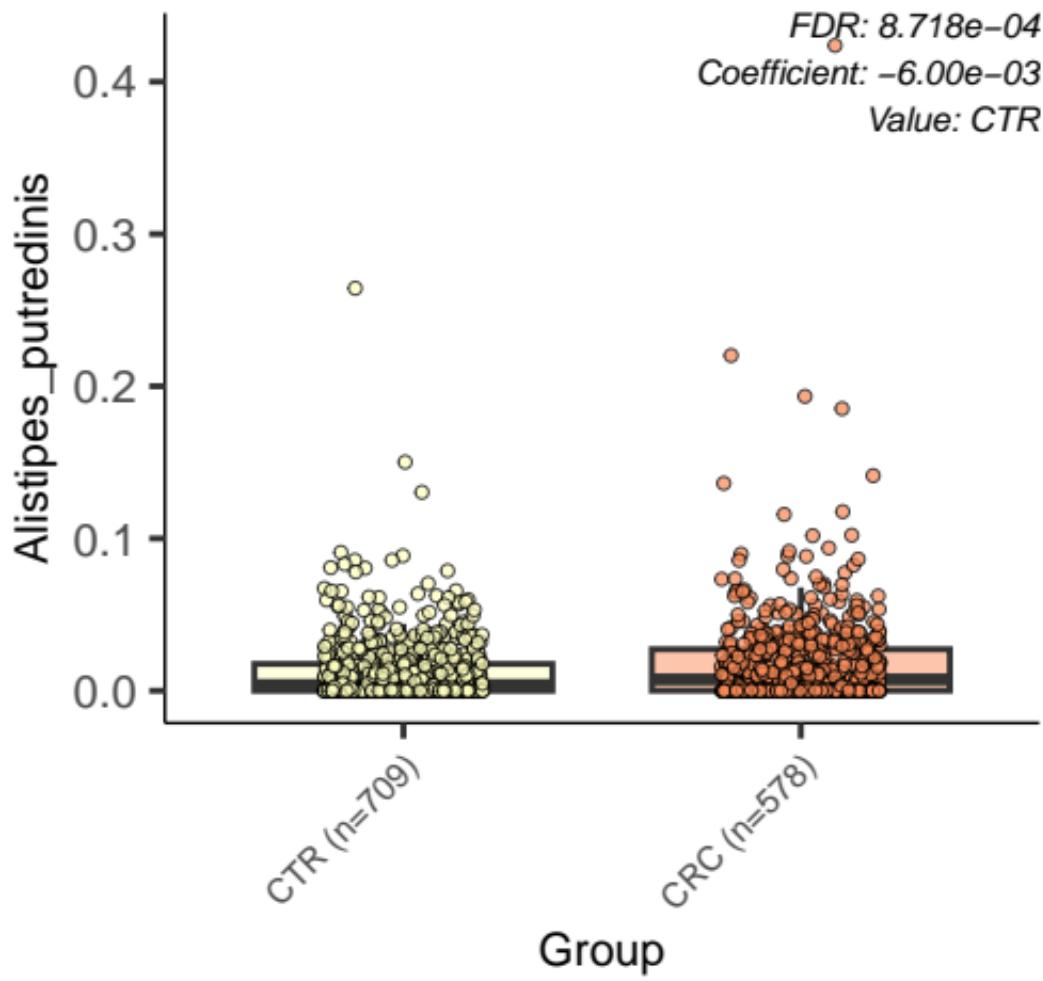


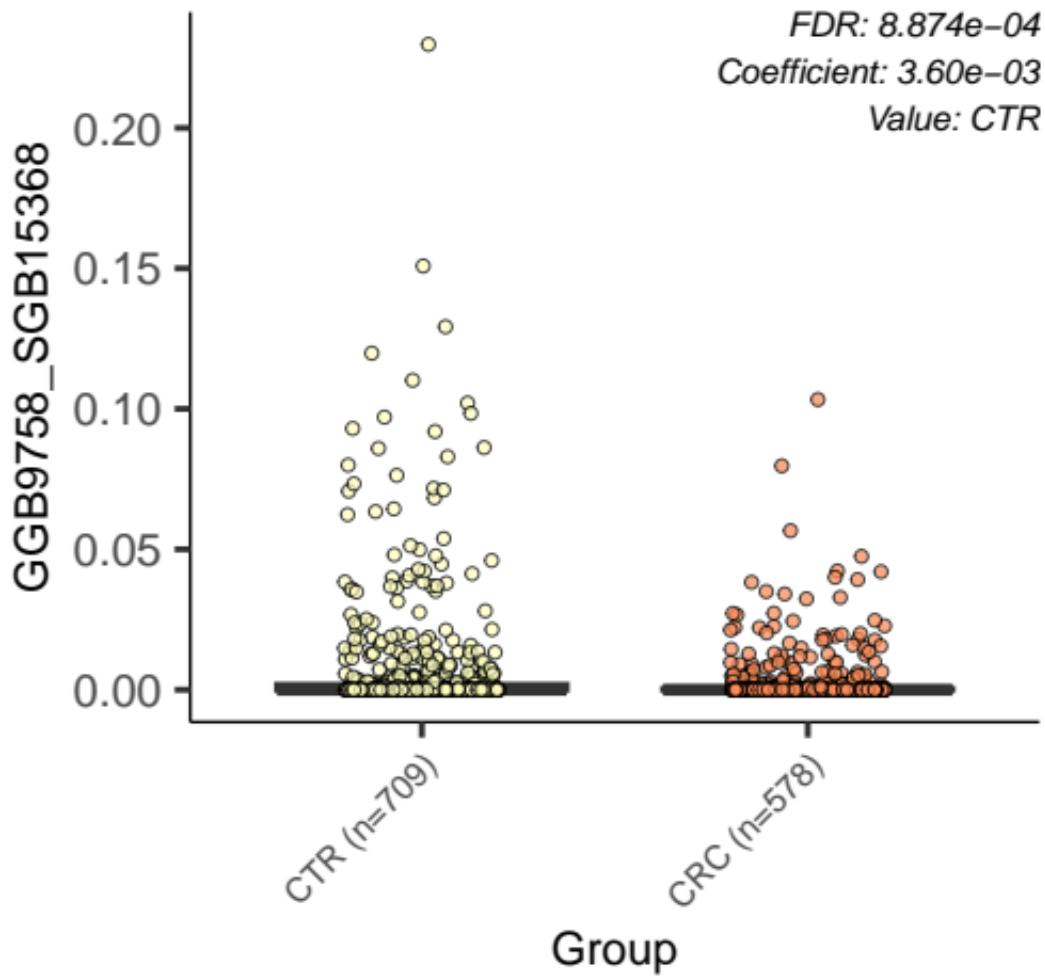


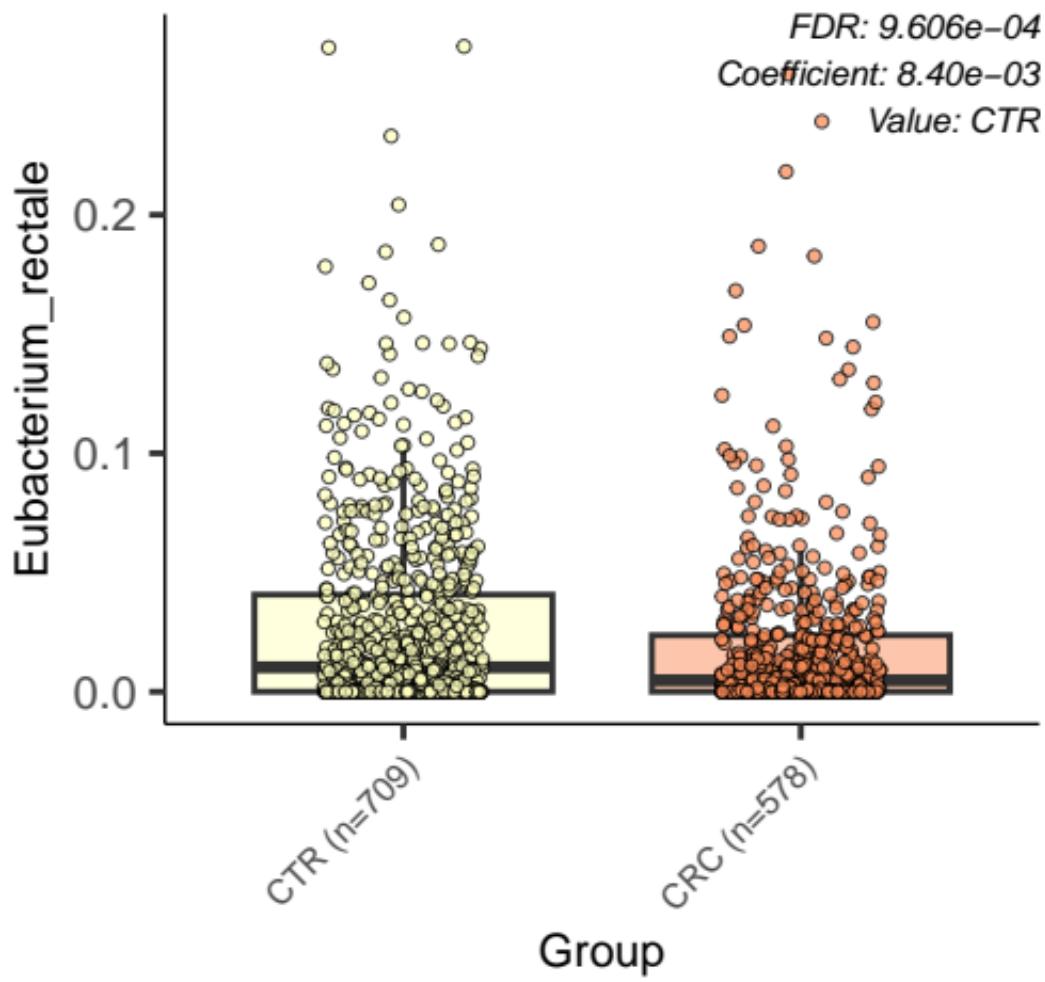


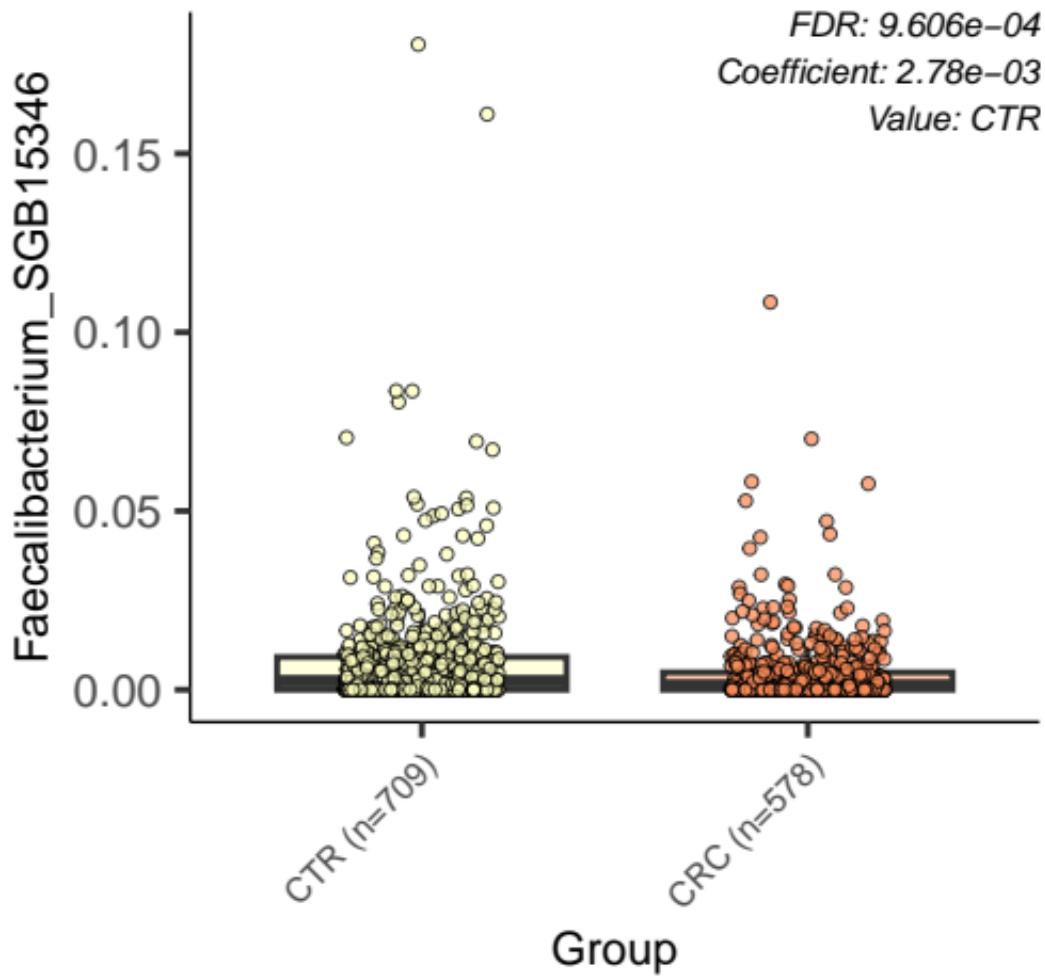


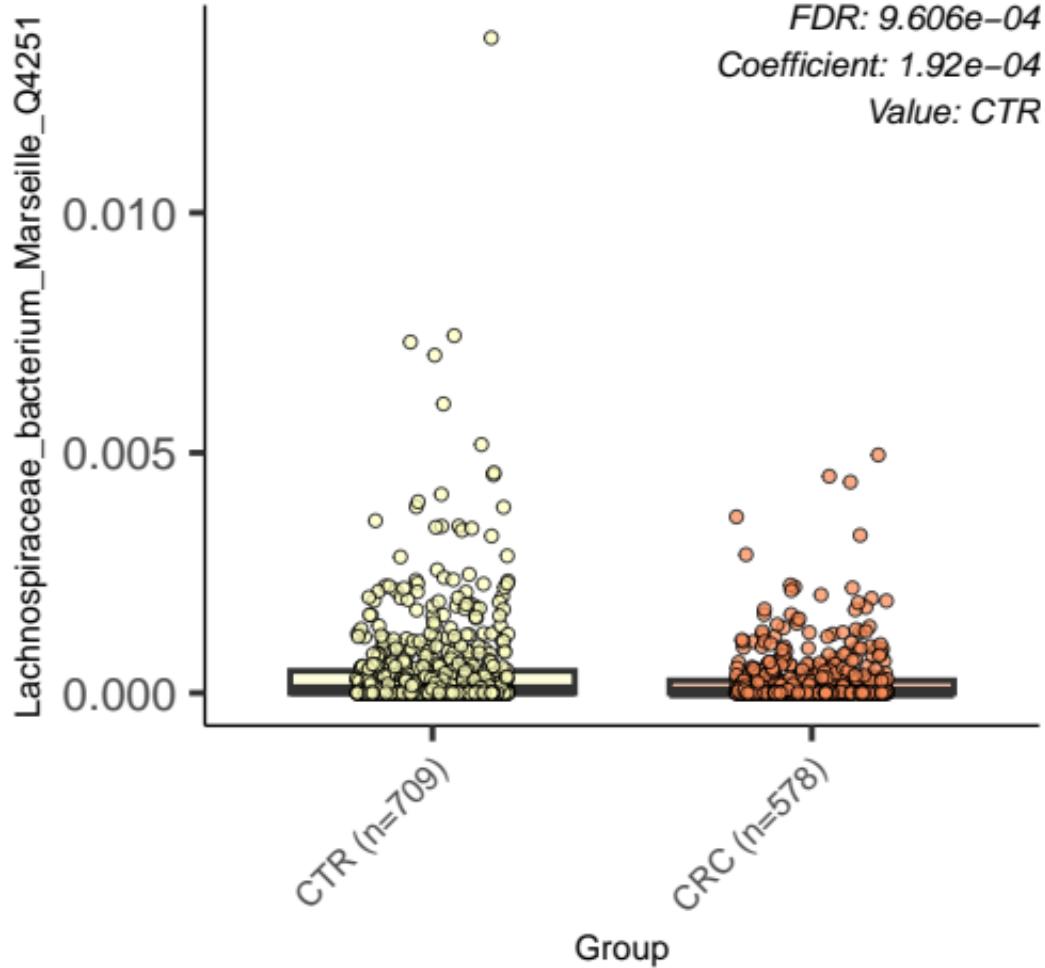


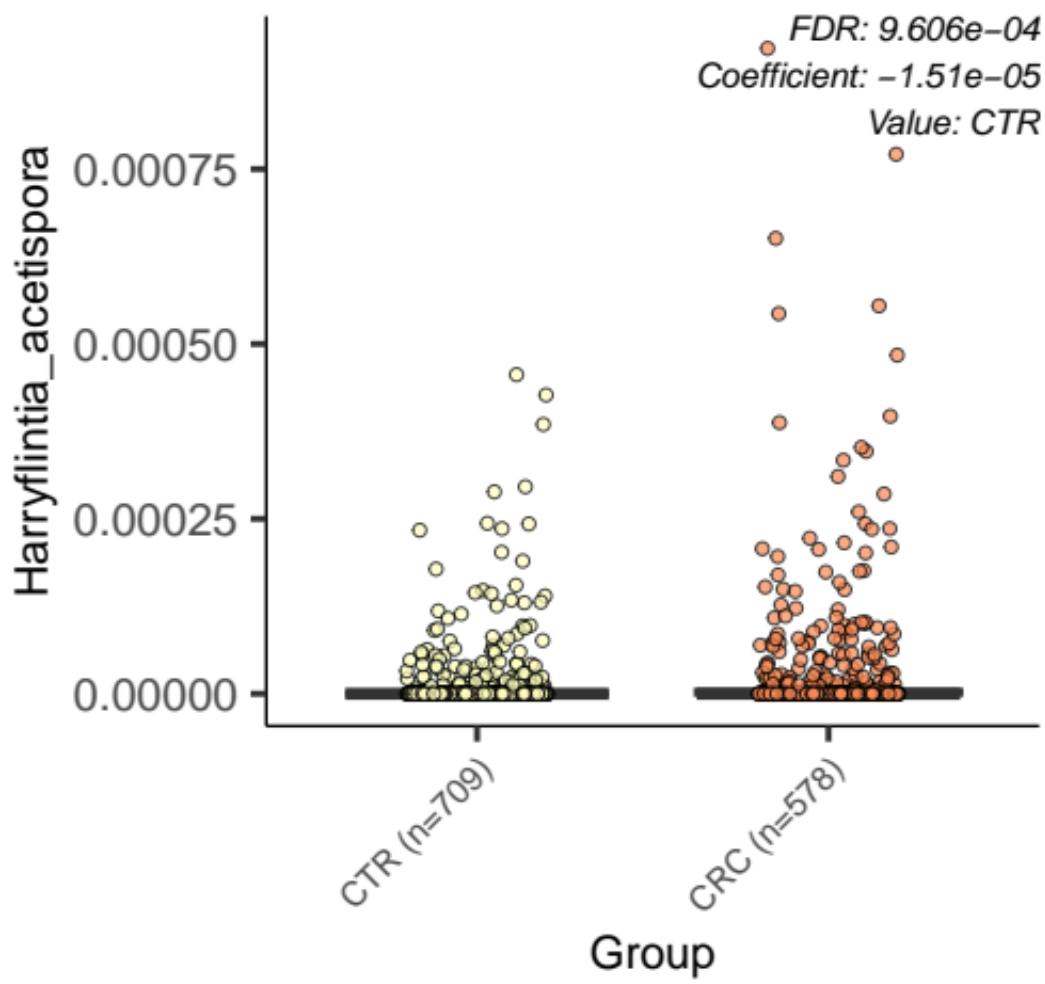


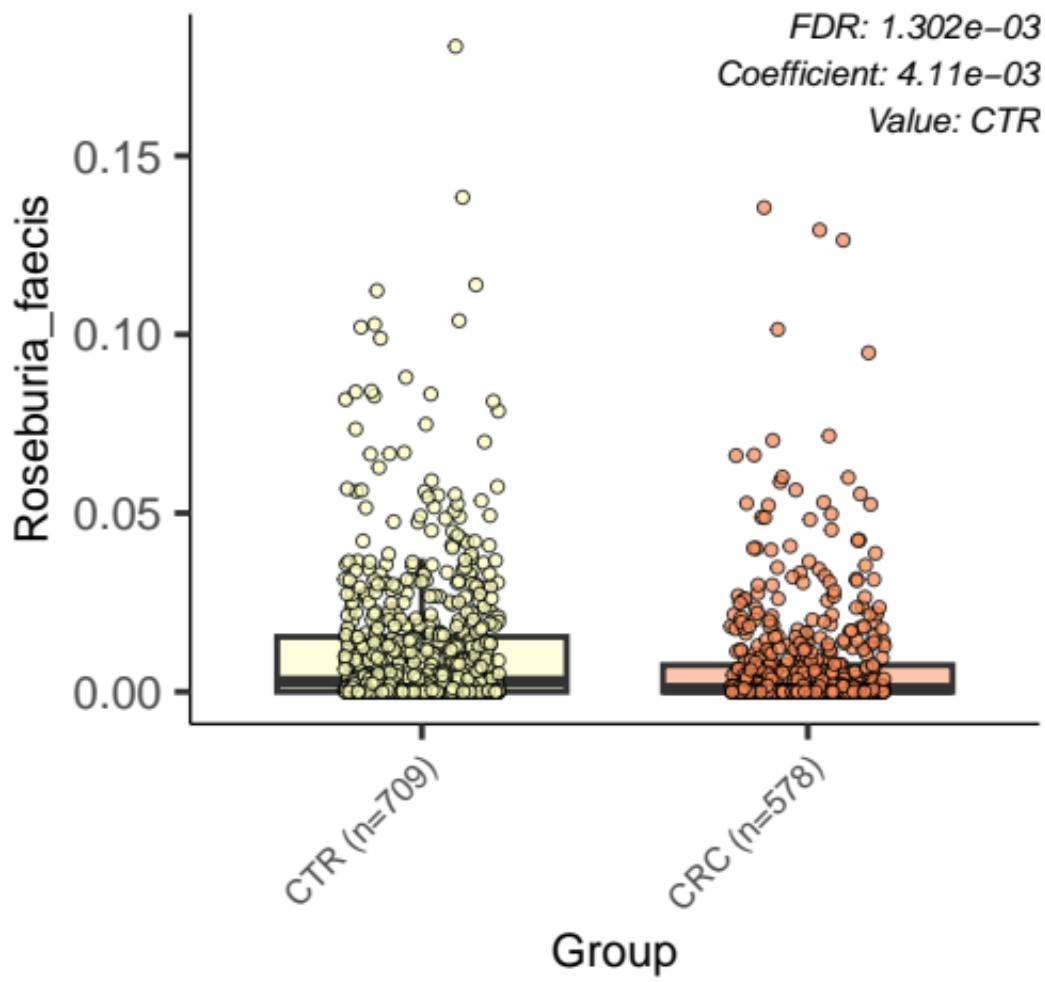


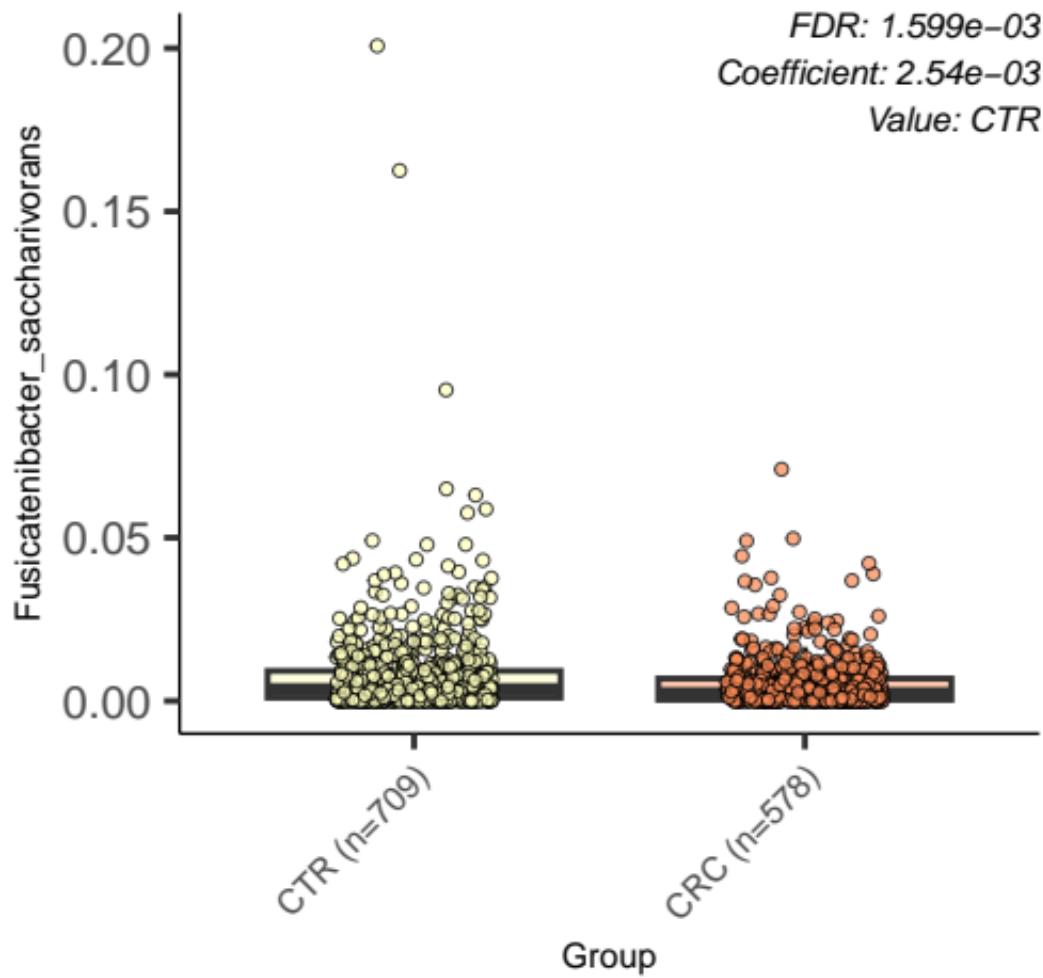












FDR: $1.640e-03$
Coefficient: $2.22e-04$
Value: CTR

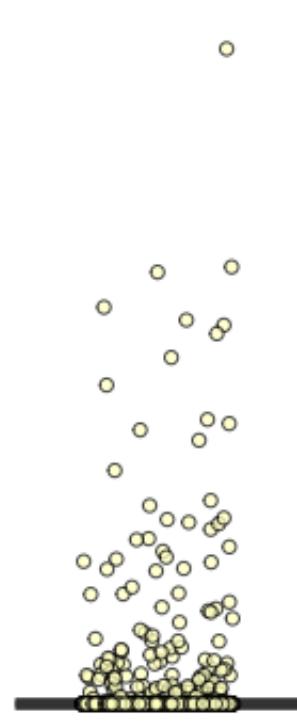
Lachnospiraceae_bacterium_BX3

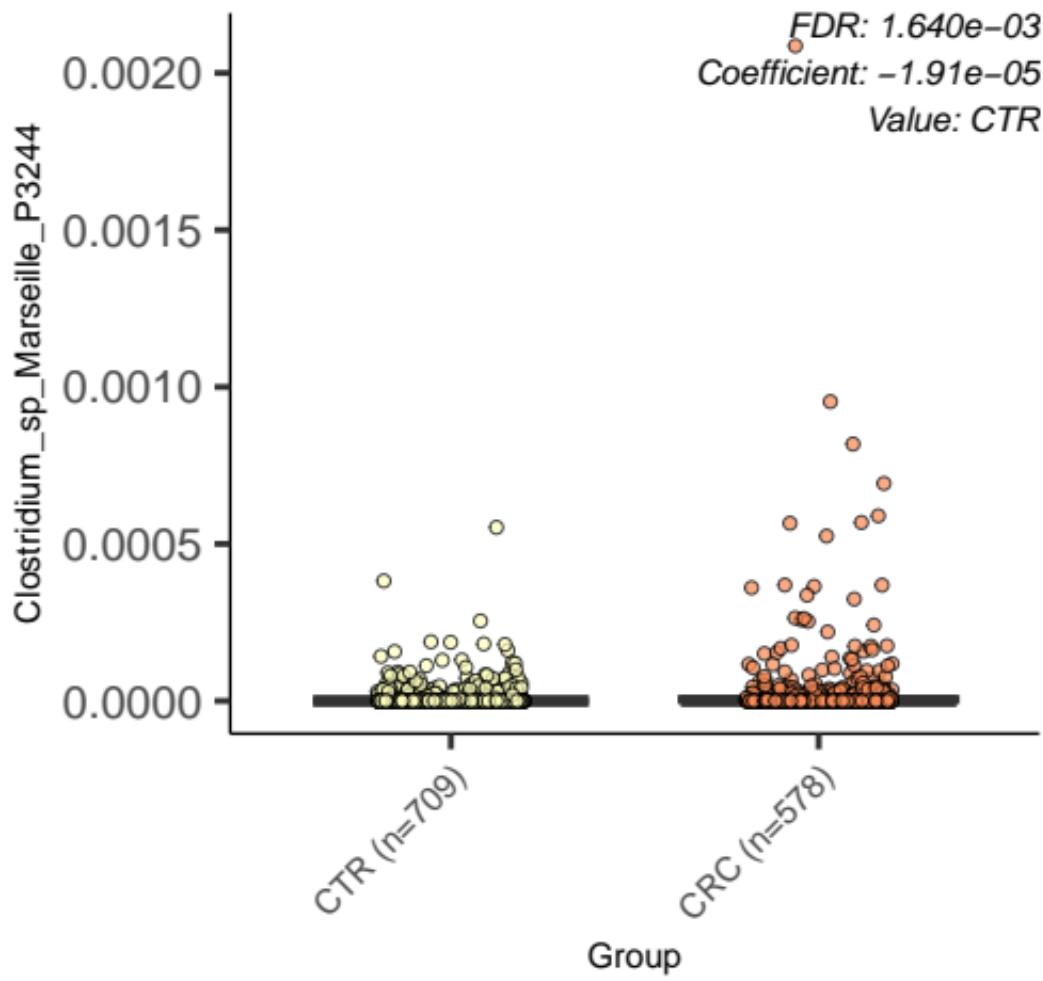
0.010
0.005
0.000

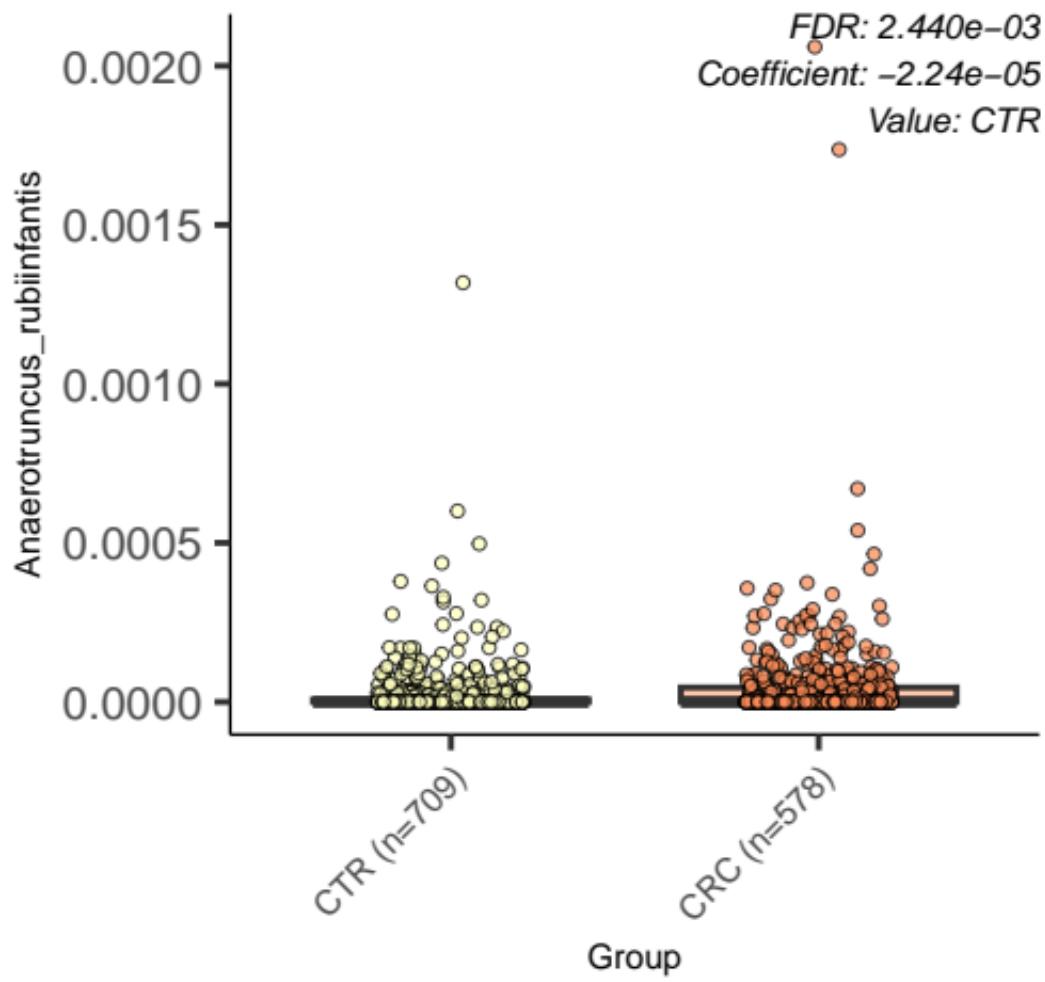
CTR (n=709)

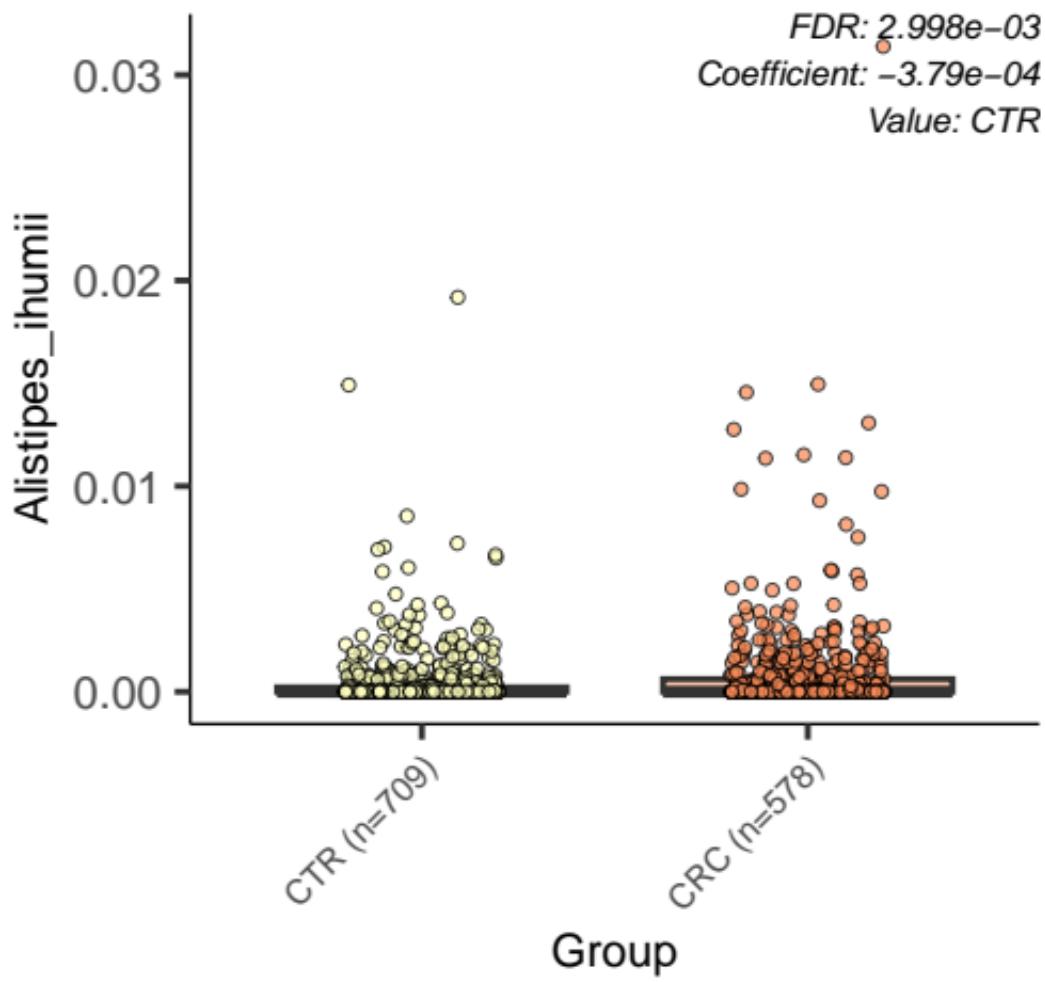
Group

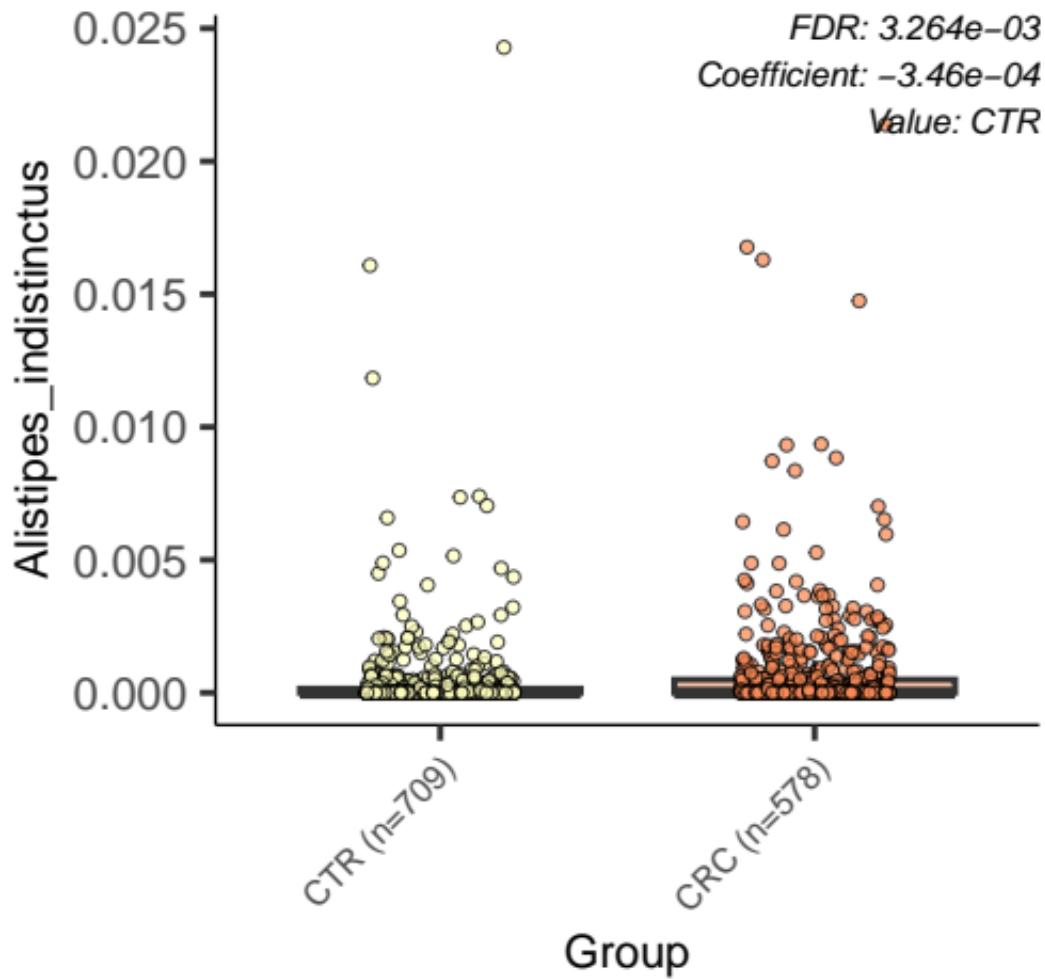
CRC (n=578)

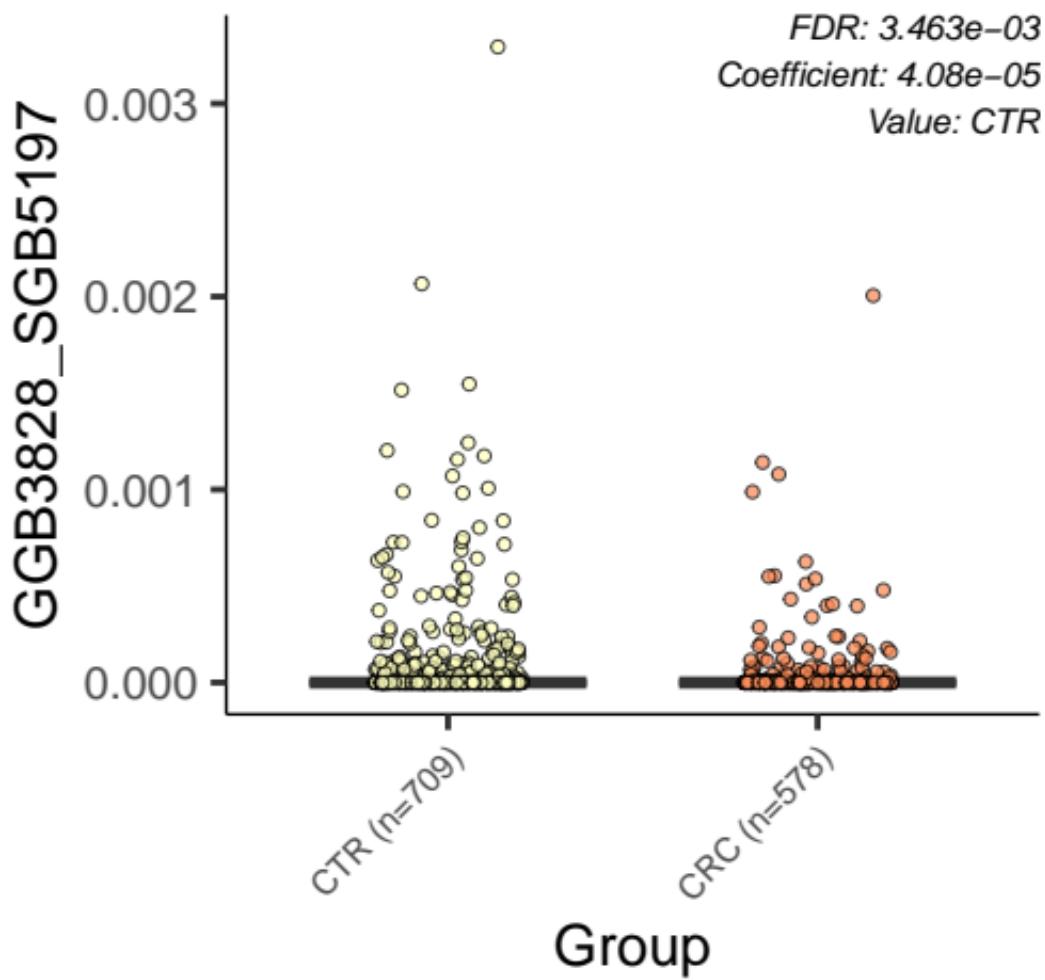


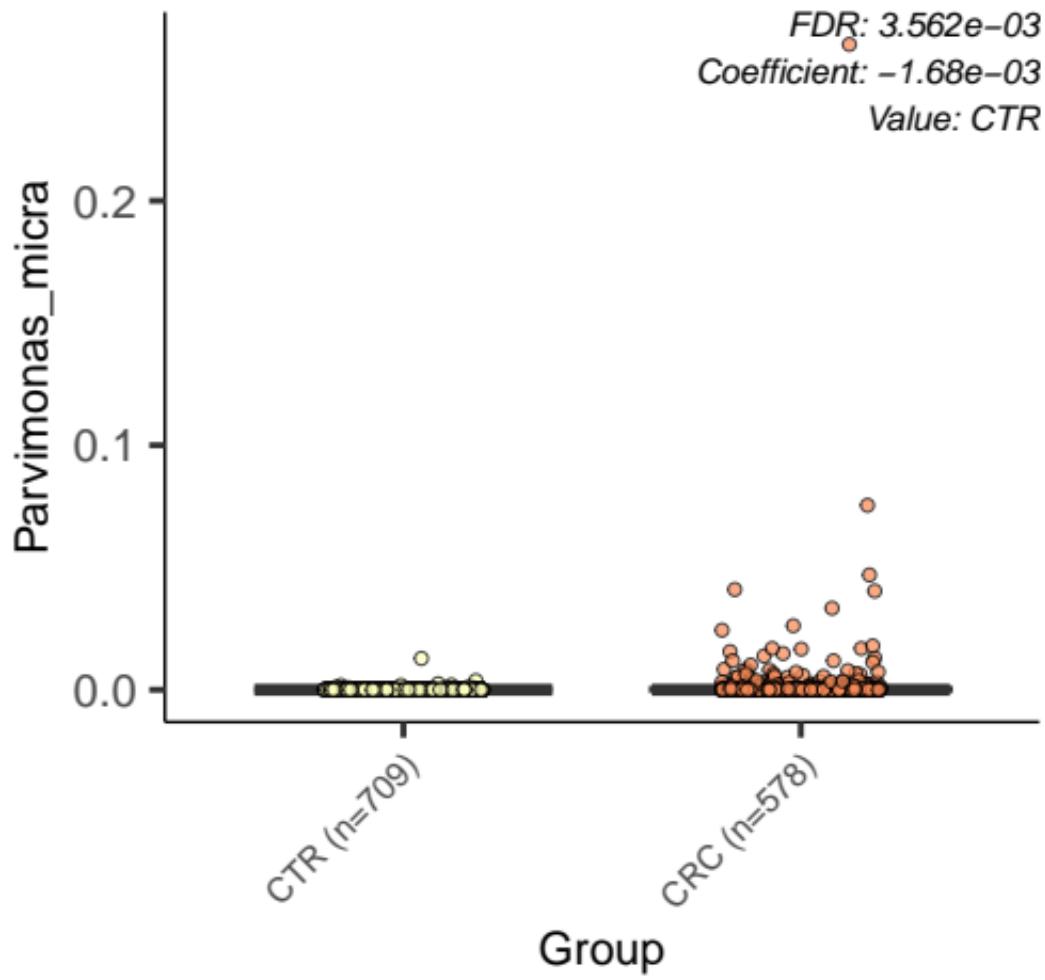


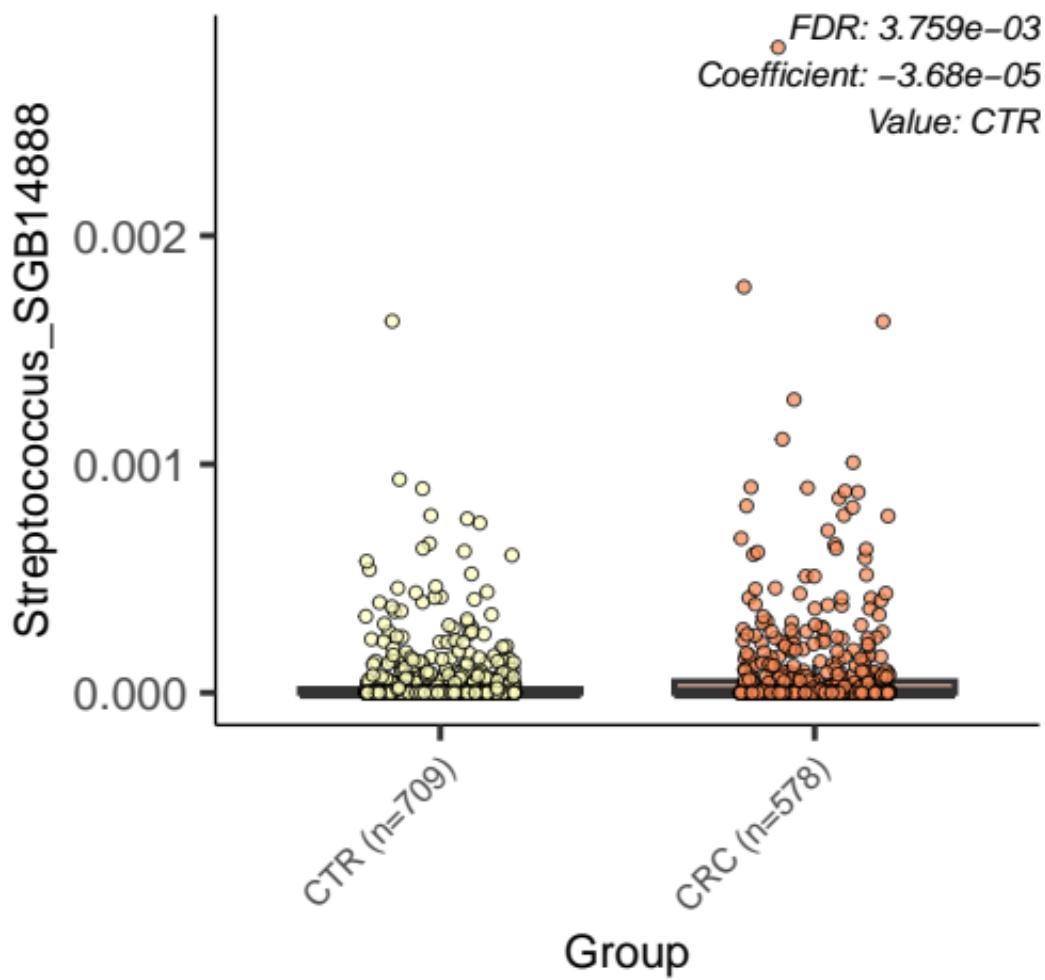


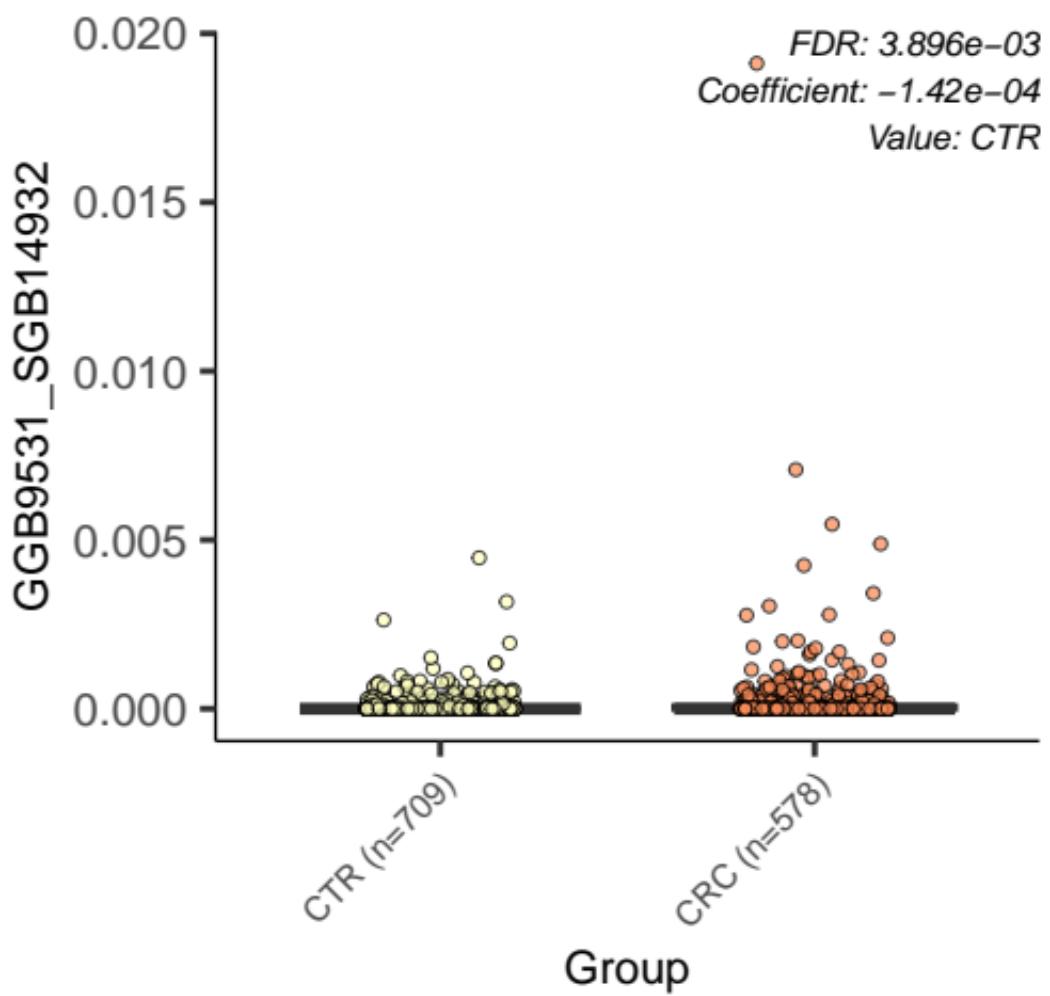


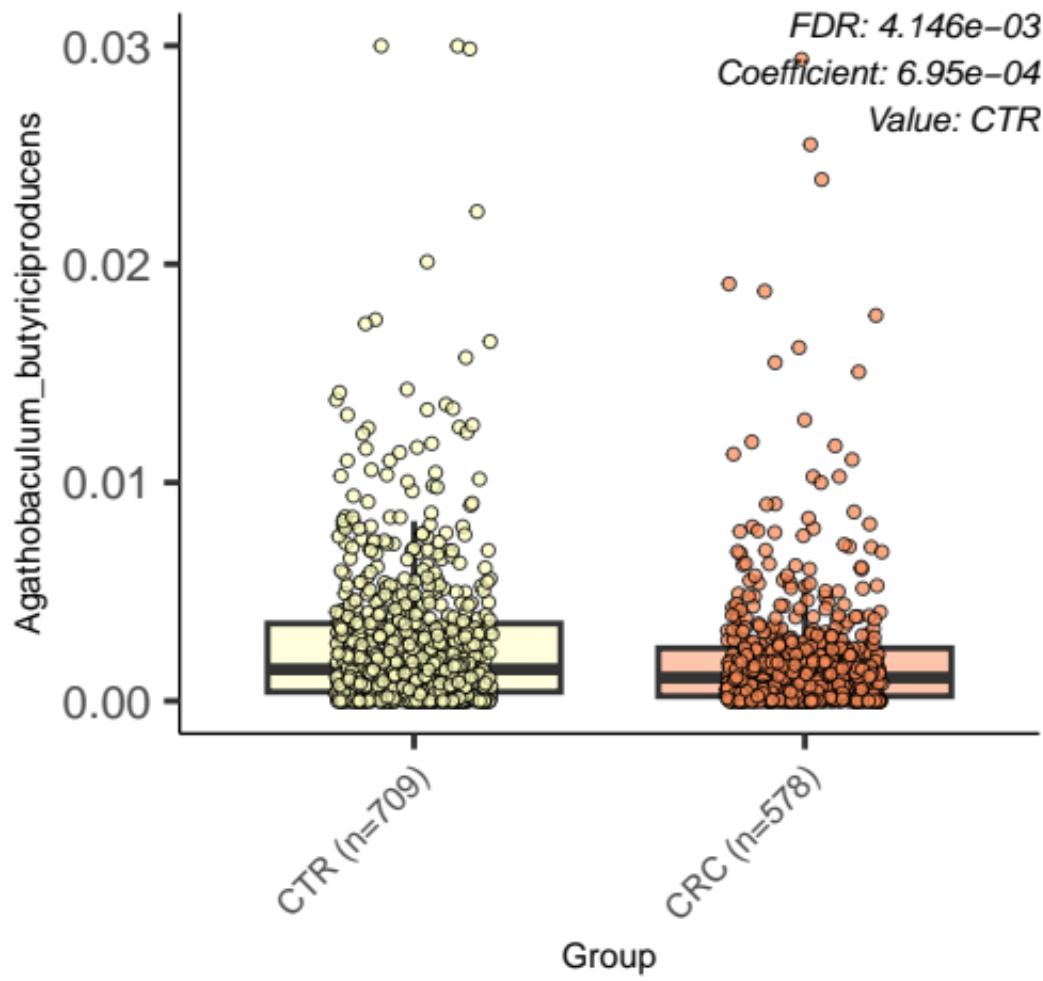


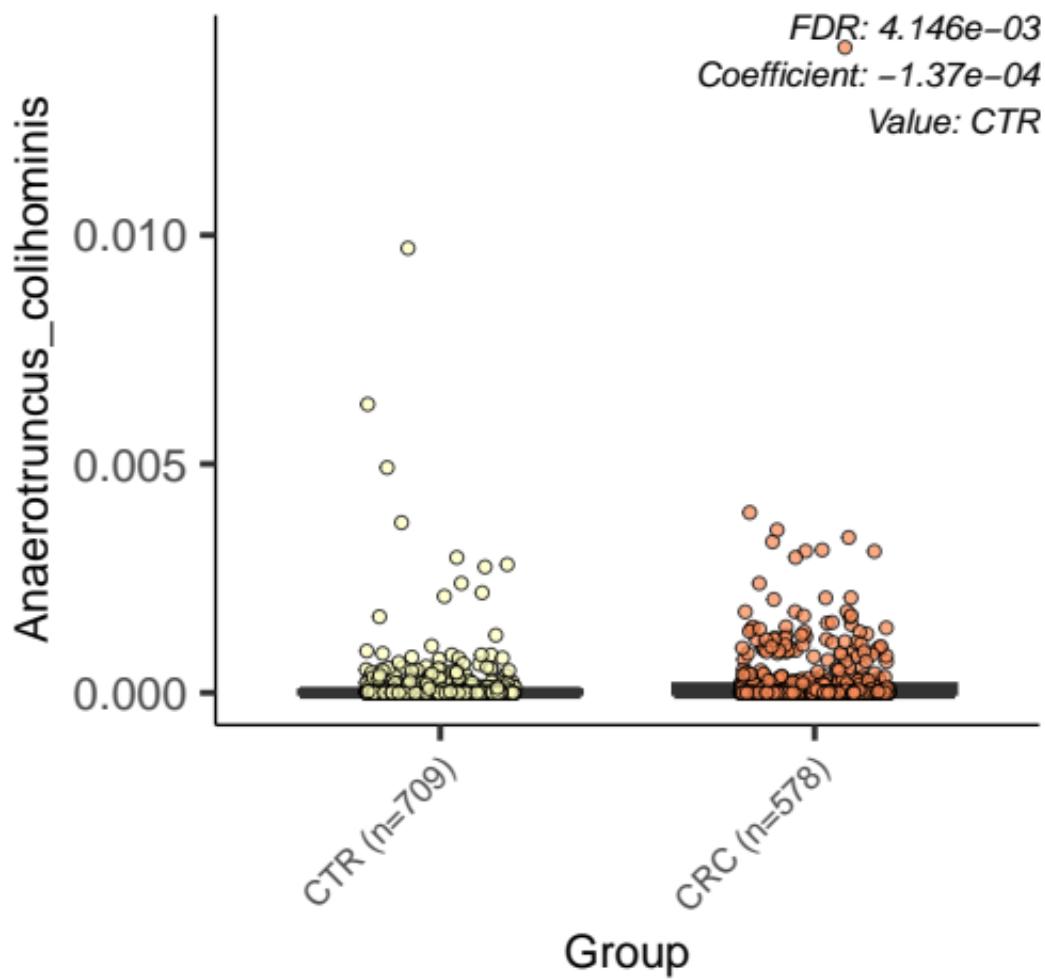


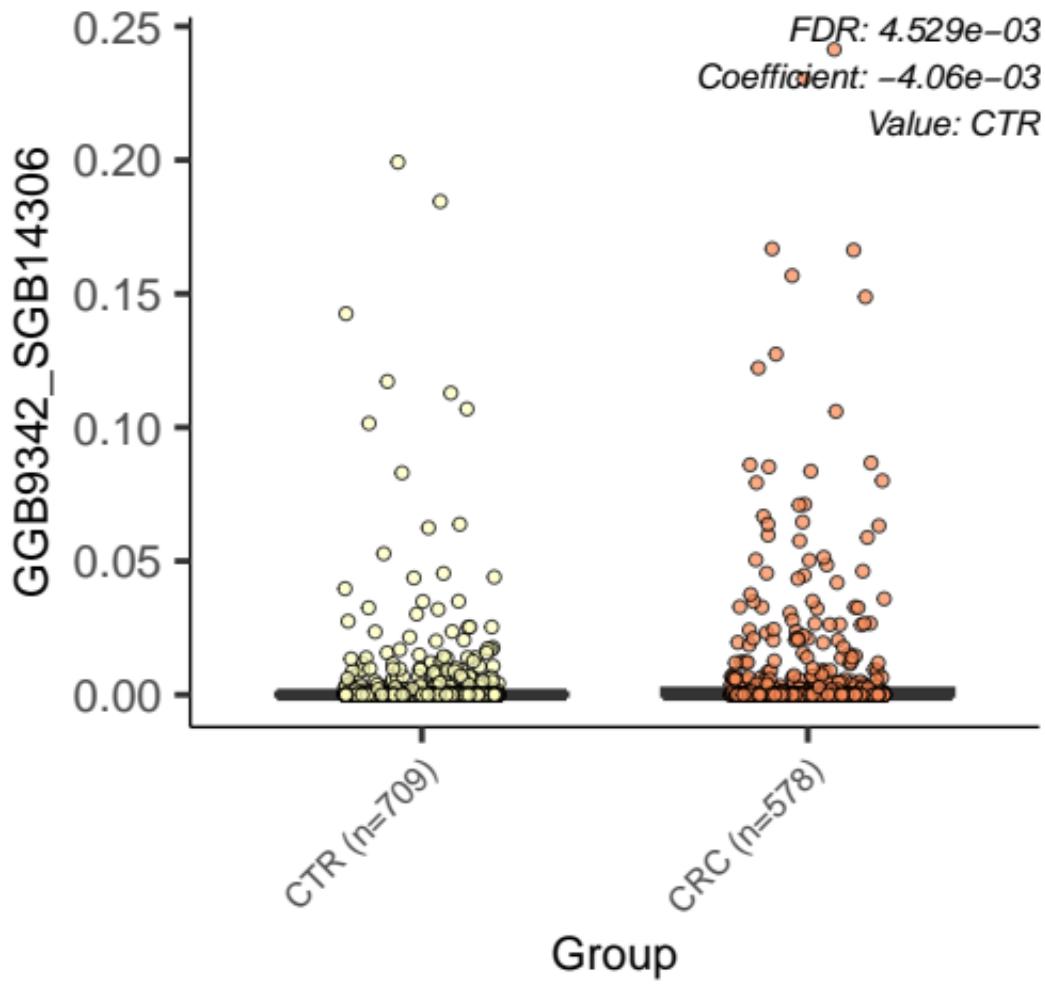


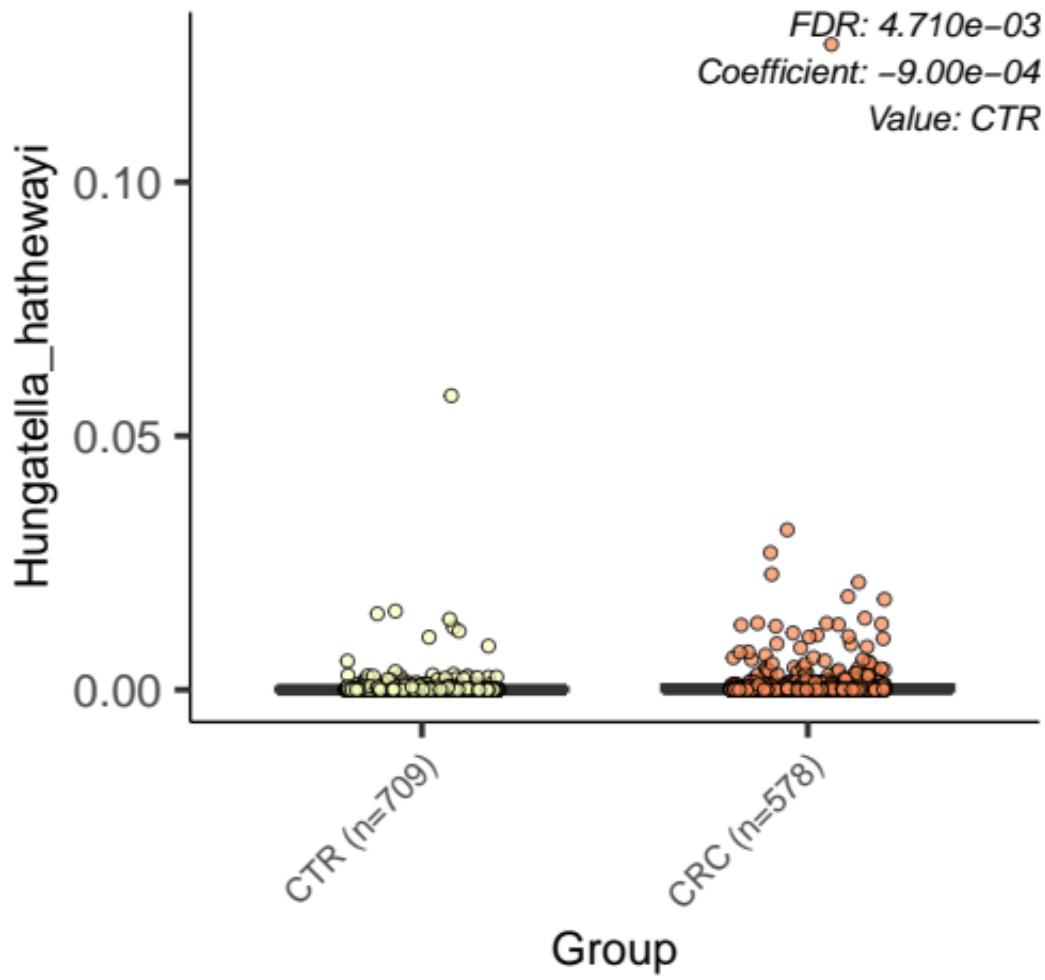


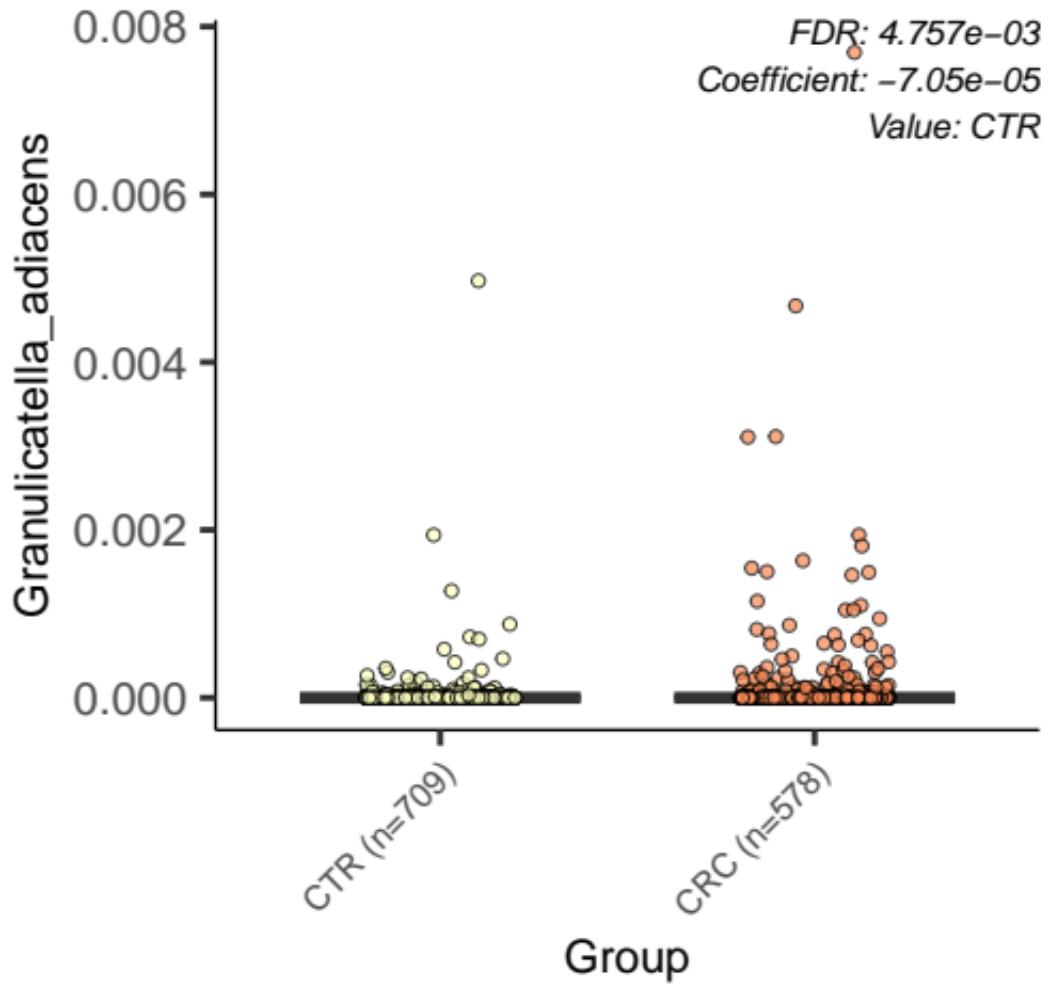


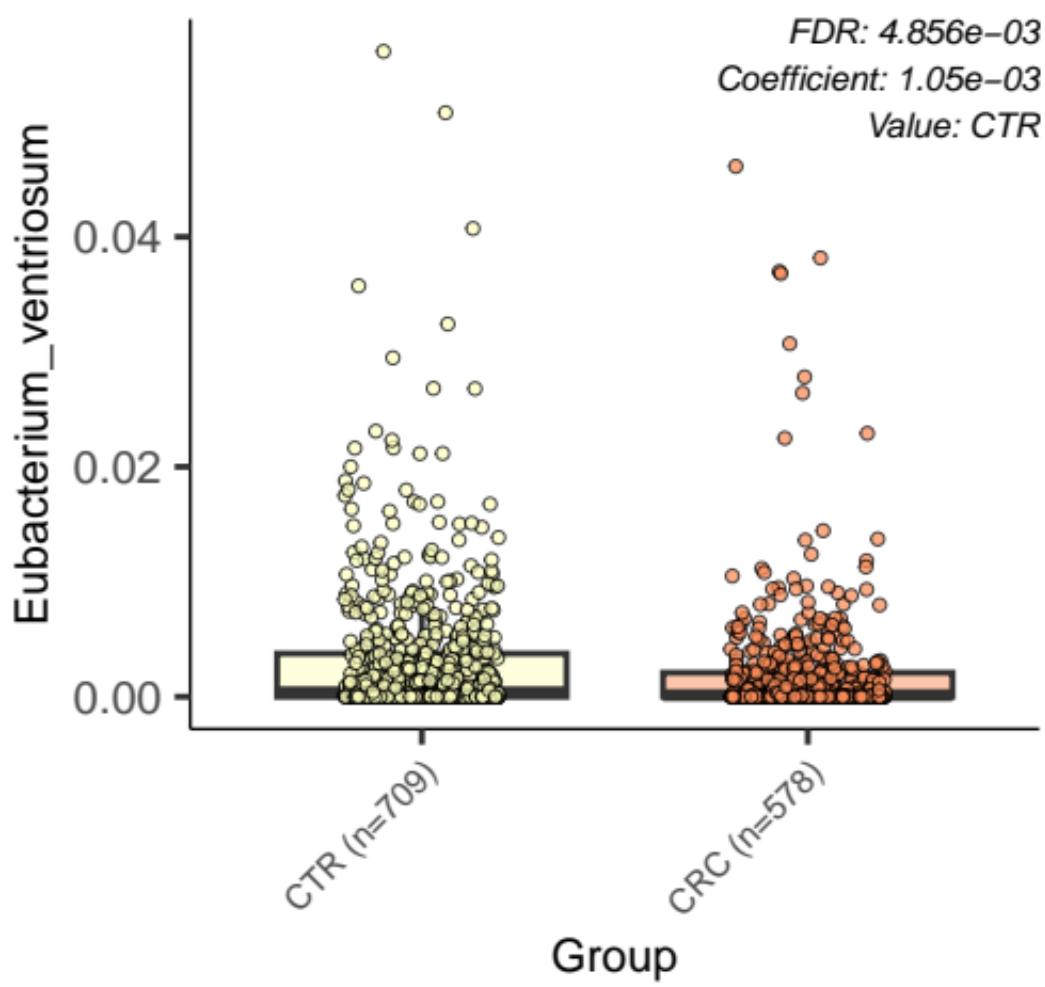


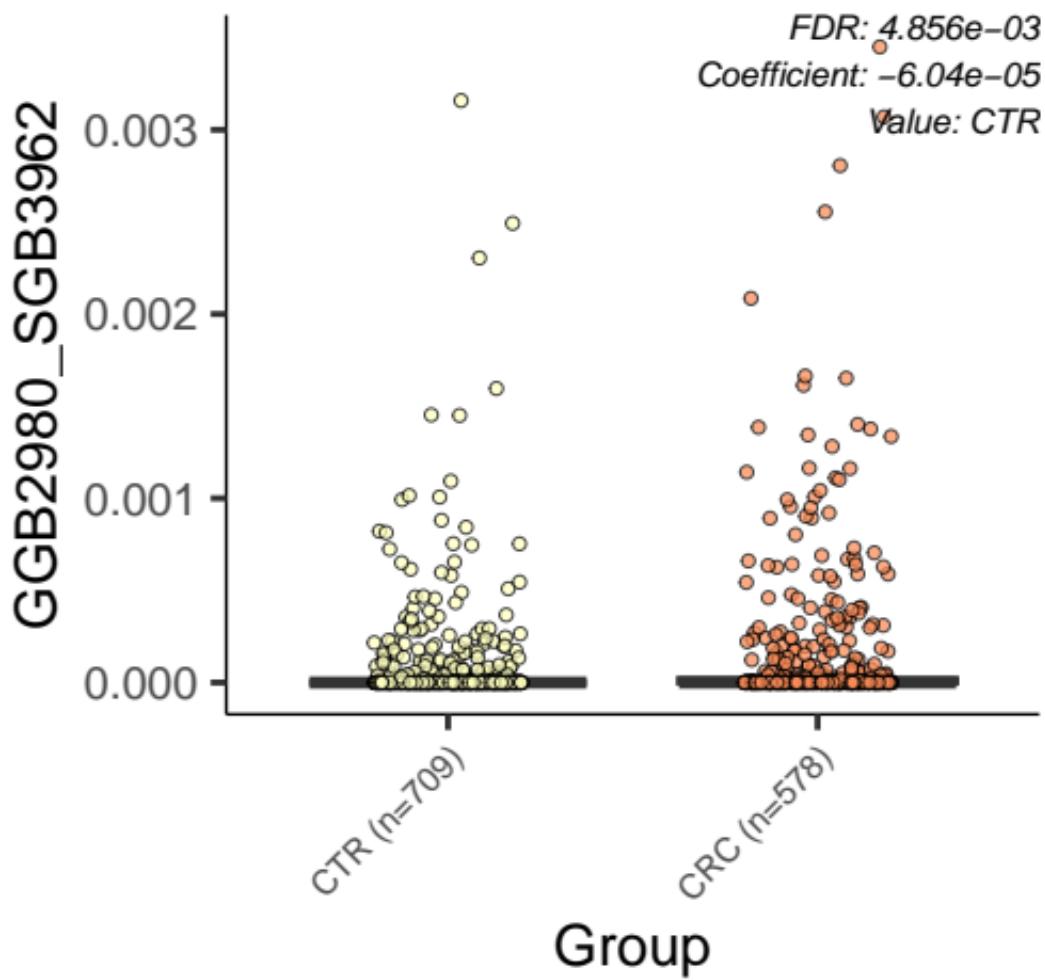


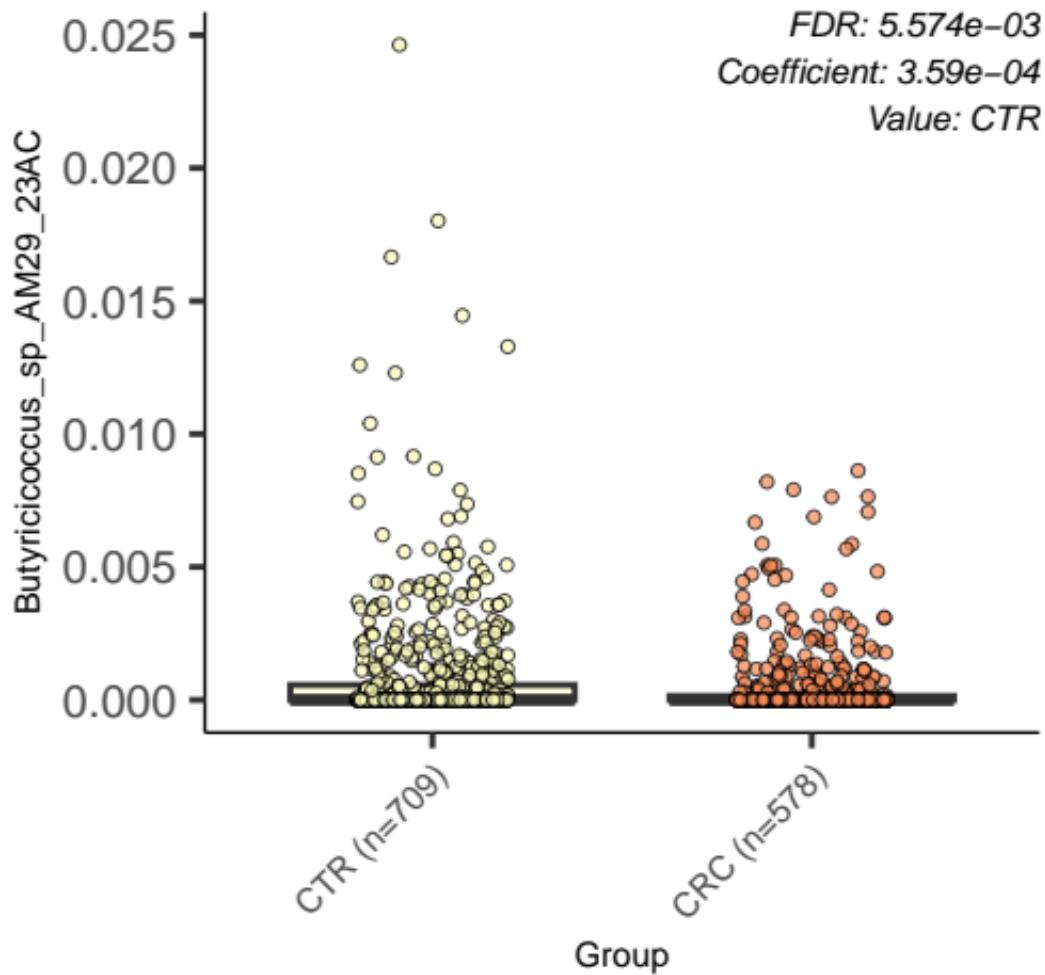


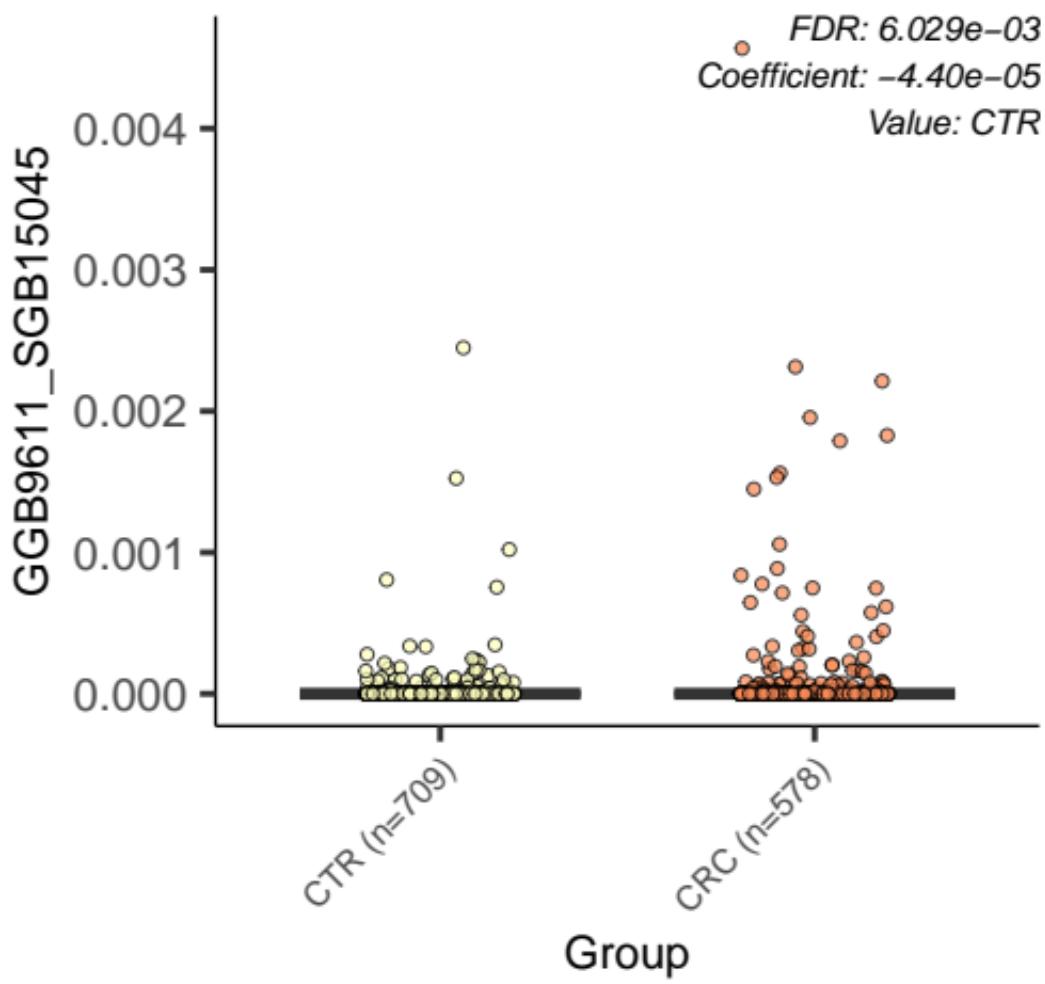


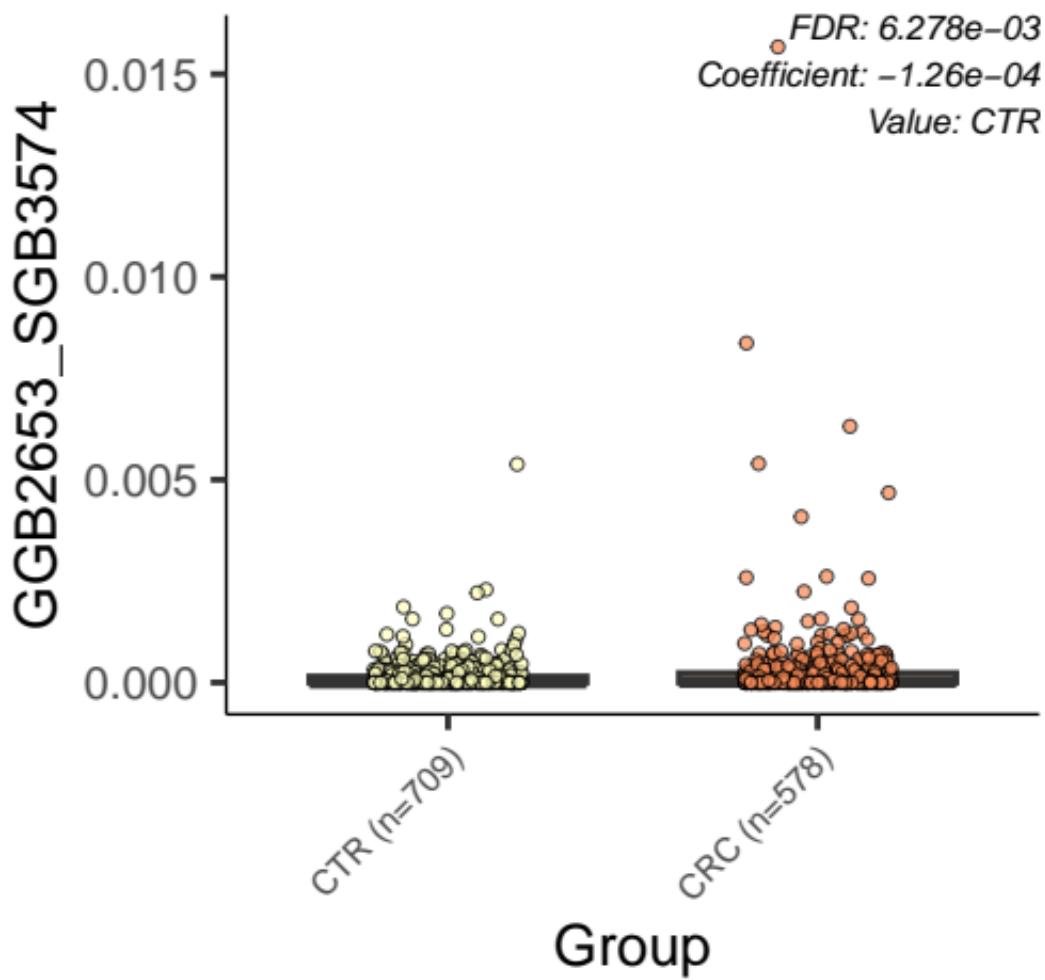


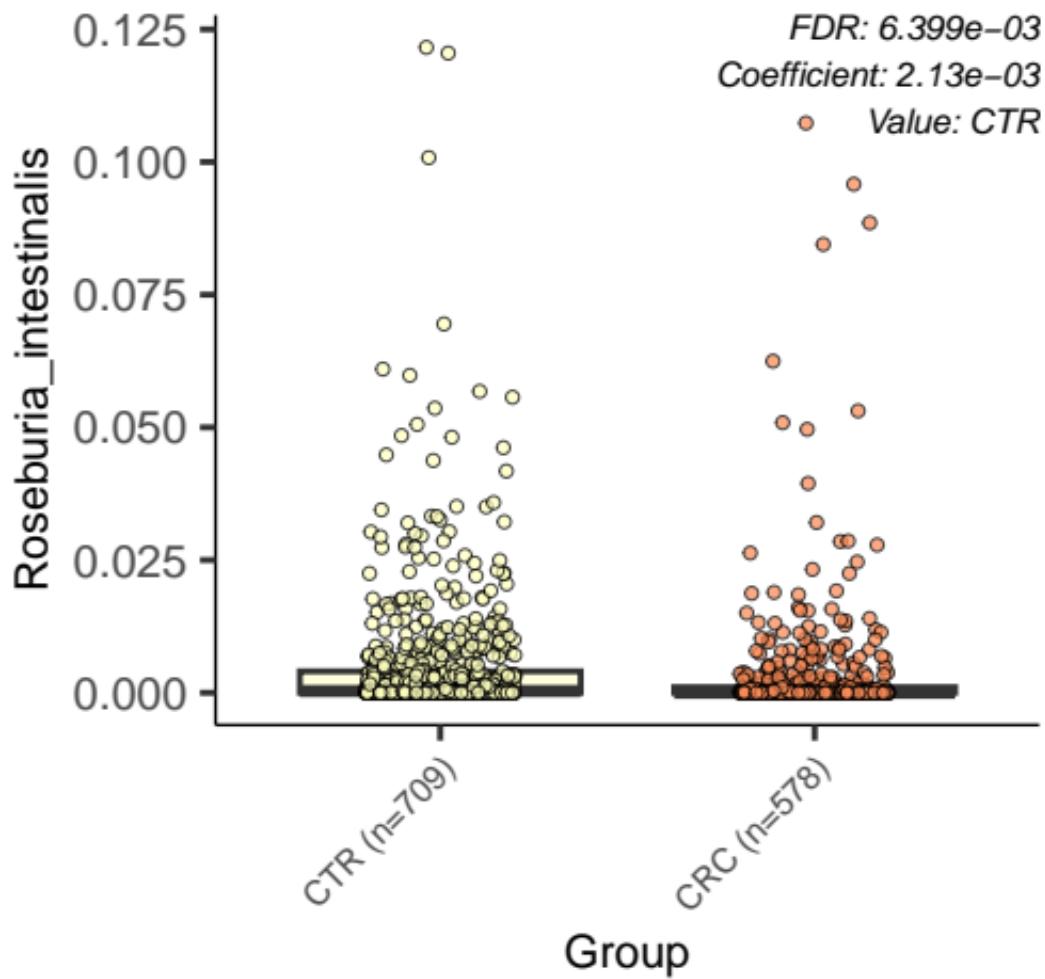


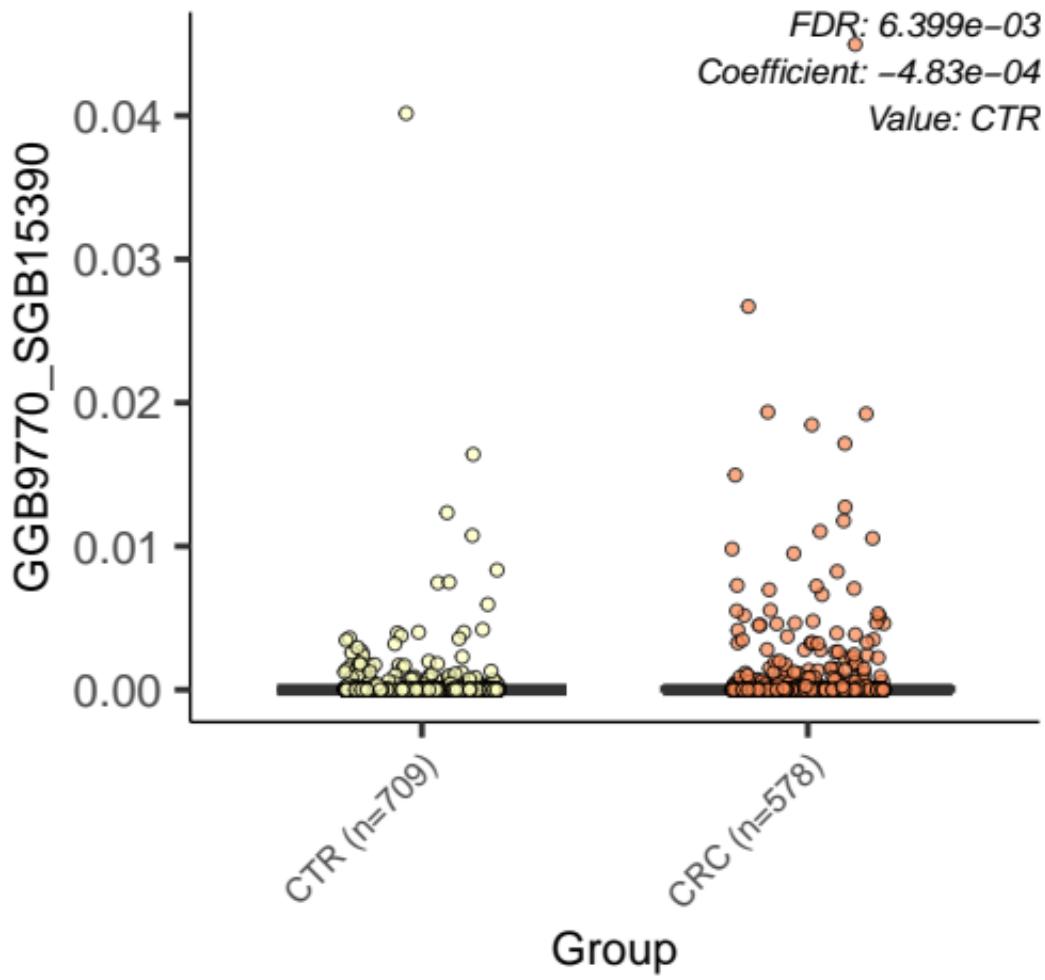


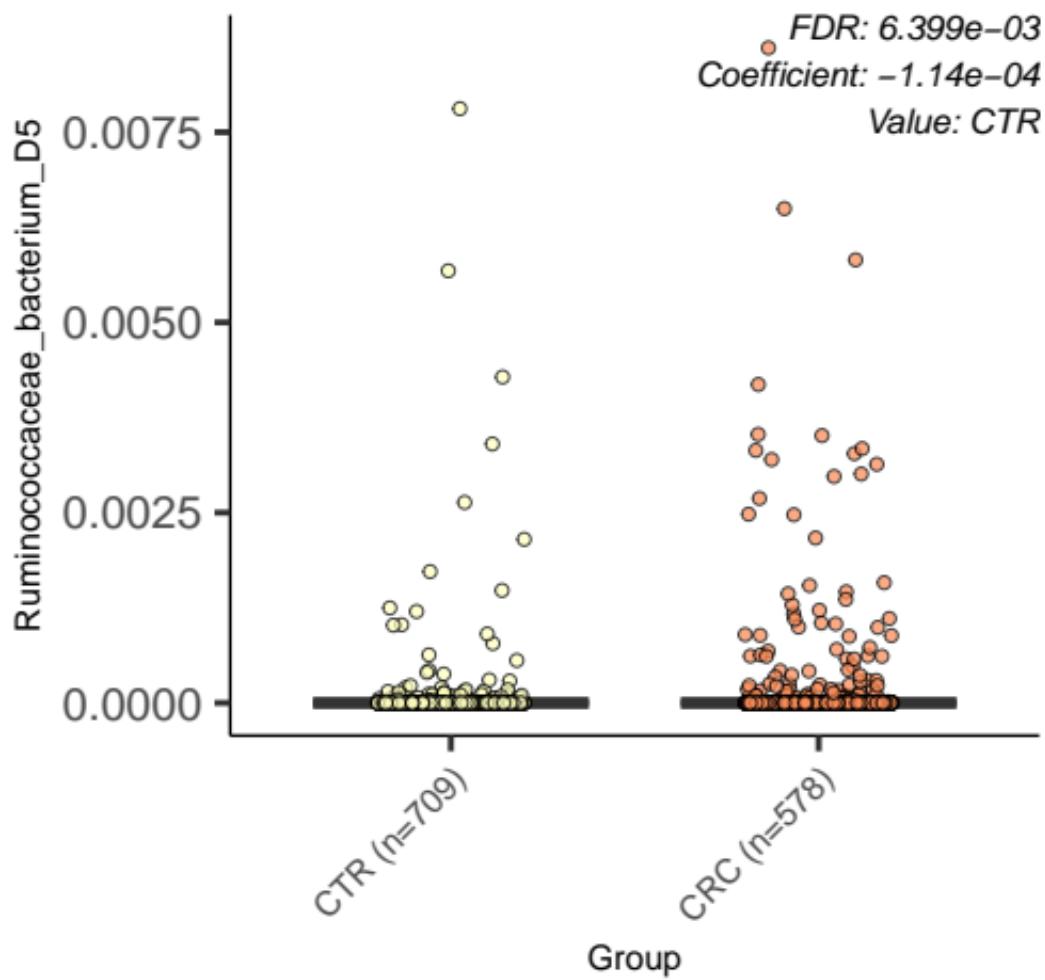


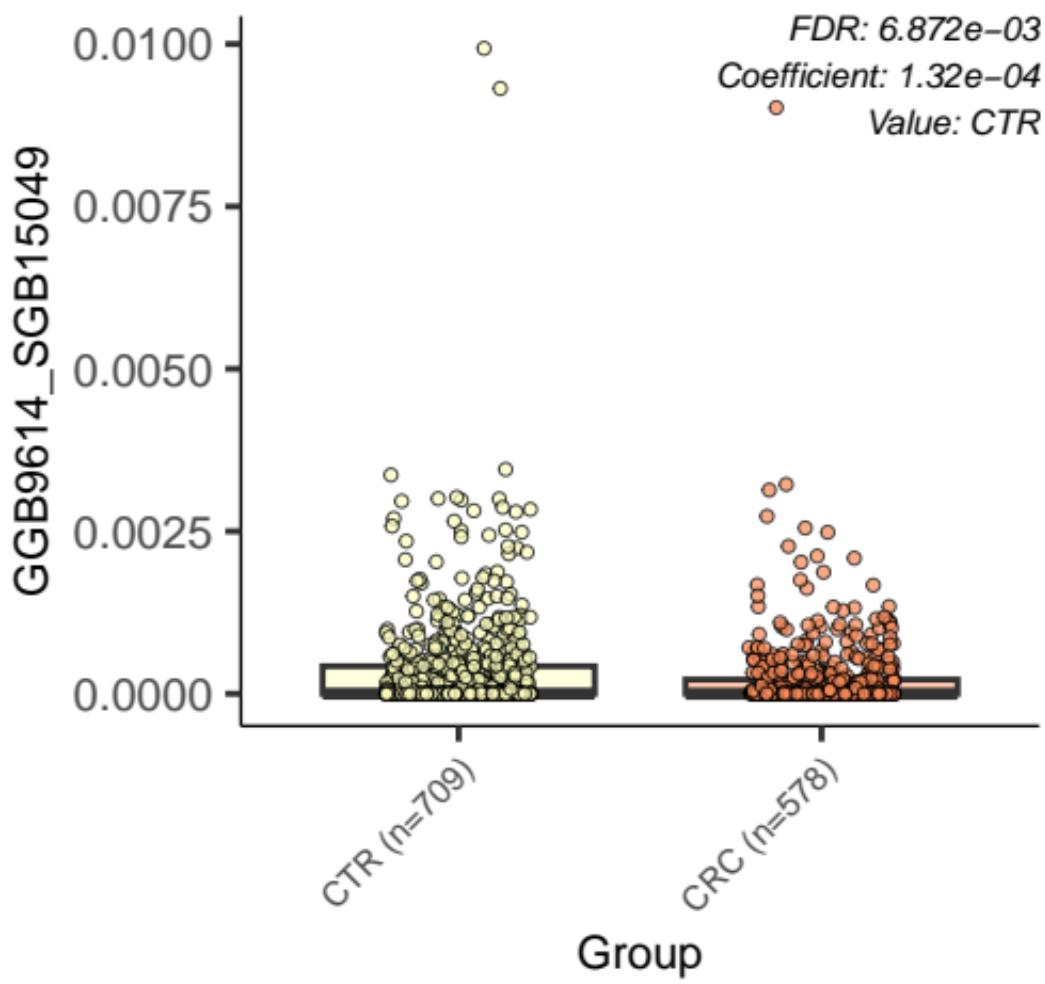


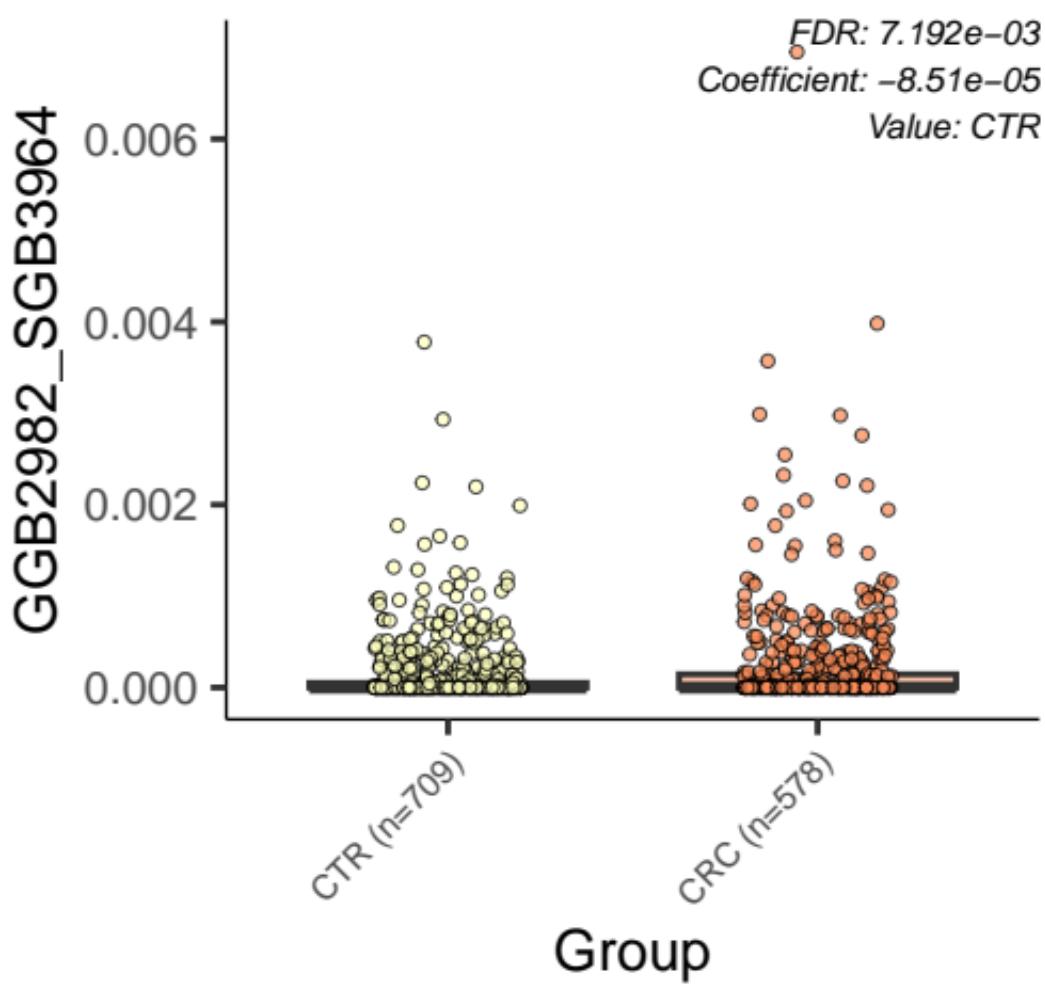


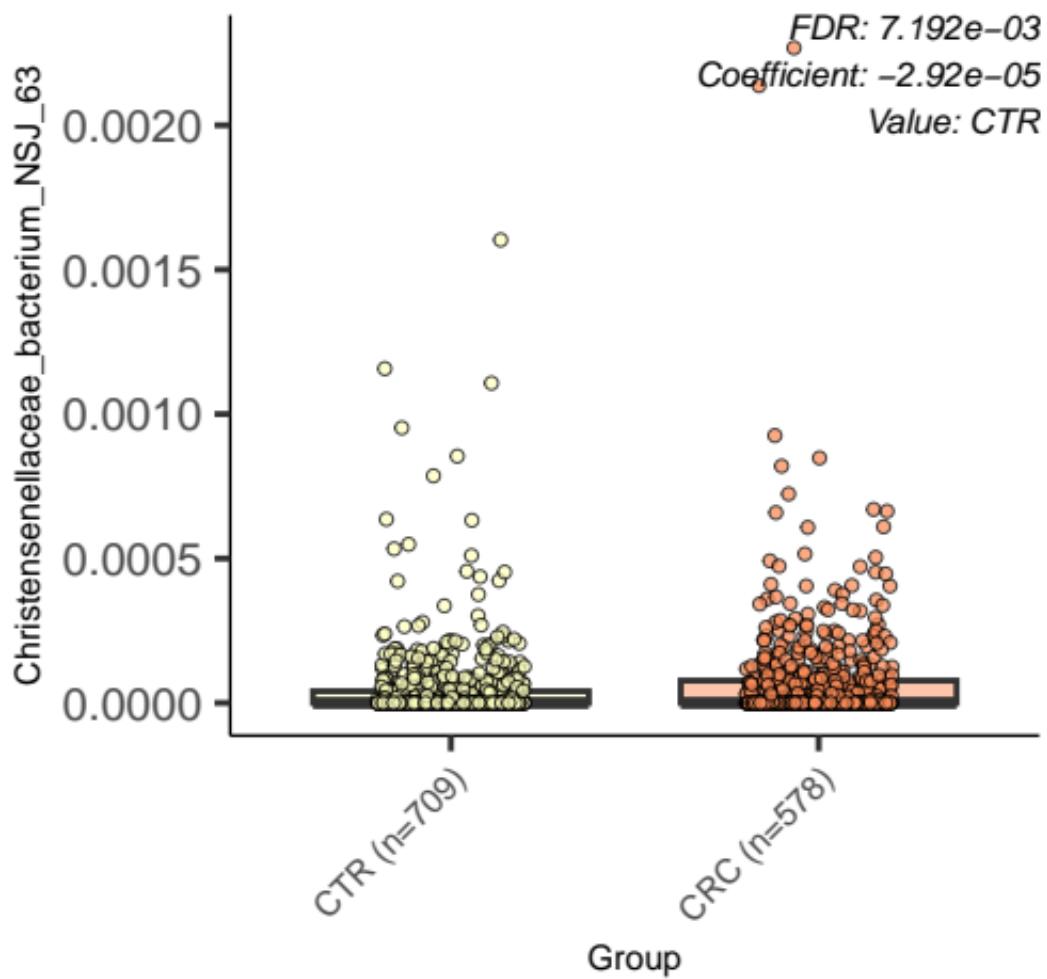


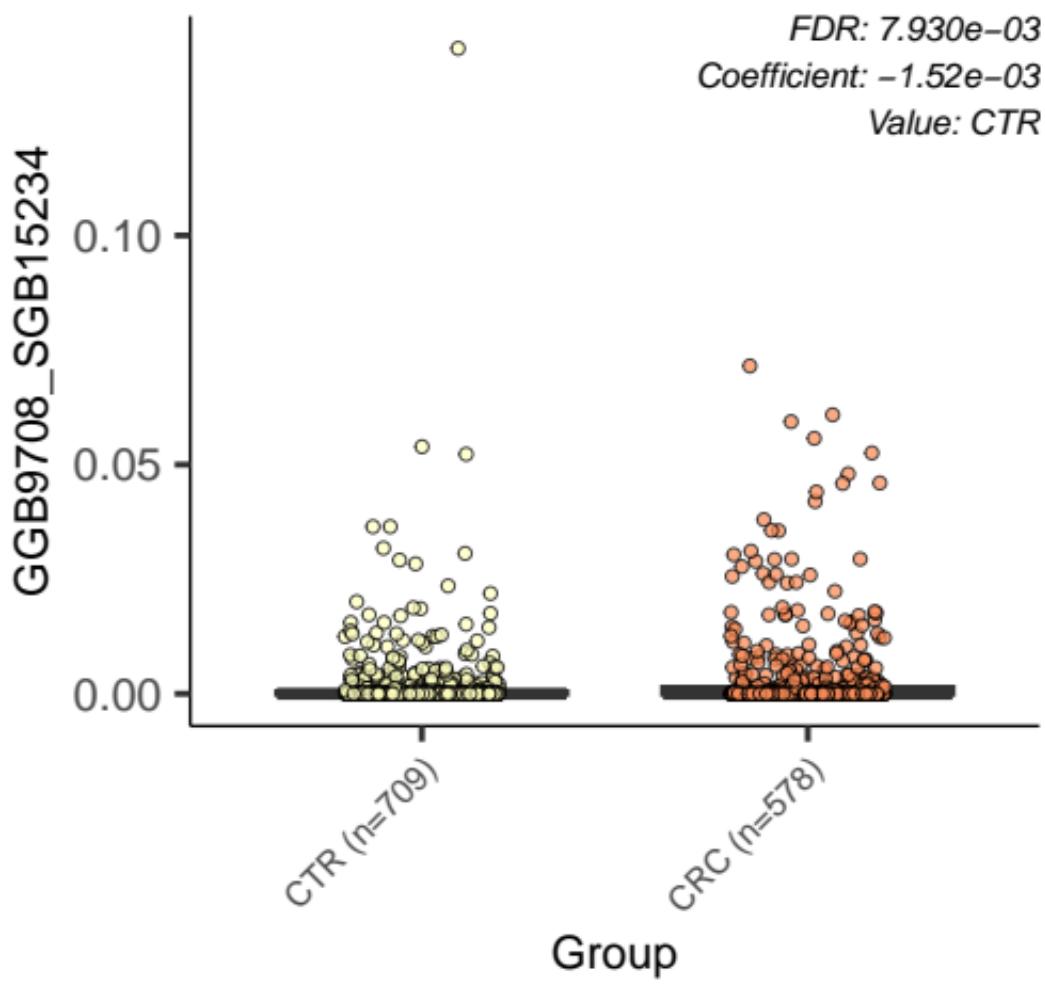


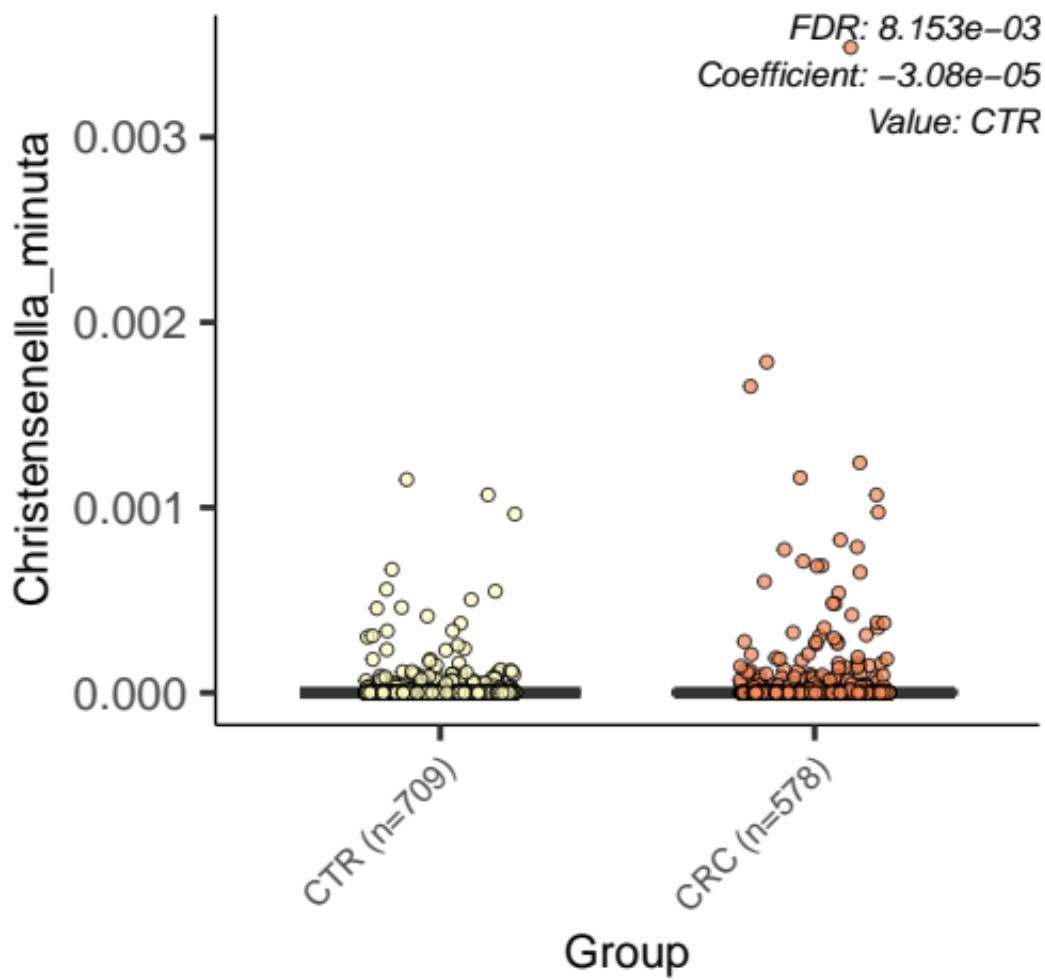


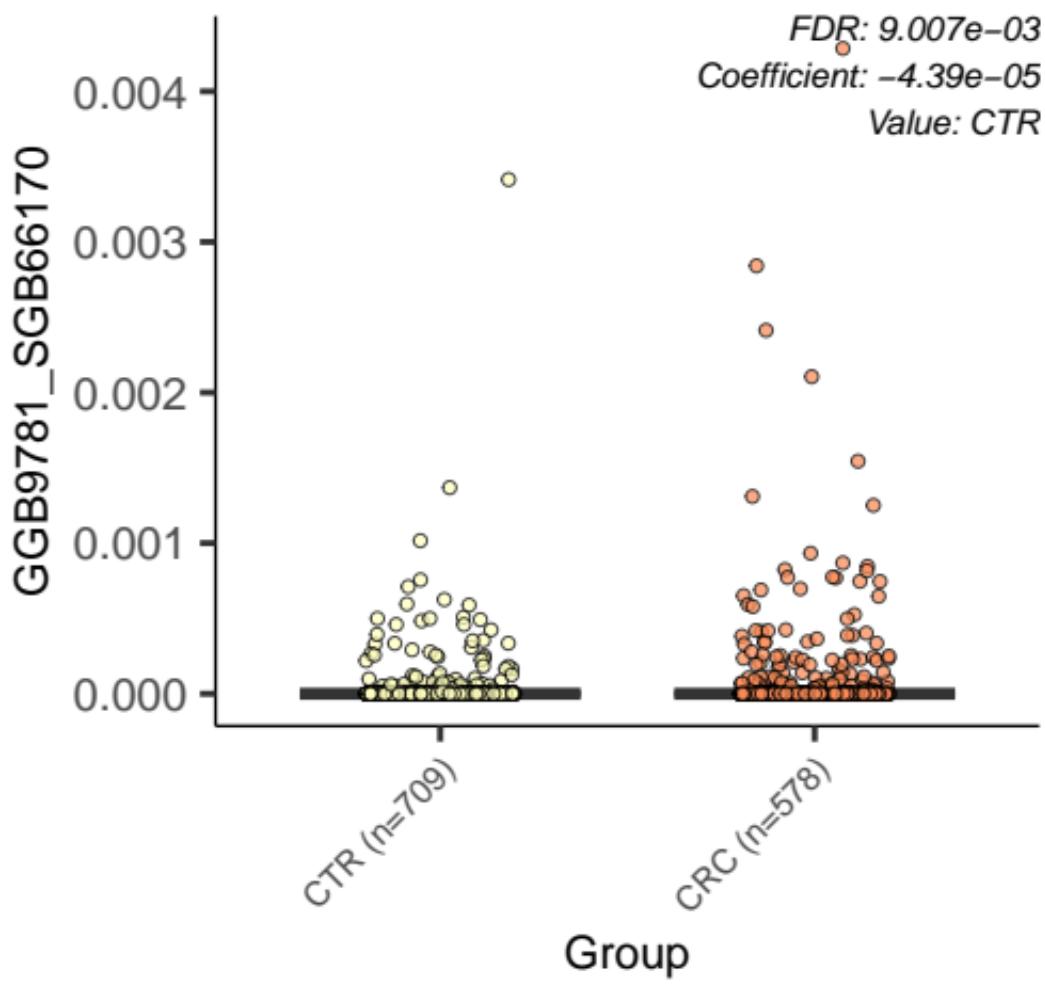


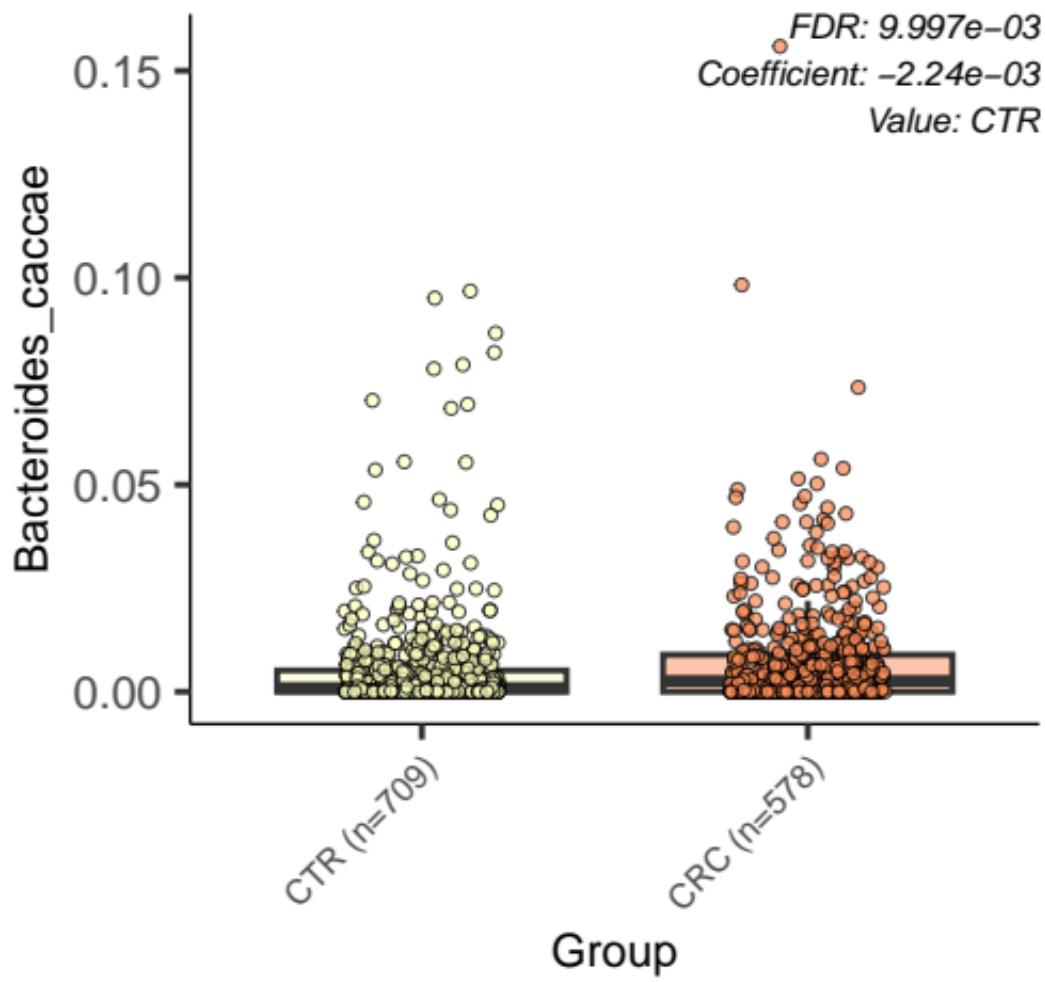


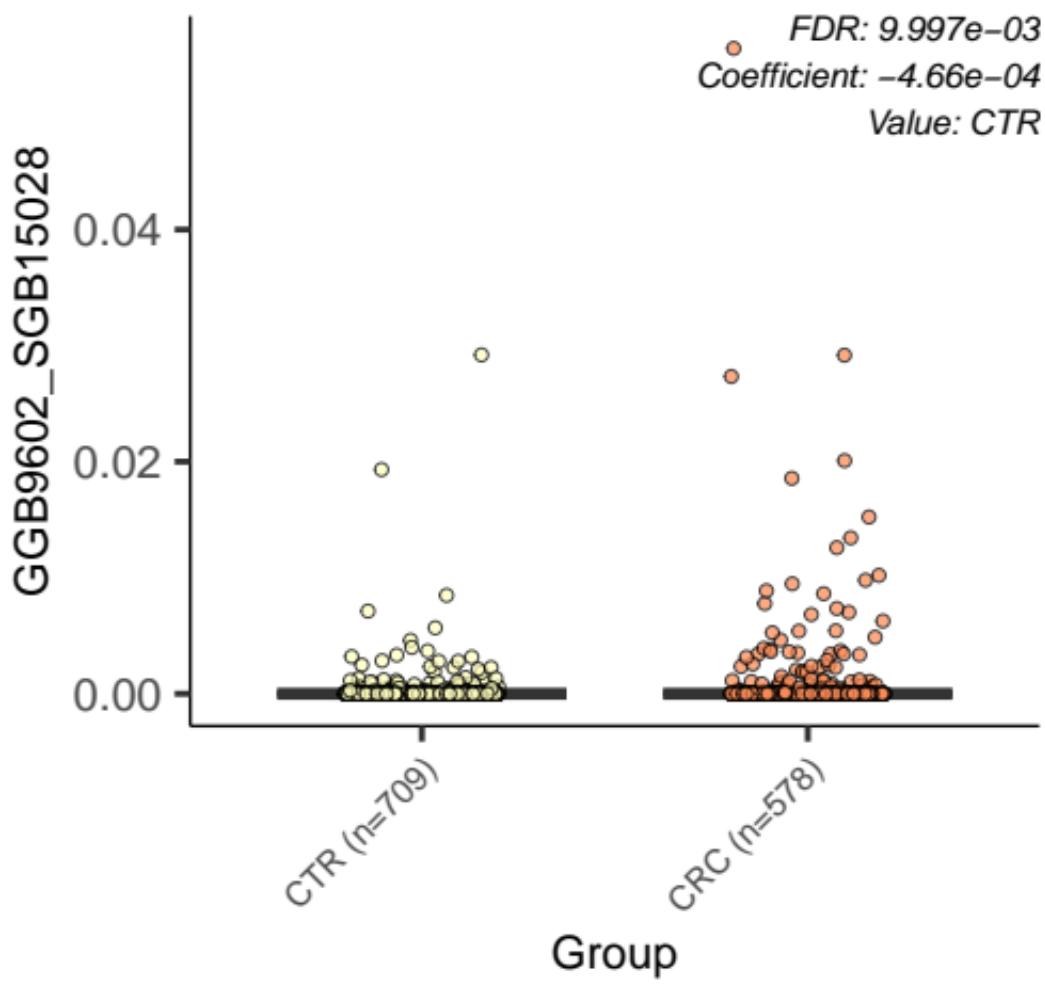


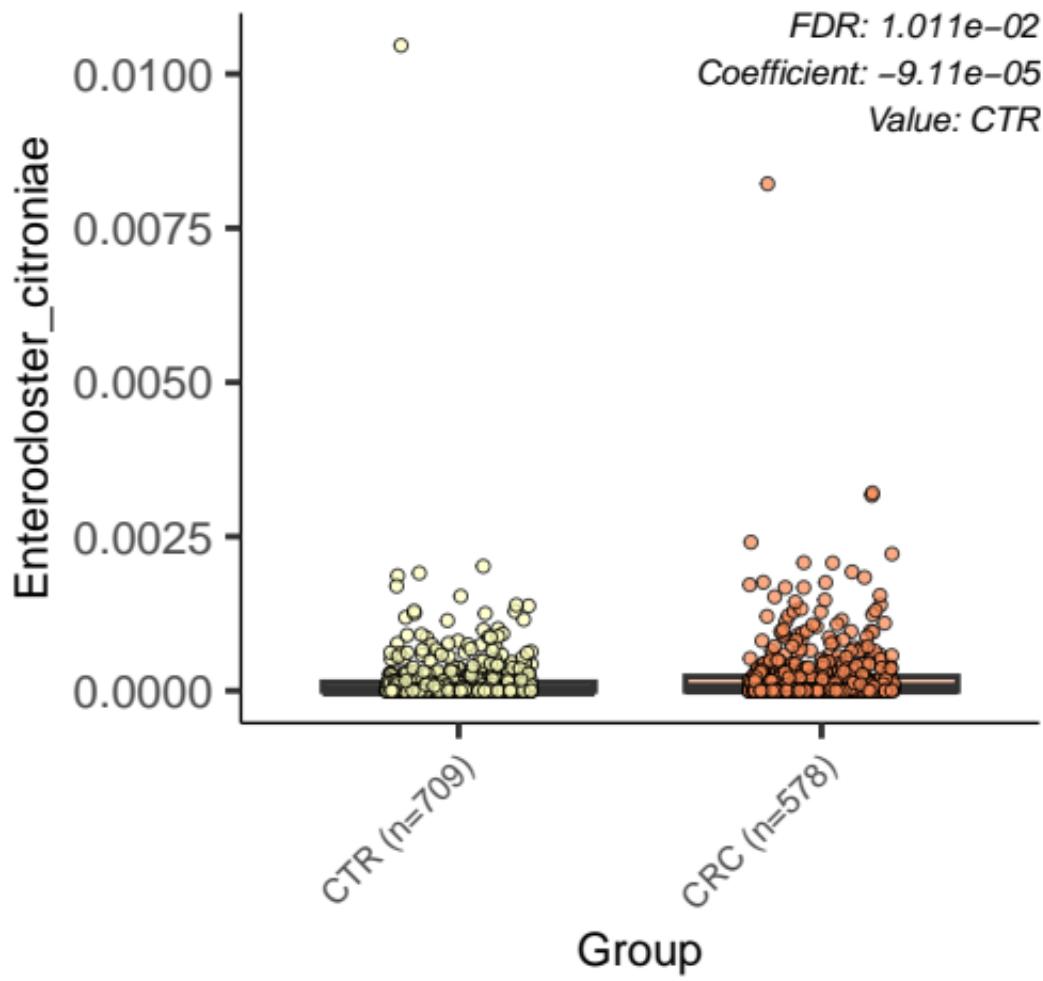


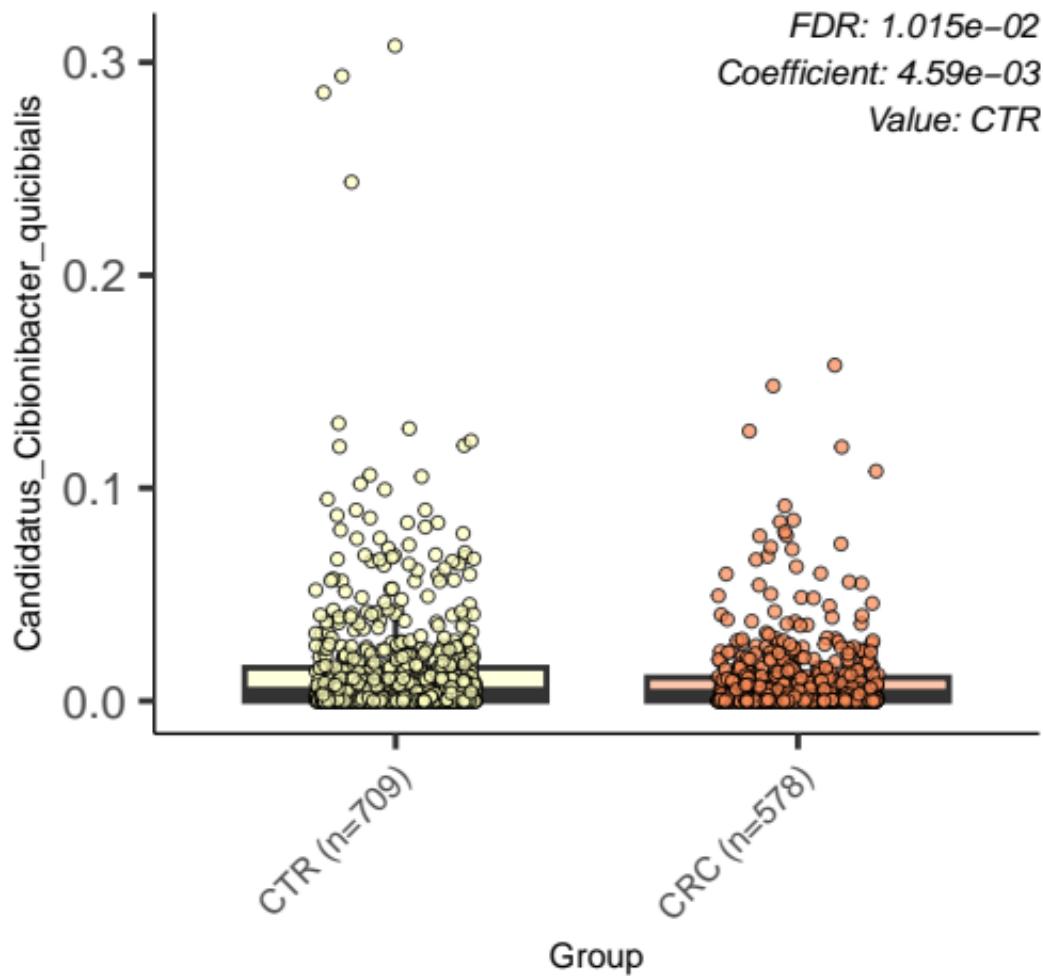


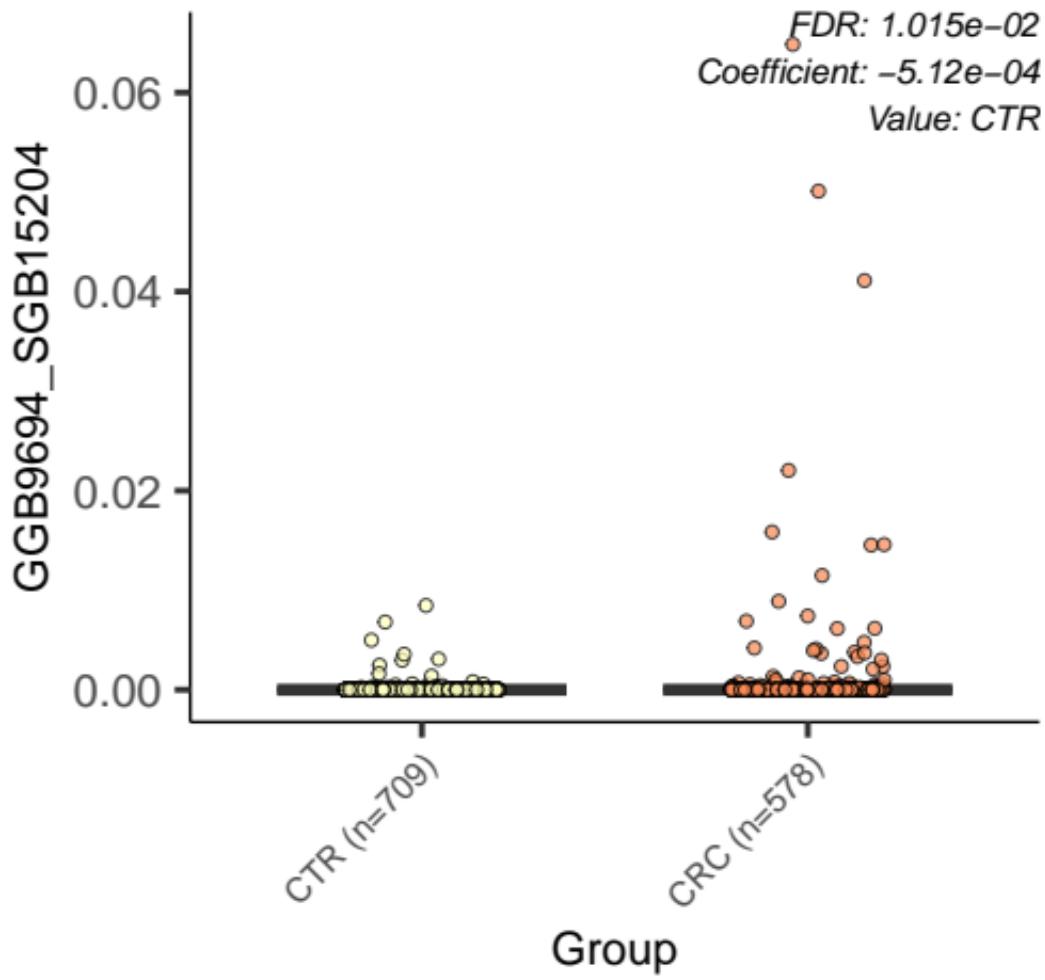












Clostridiales_bacterium_KLE1615

FDR: 1.020e-02
Coefficient: 5.94e-04
Value: CTR

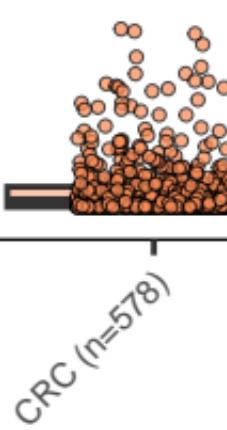
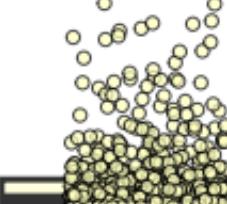
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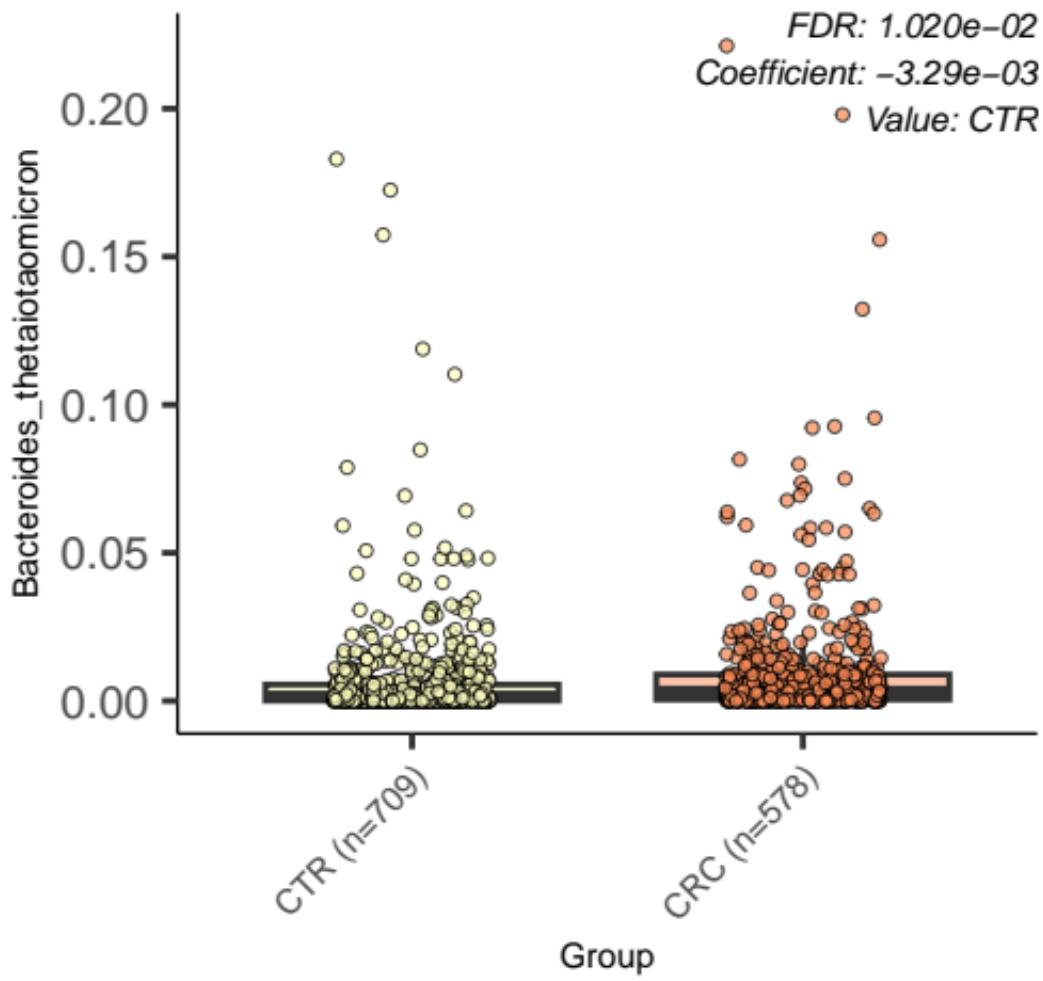
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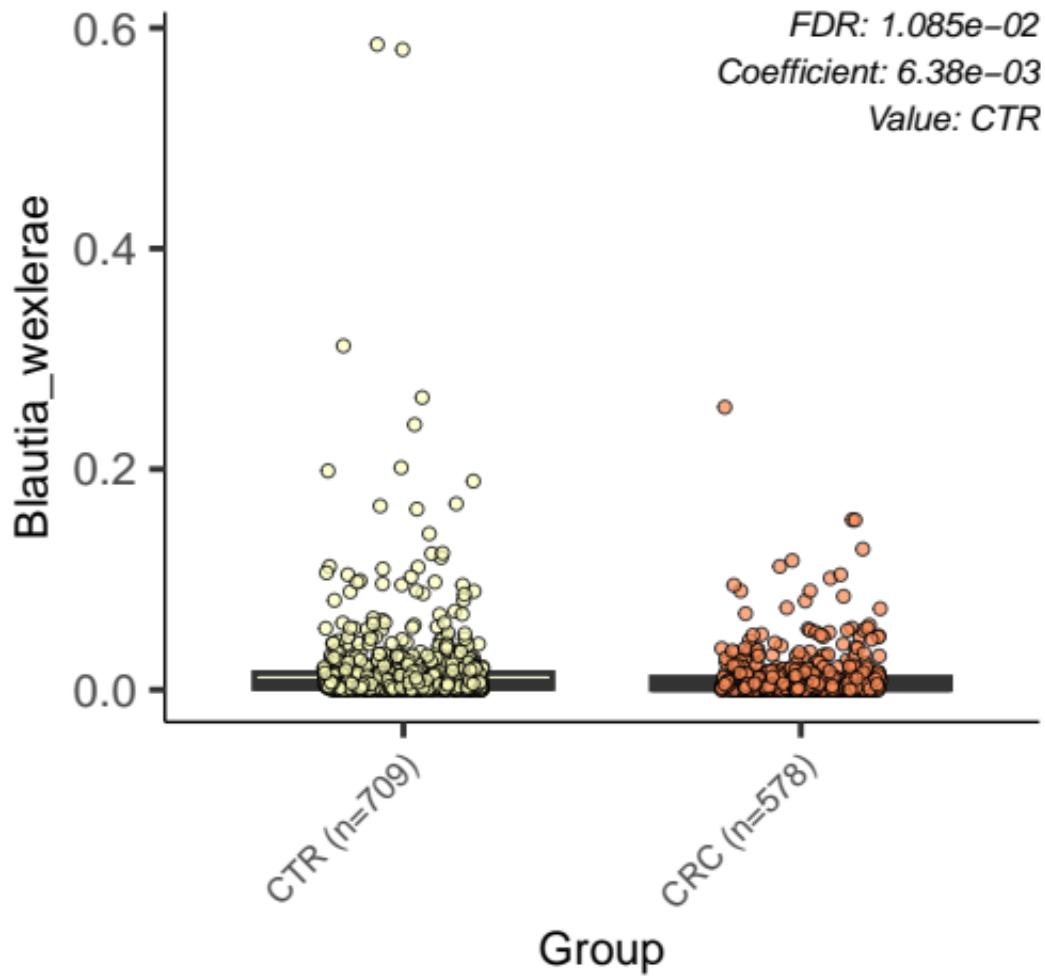
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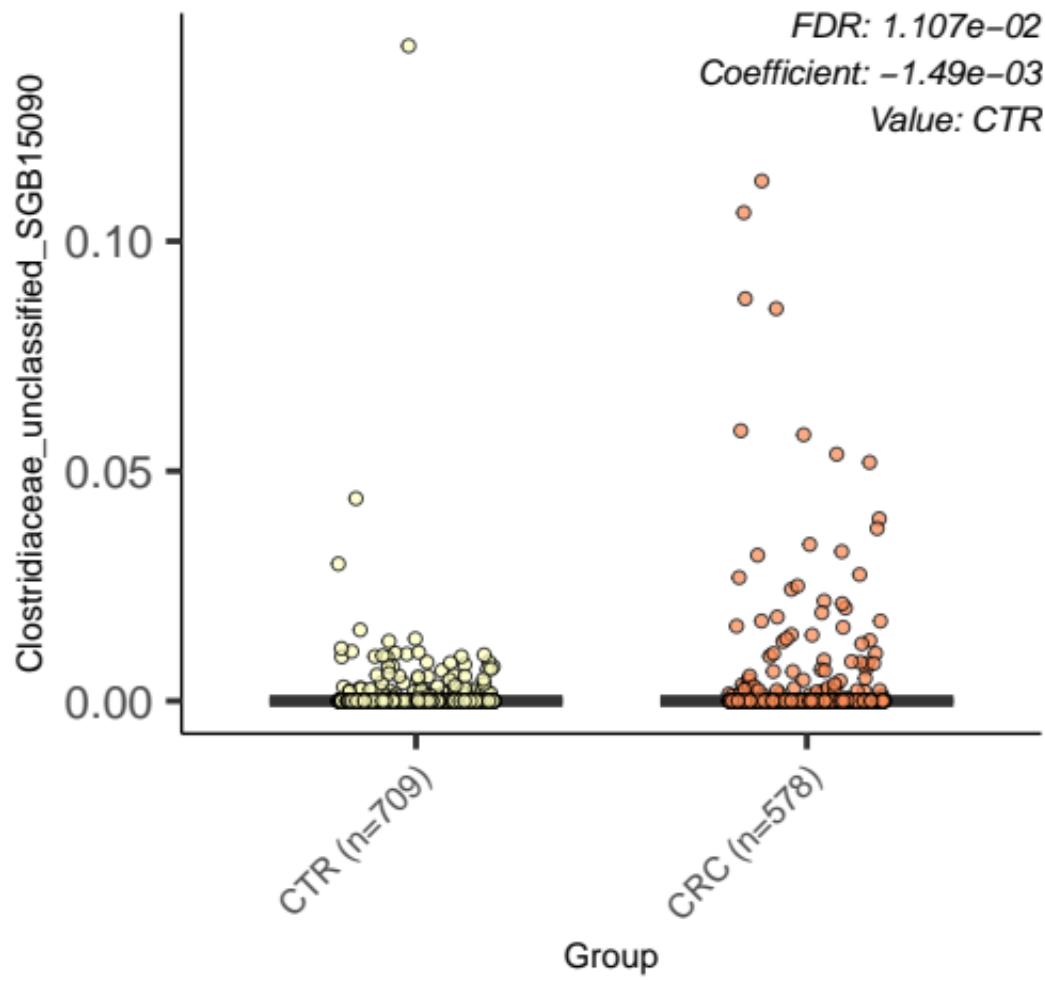
CTR (n=709)

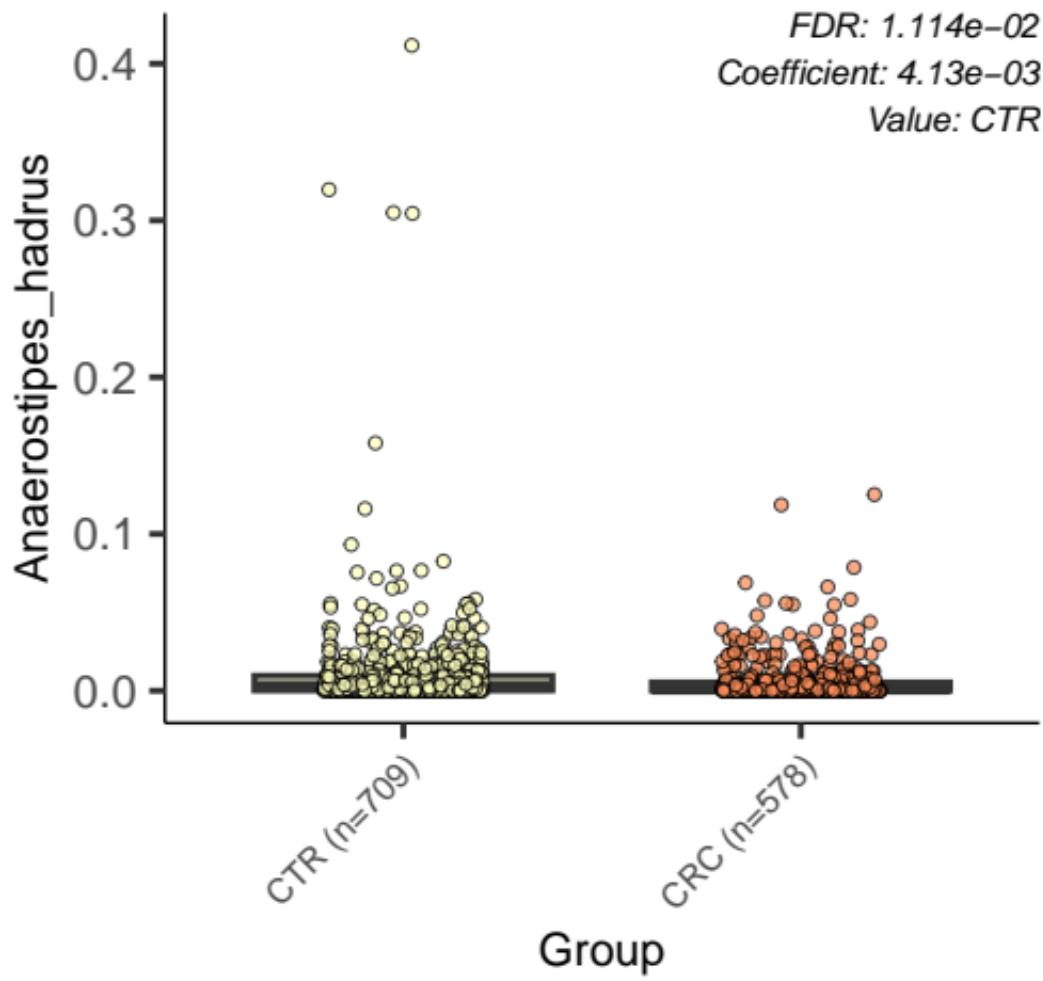
Group

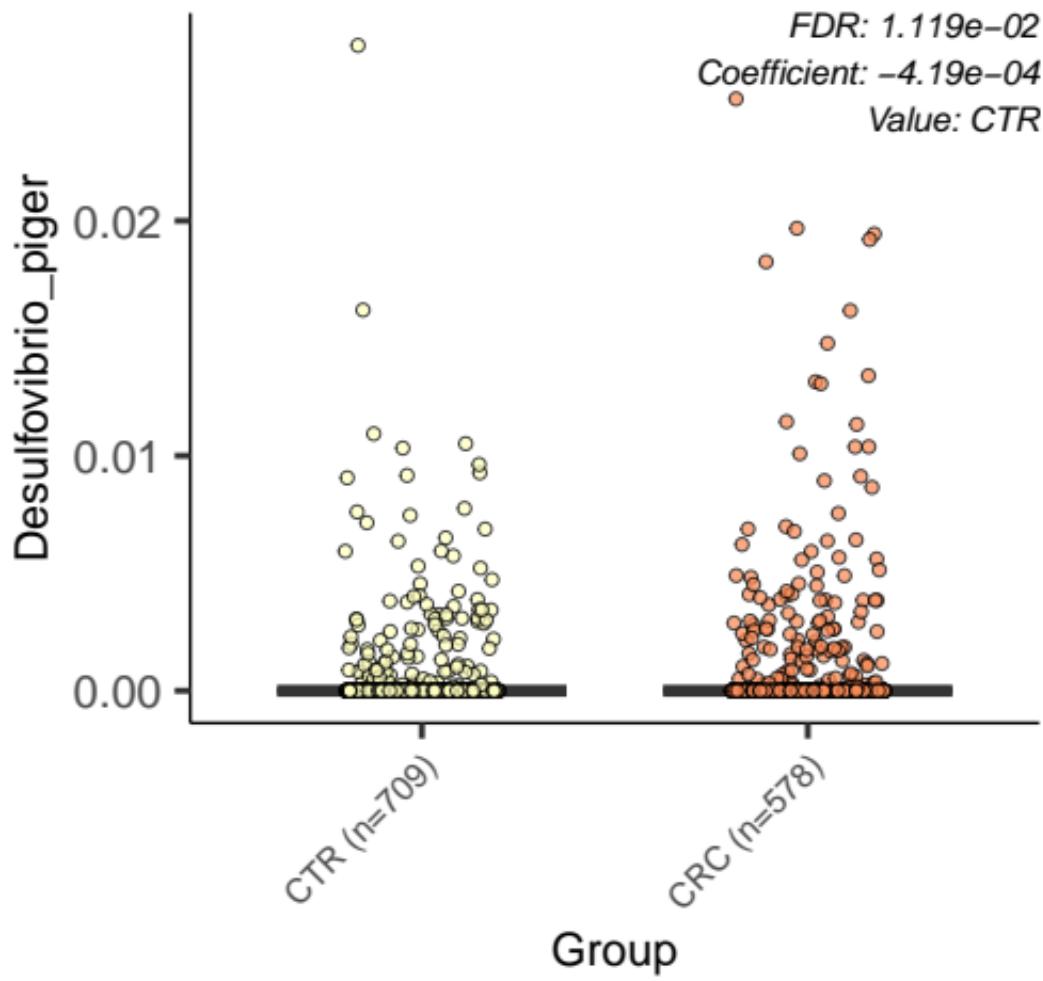












GGB3033_SGB4029

FDR: 1.172e-02
Coefficient: -1.35e-04
Value: CTR

0.010

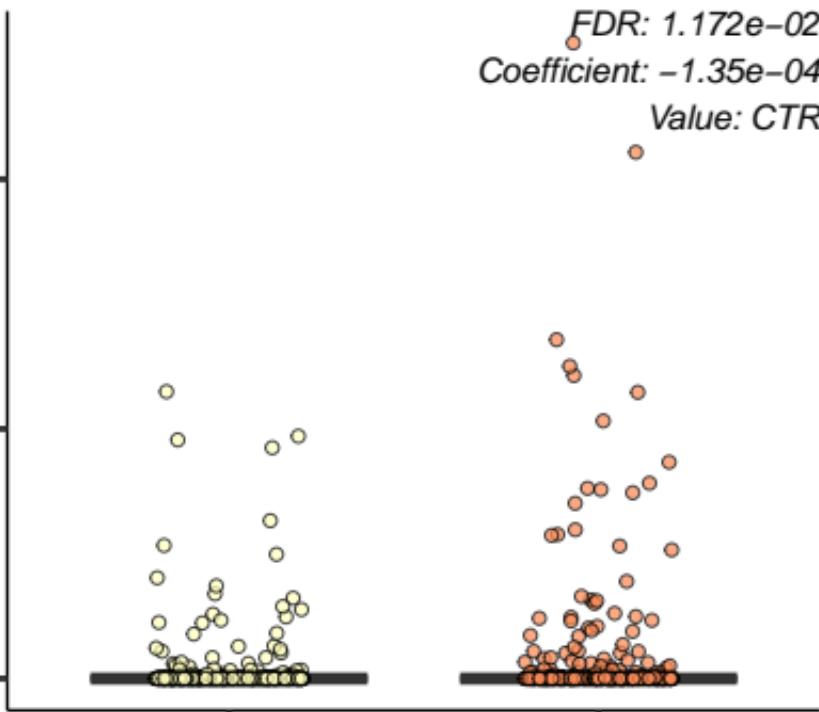
0.005

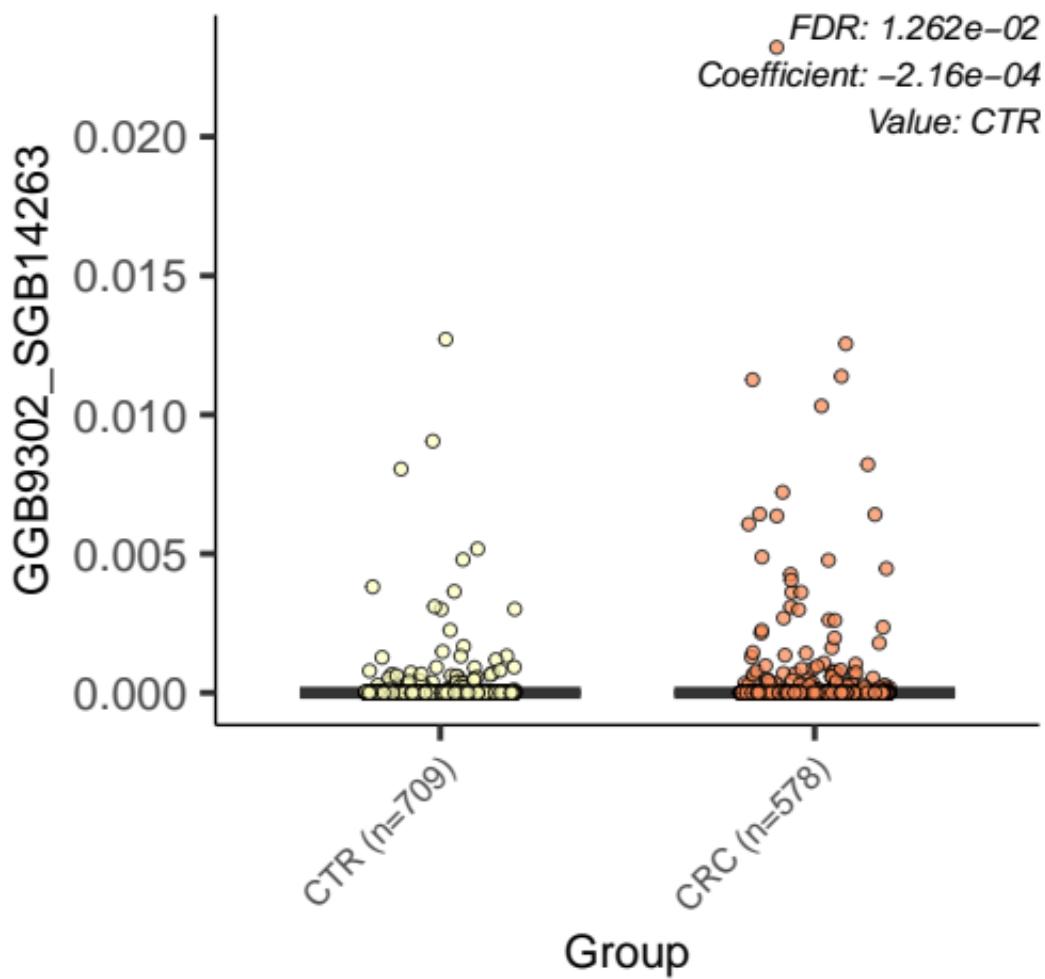
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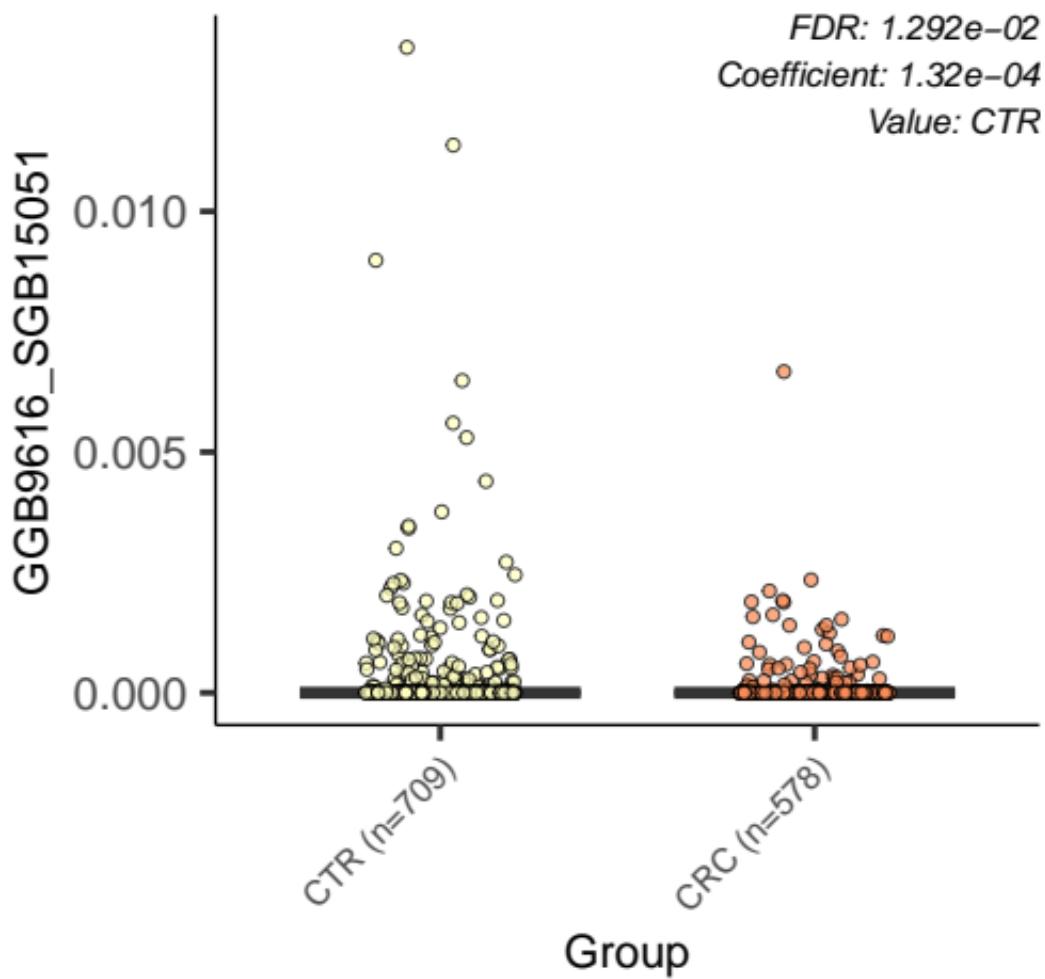
CTR (n=709)

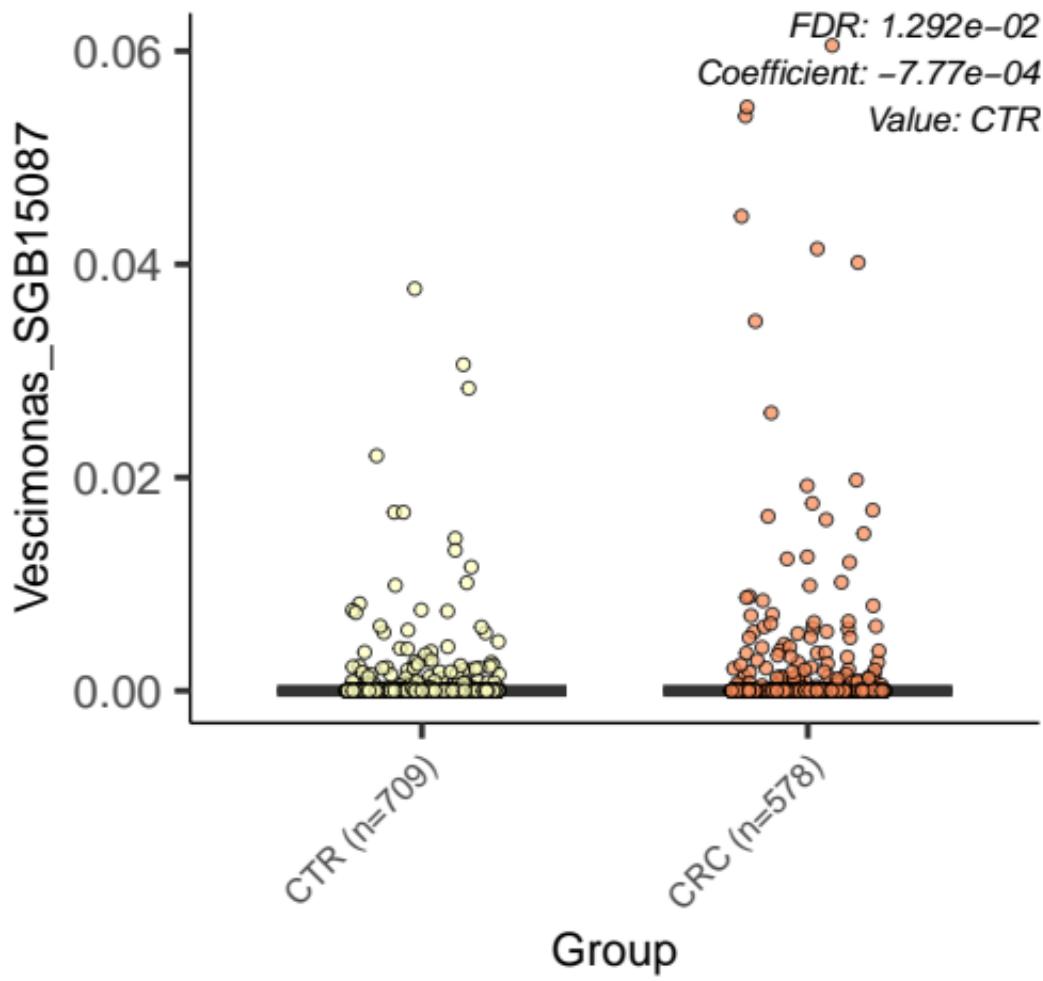
CRC (n=578)

Group

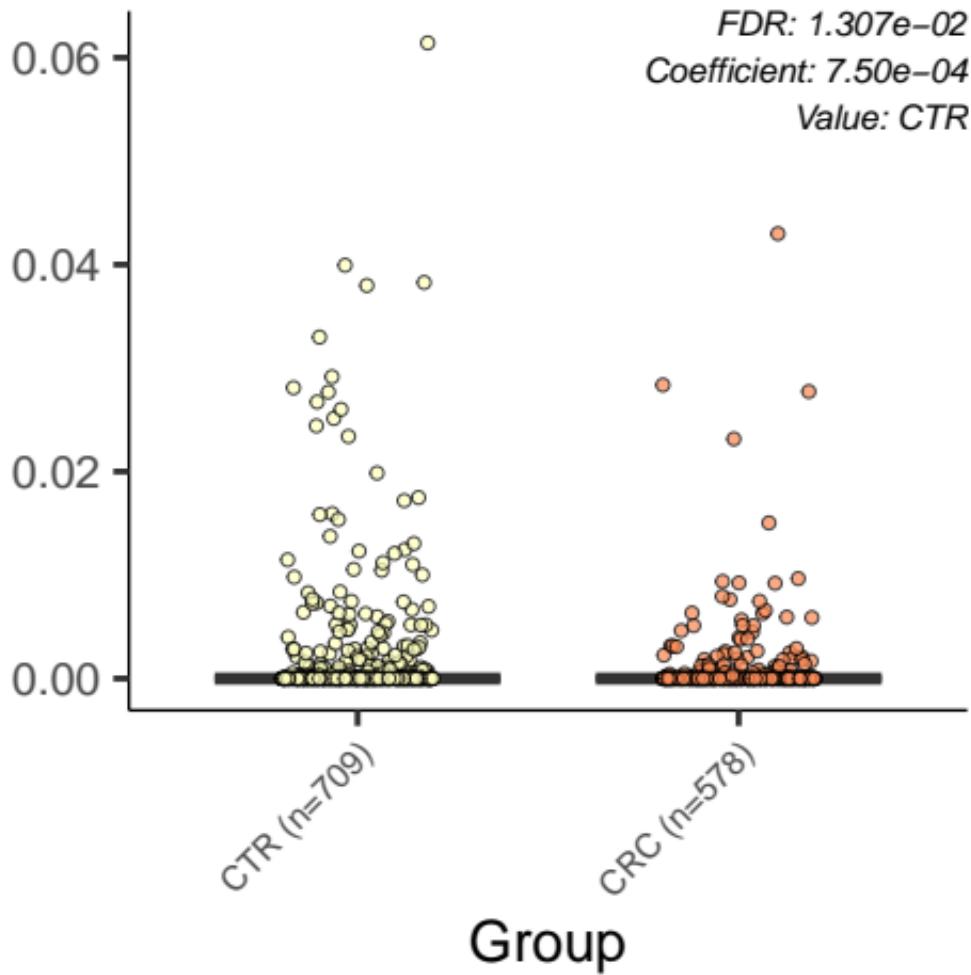


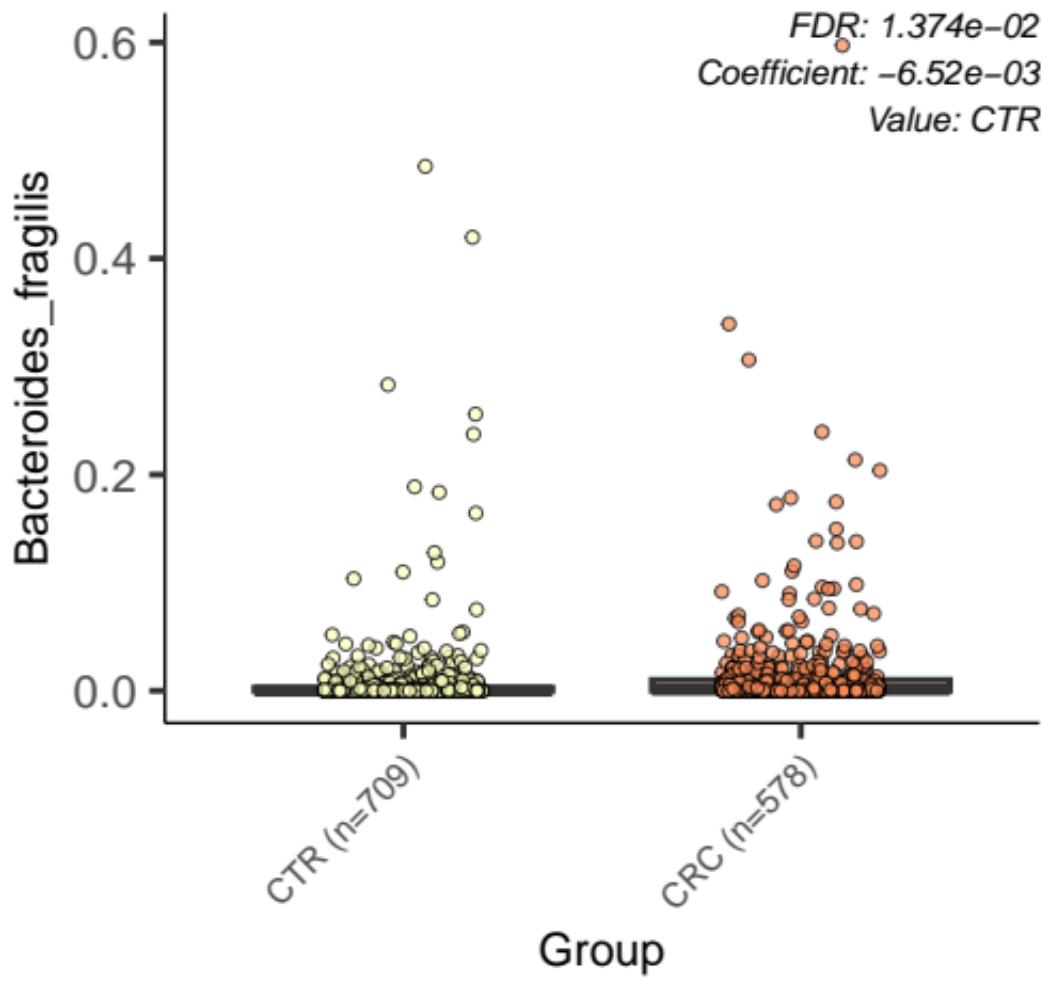


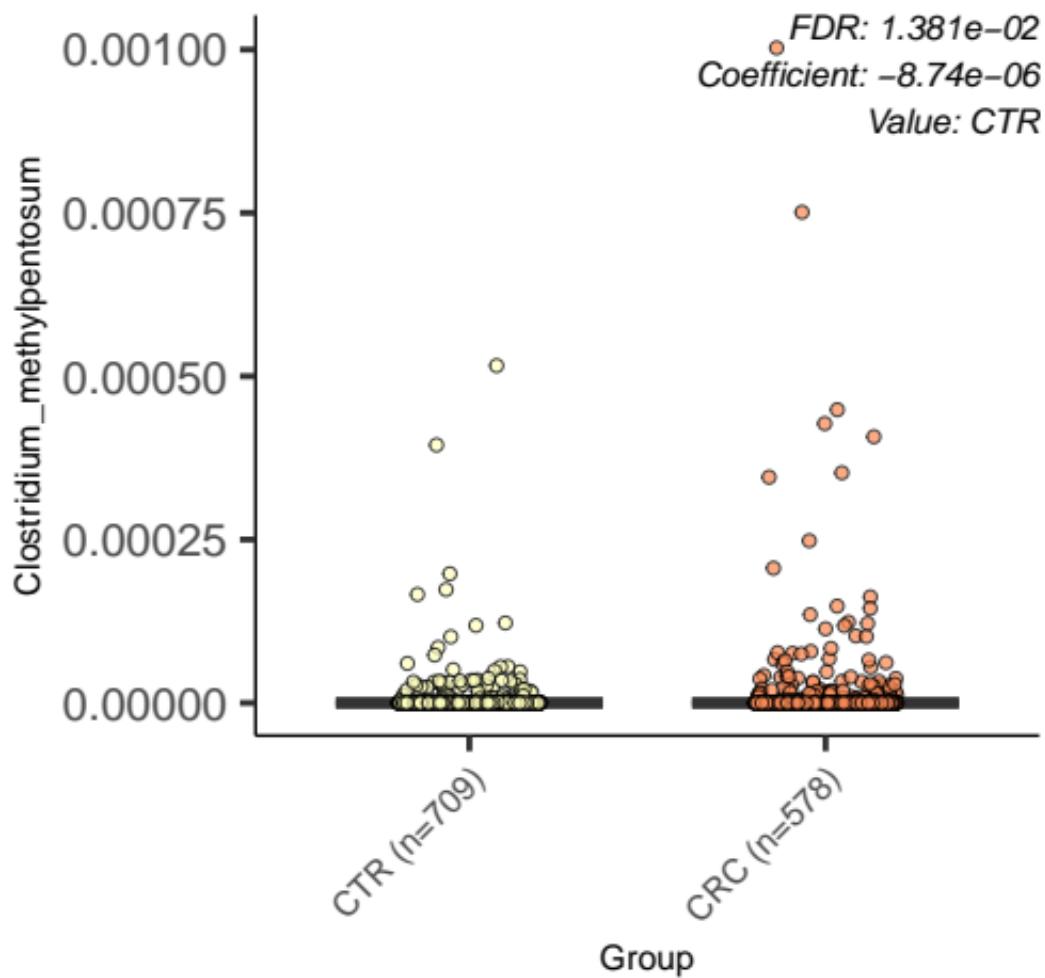


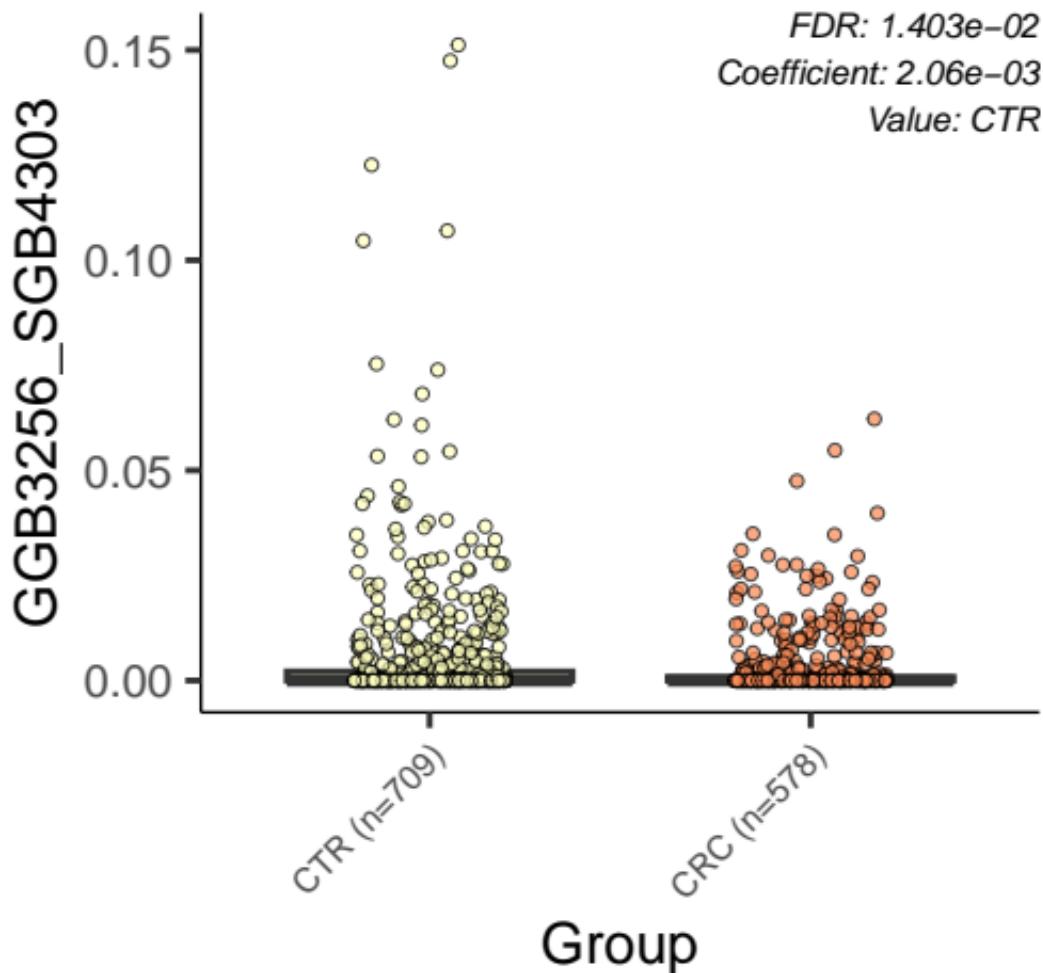


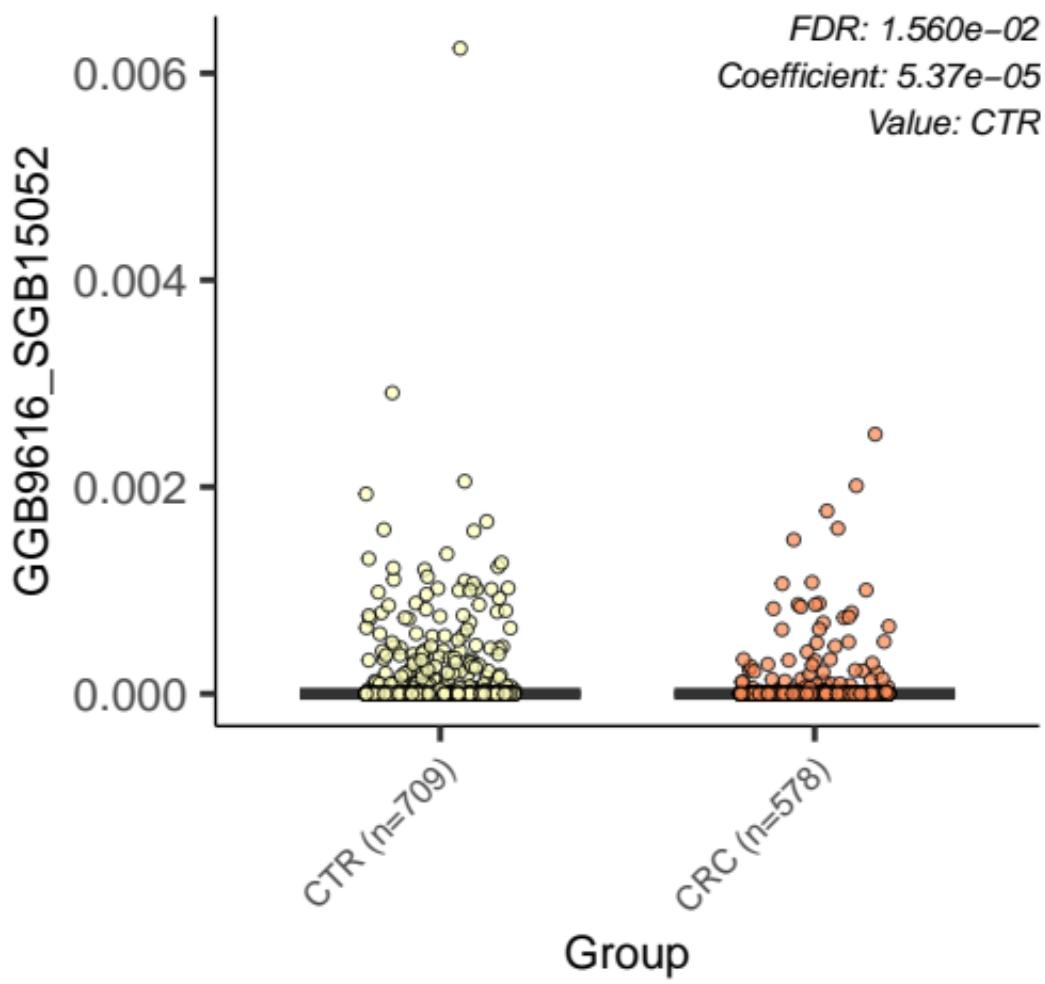
GGB3175_SGB4191

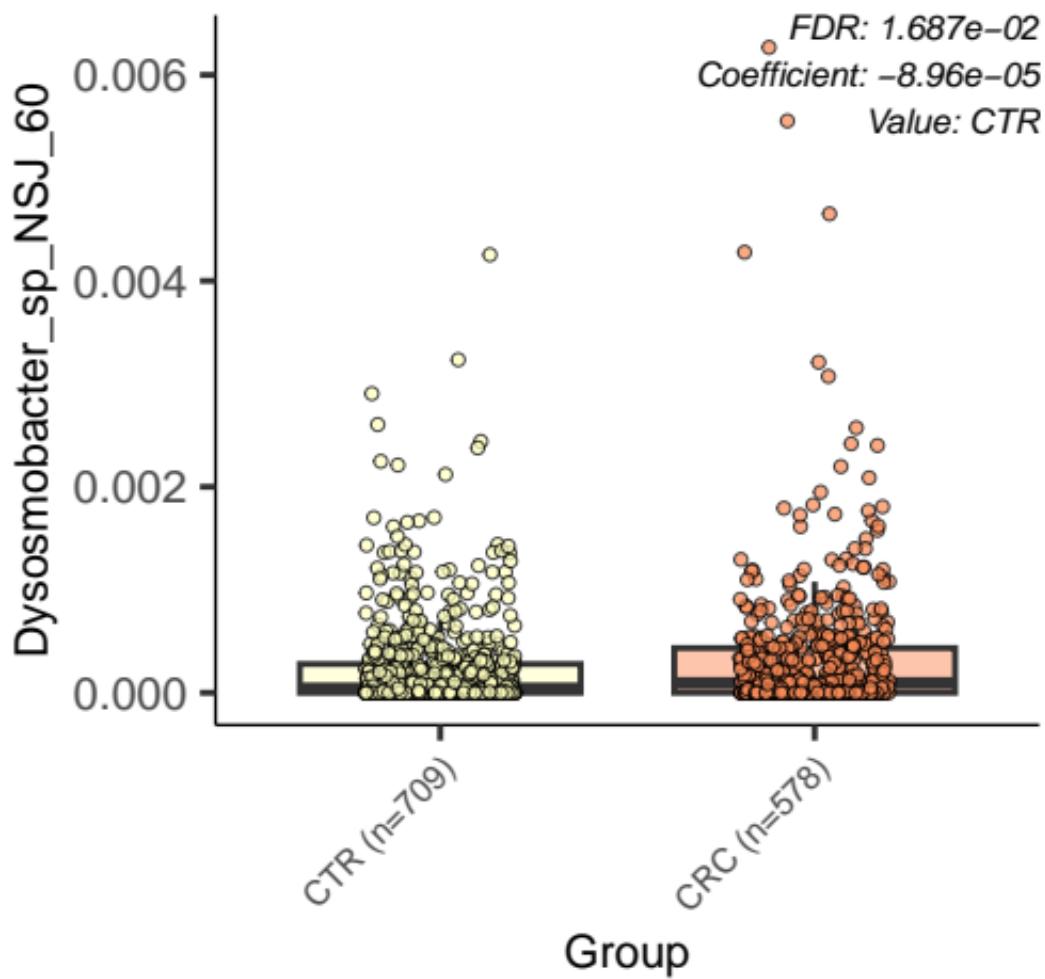


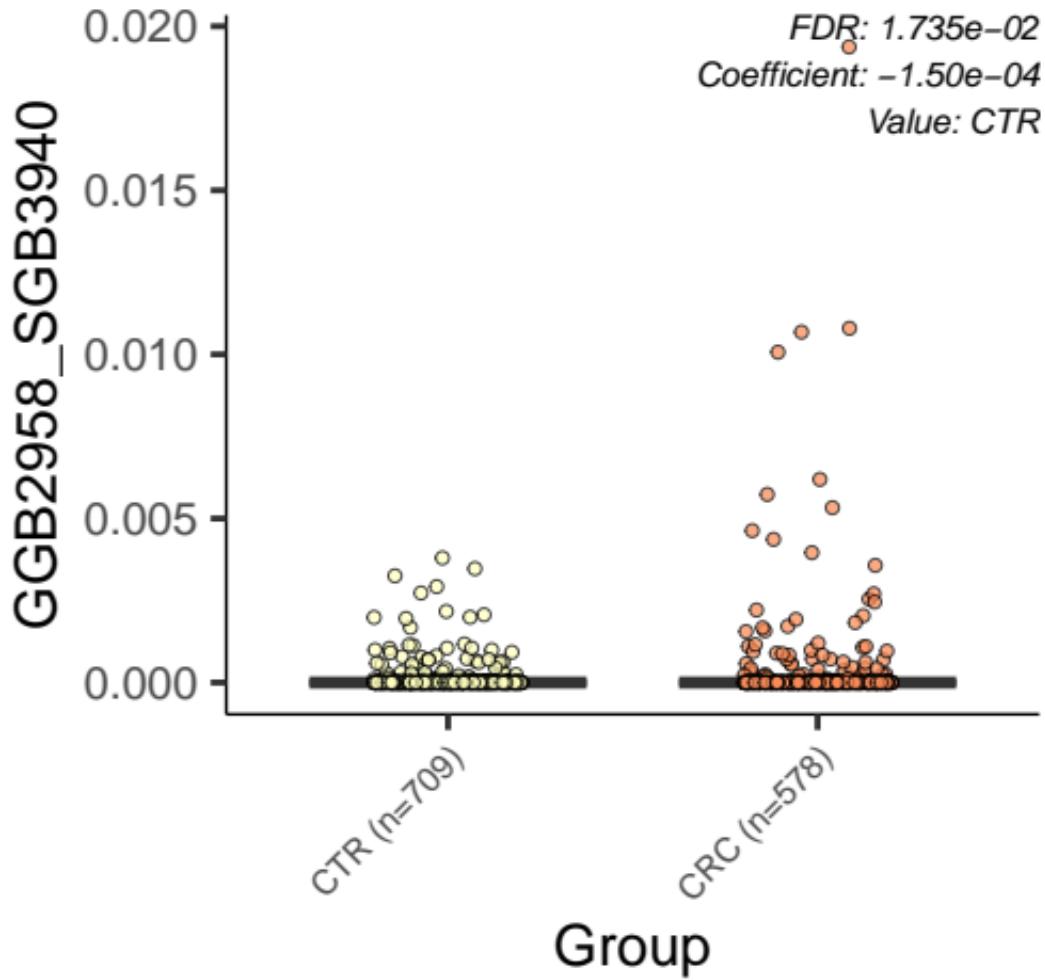


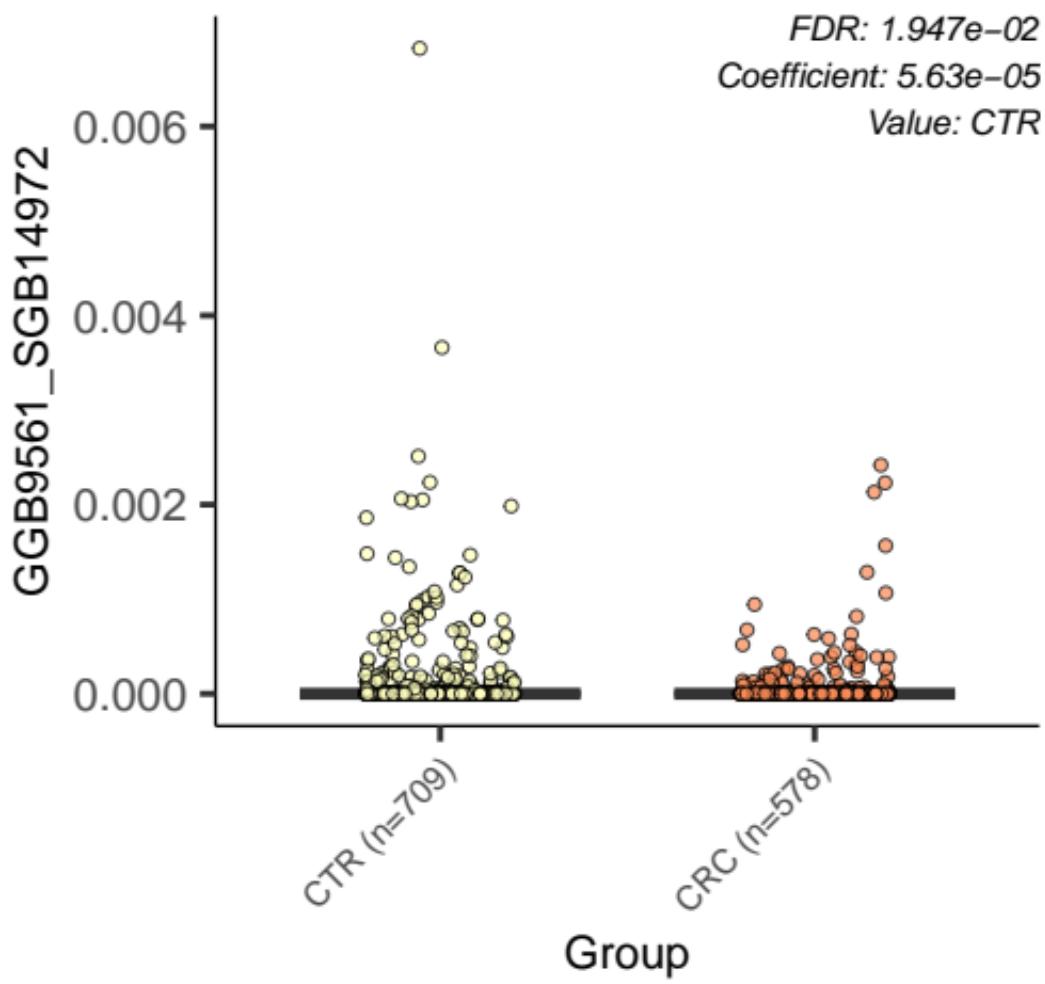


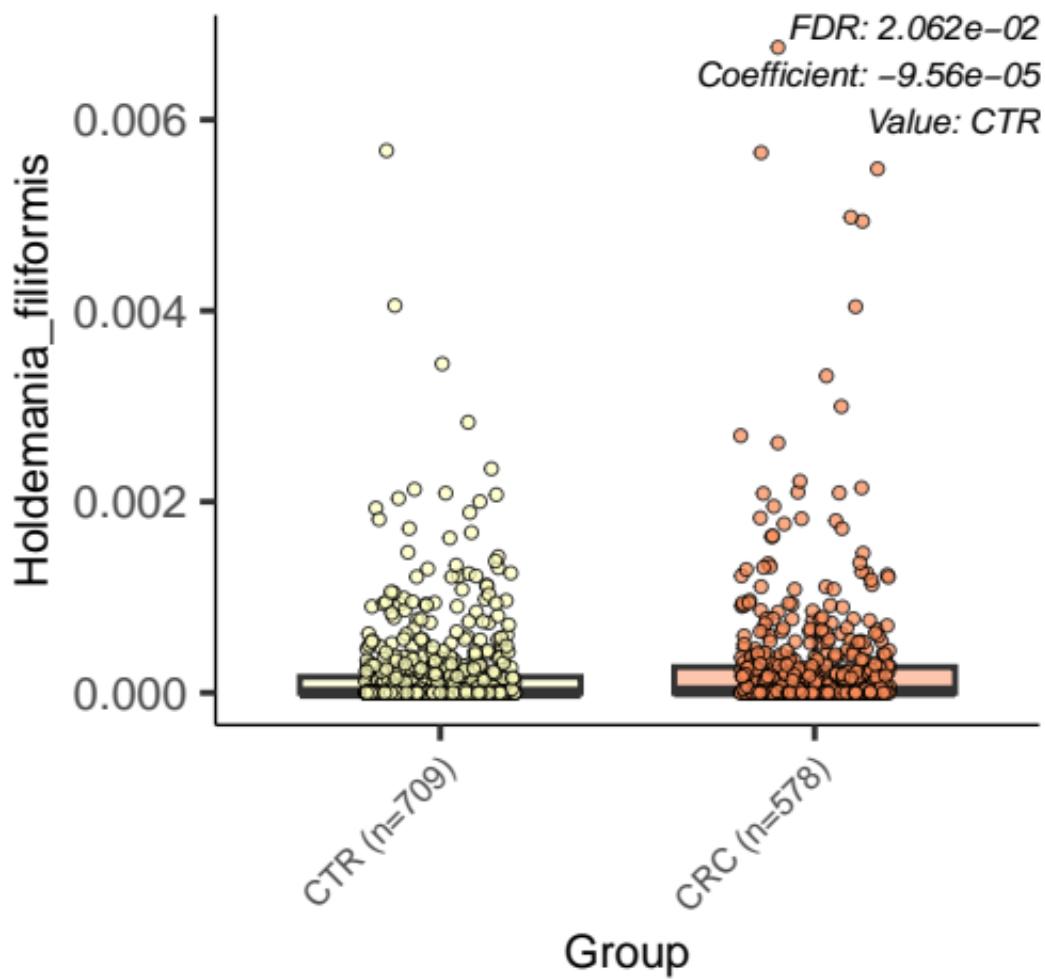


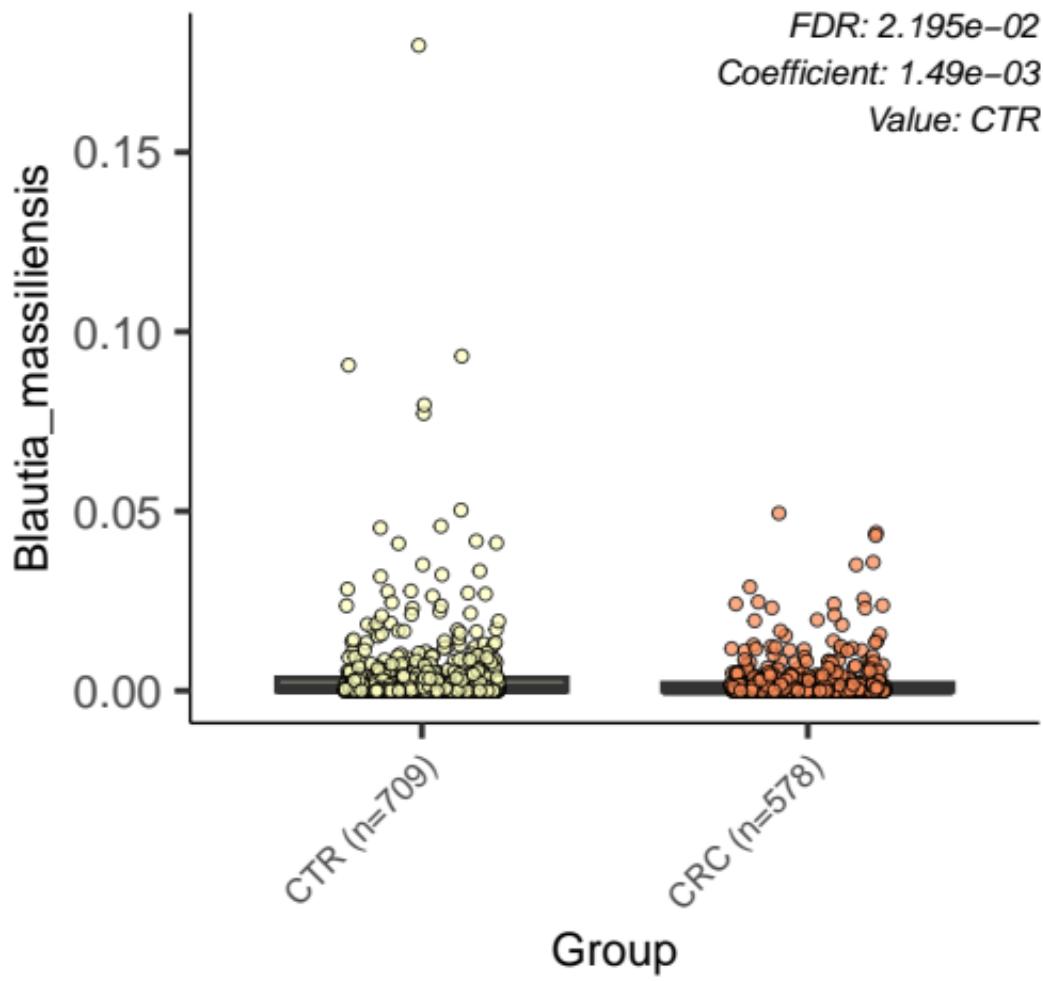


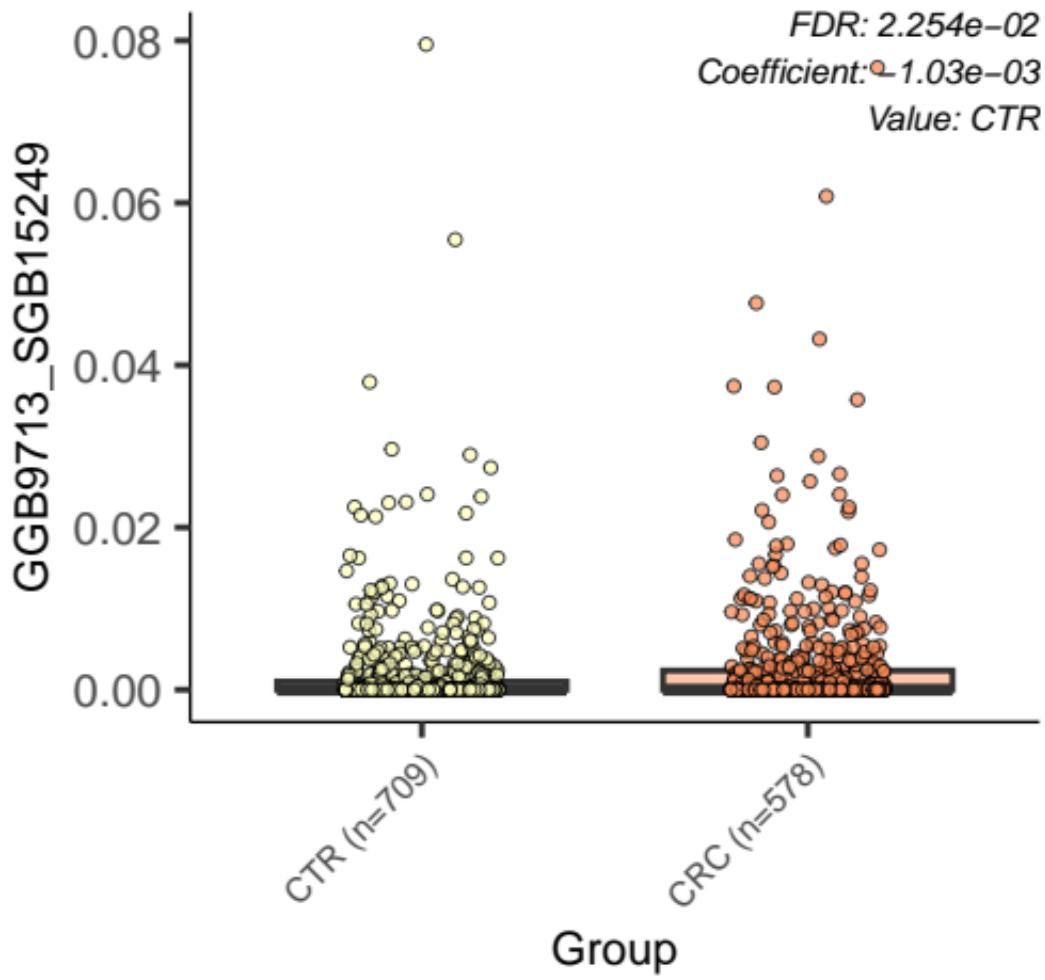












FDR: 2.406e-02
Coefficient: 2.55e-04
Value: CTR

Clostridium_sp_AF34_13

0.02

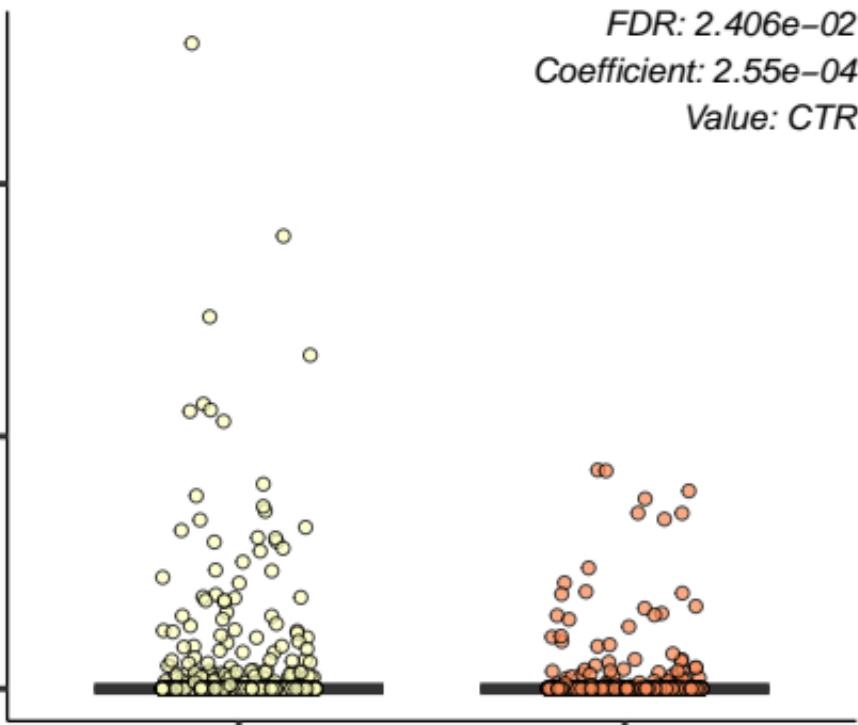
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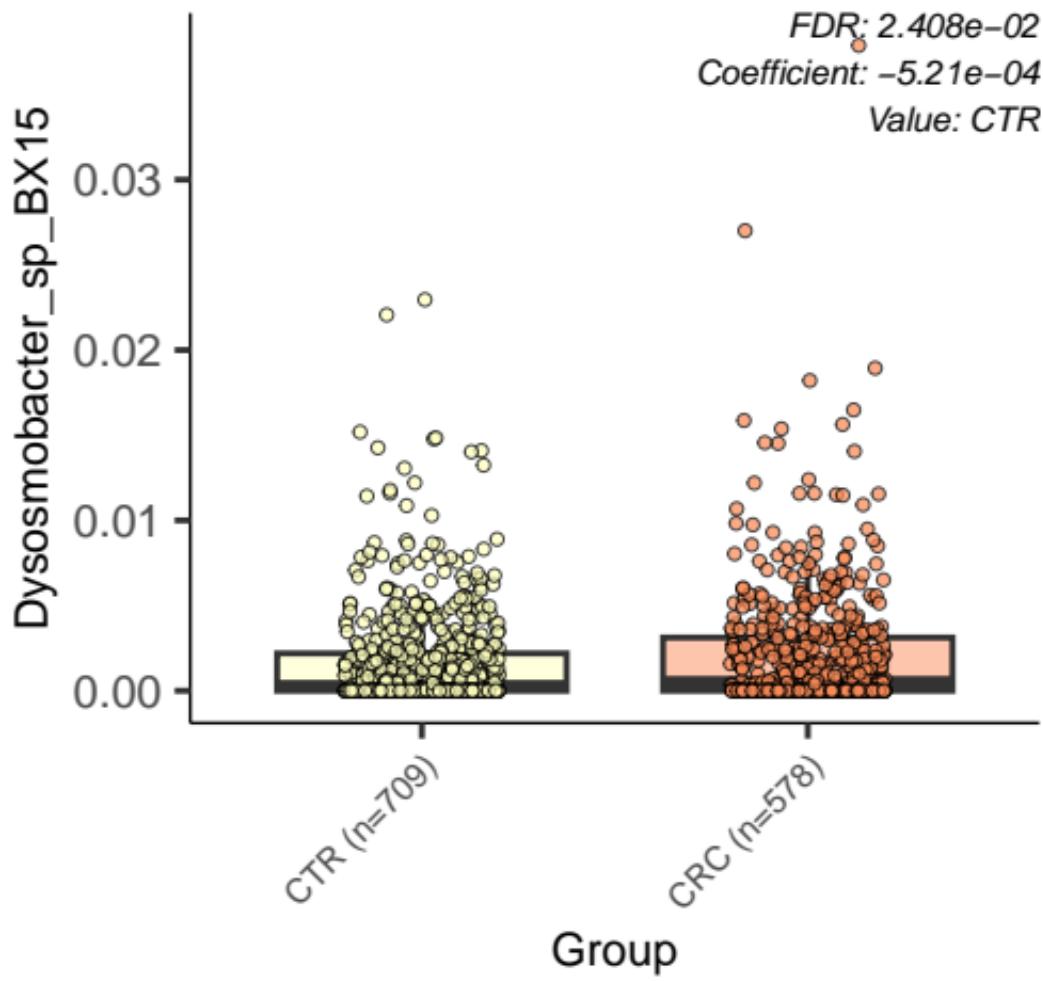
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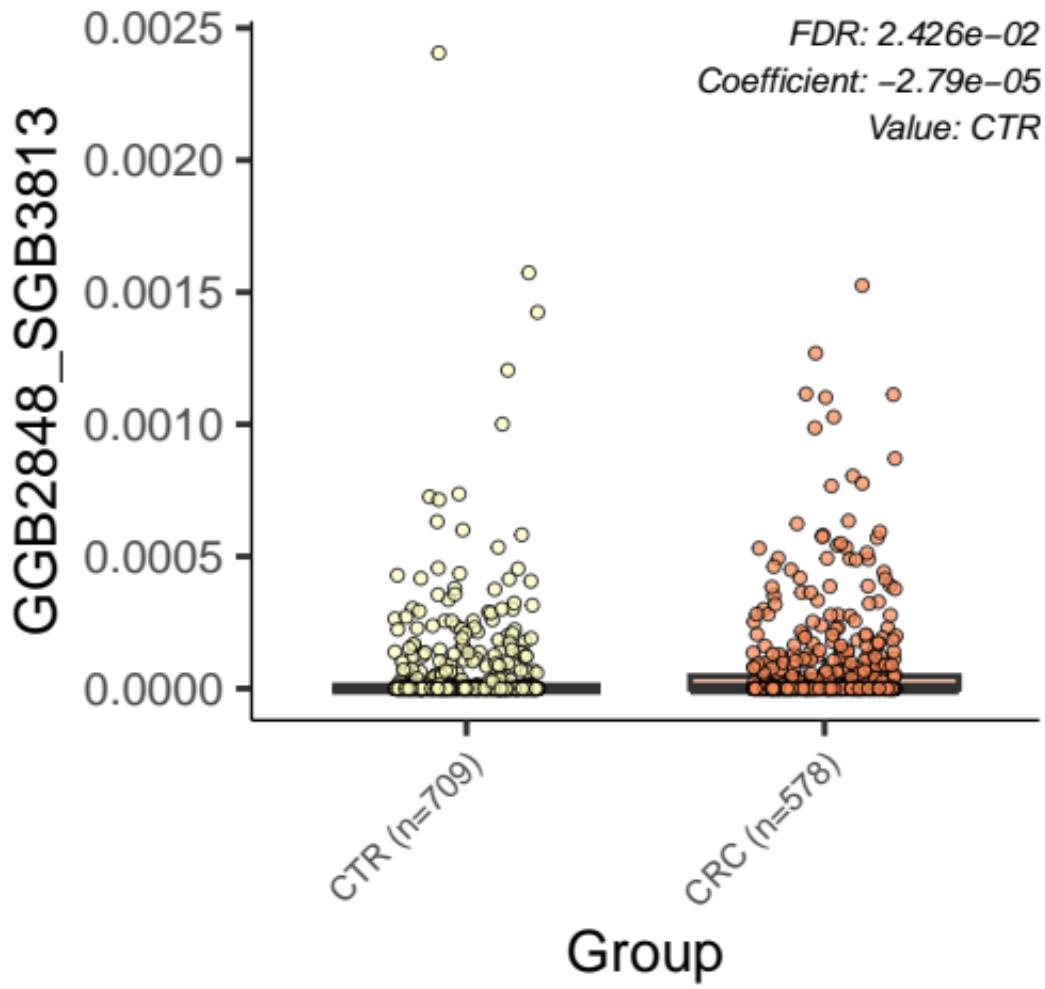
CTR (n=709)

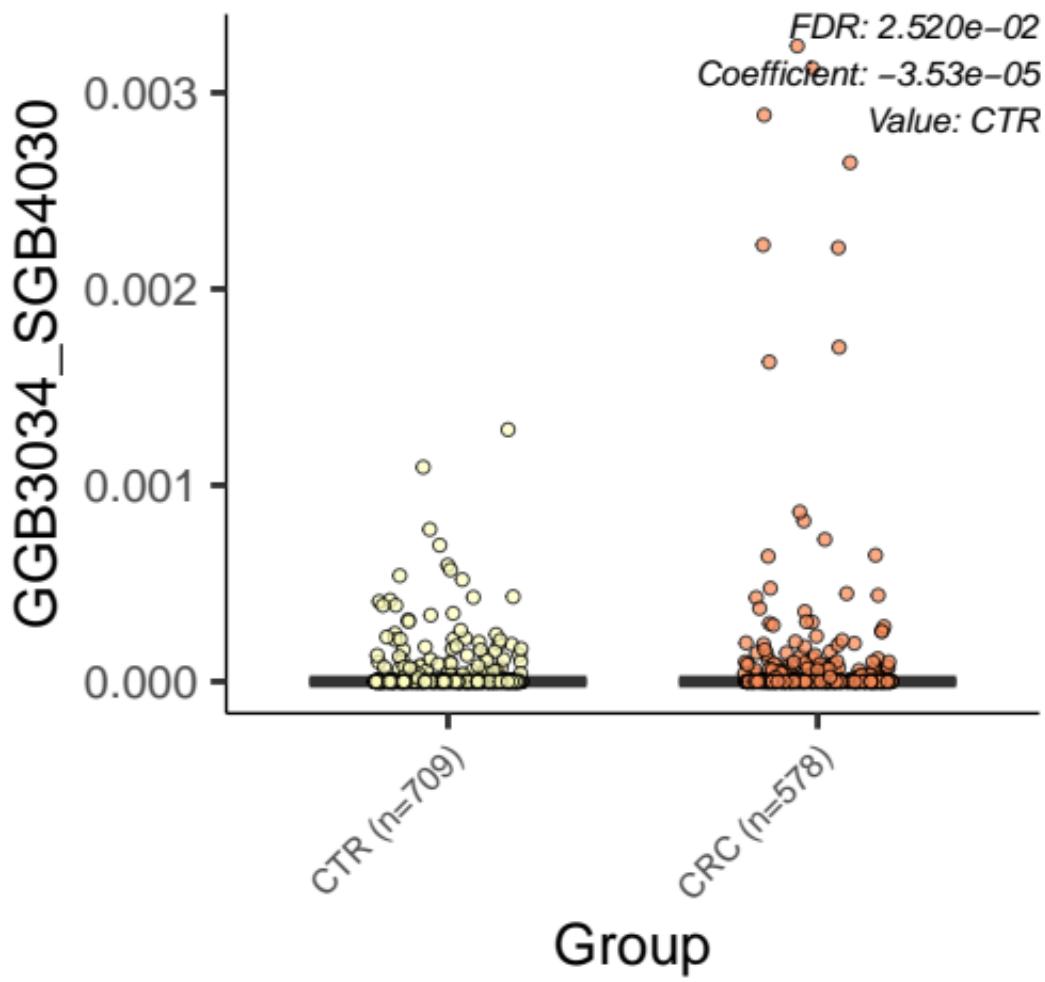
CRC (n=578)

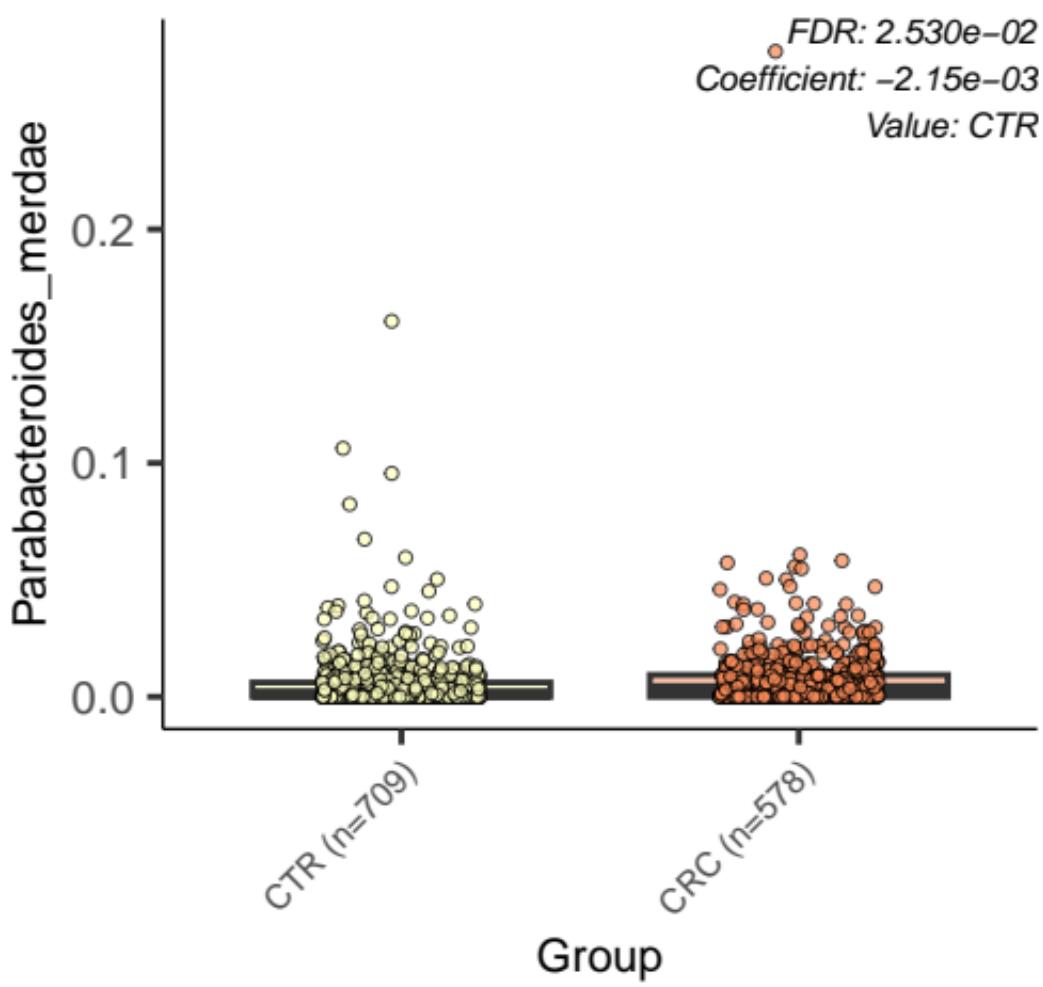
Group

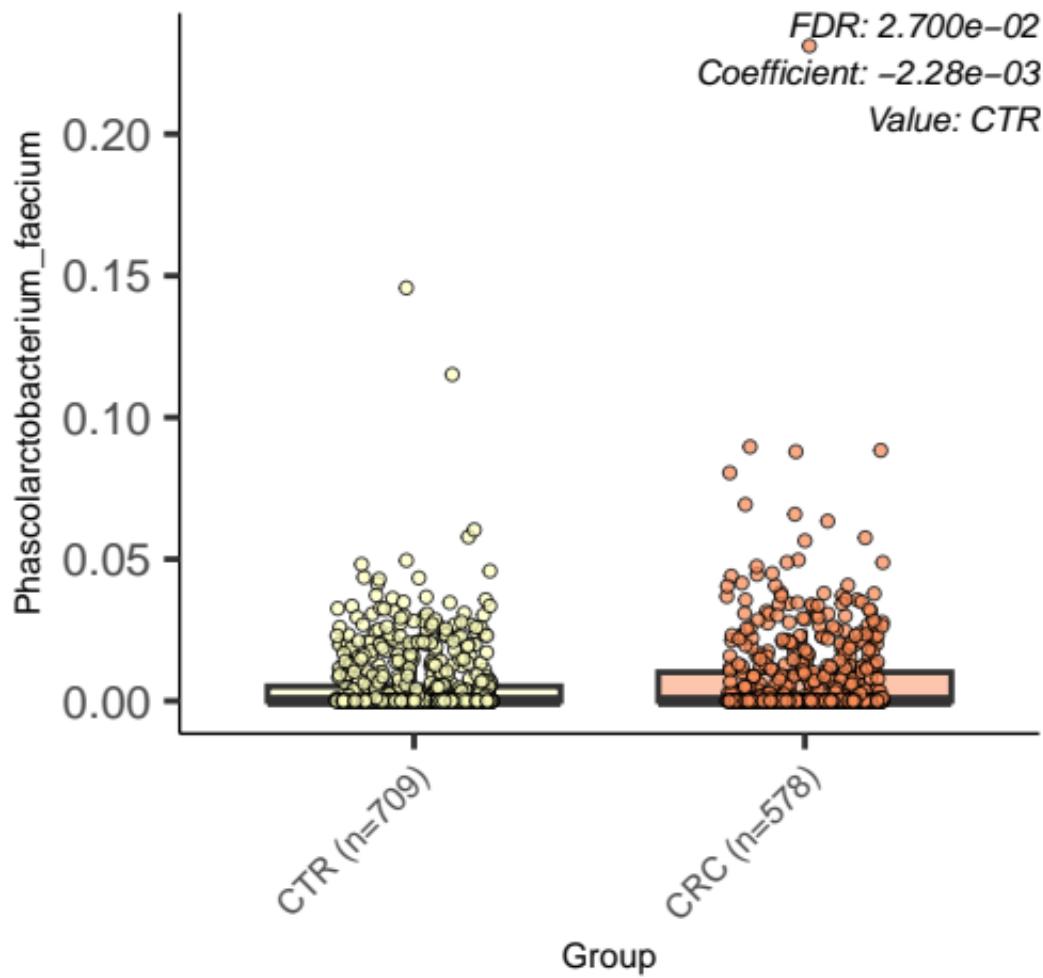


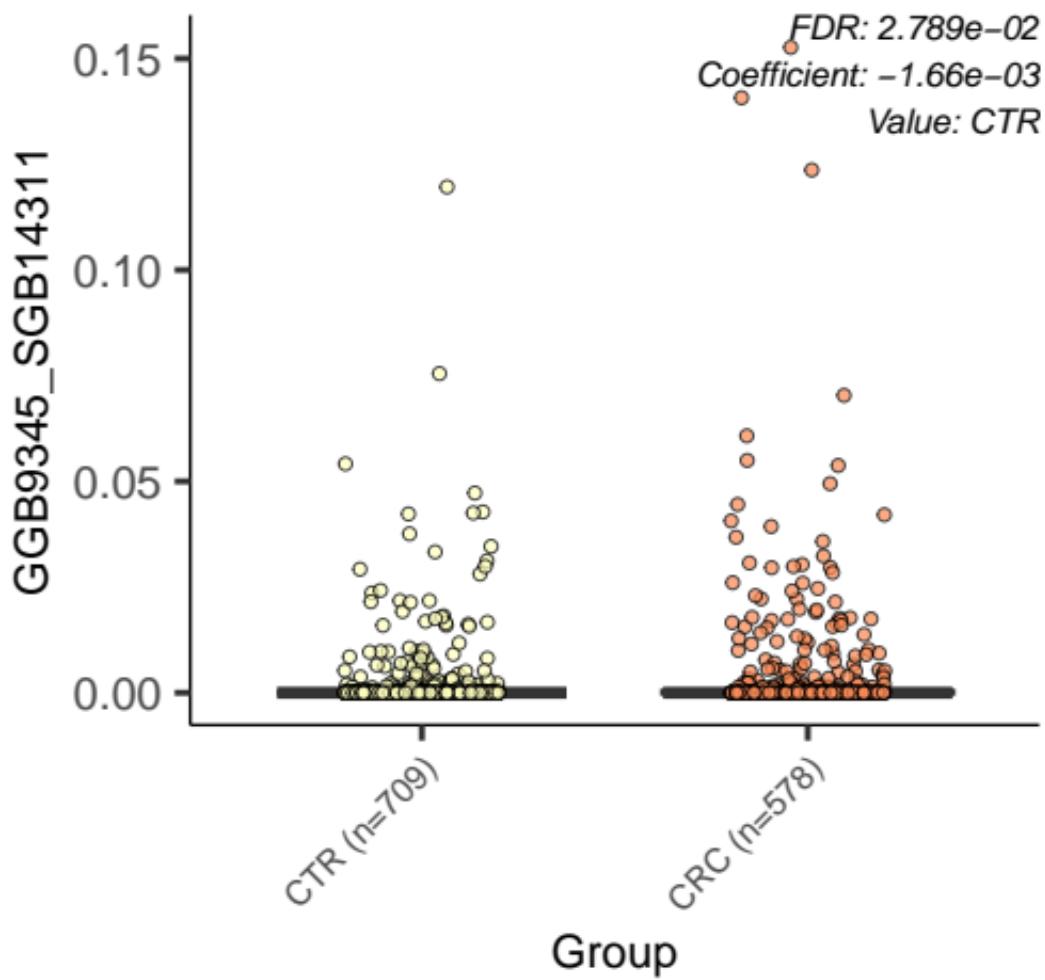


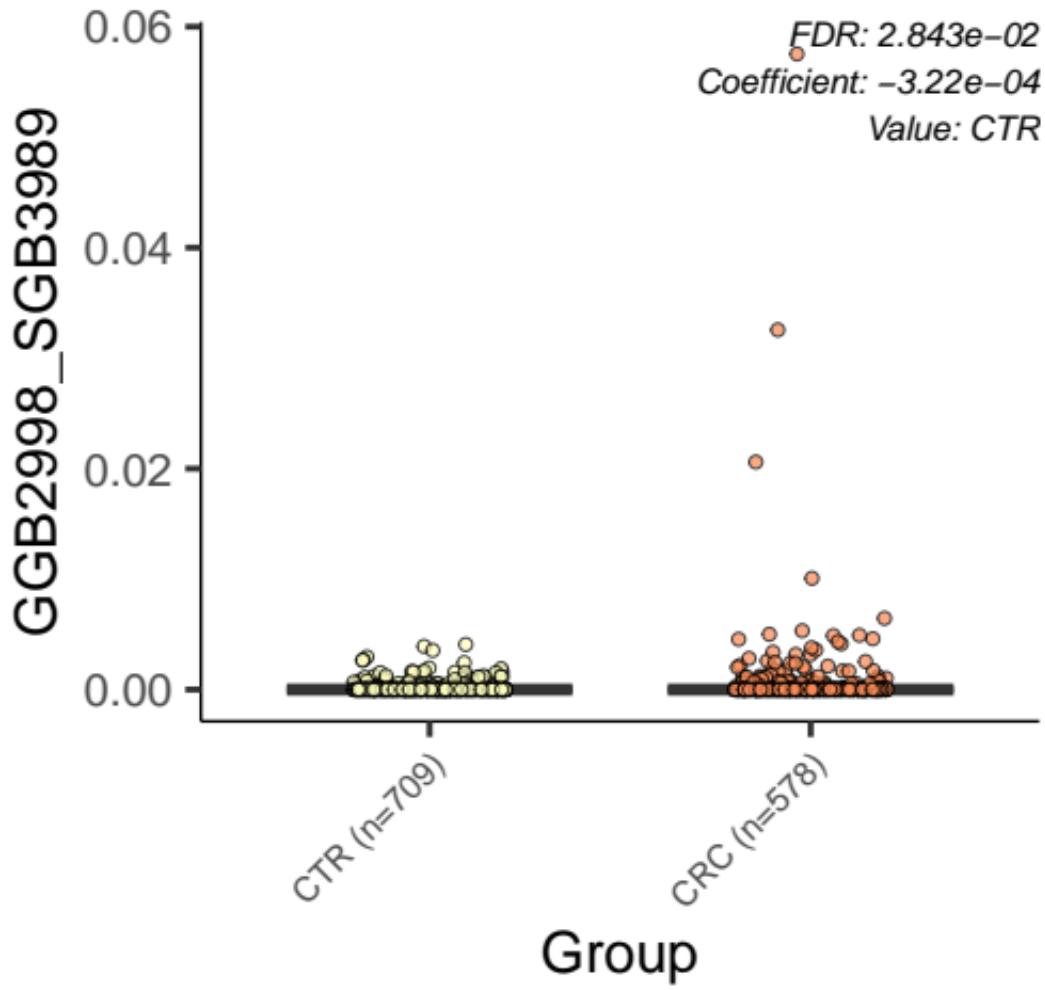


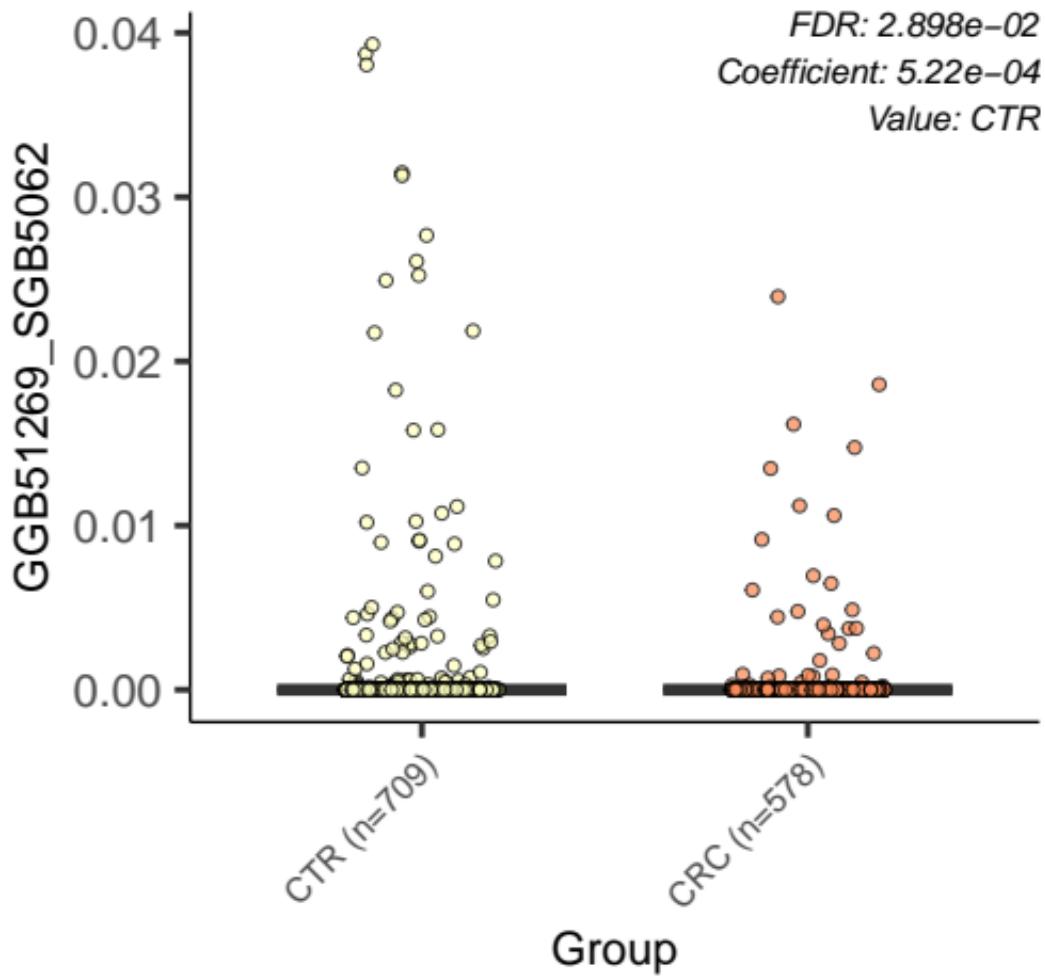


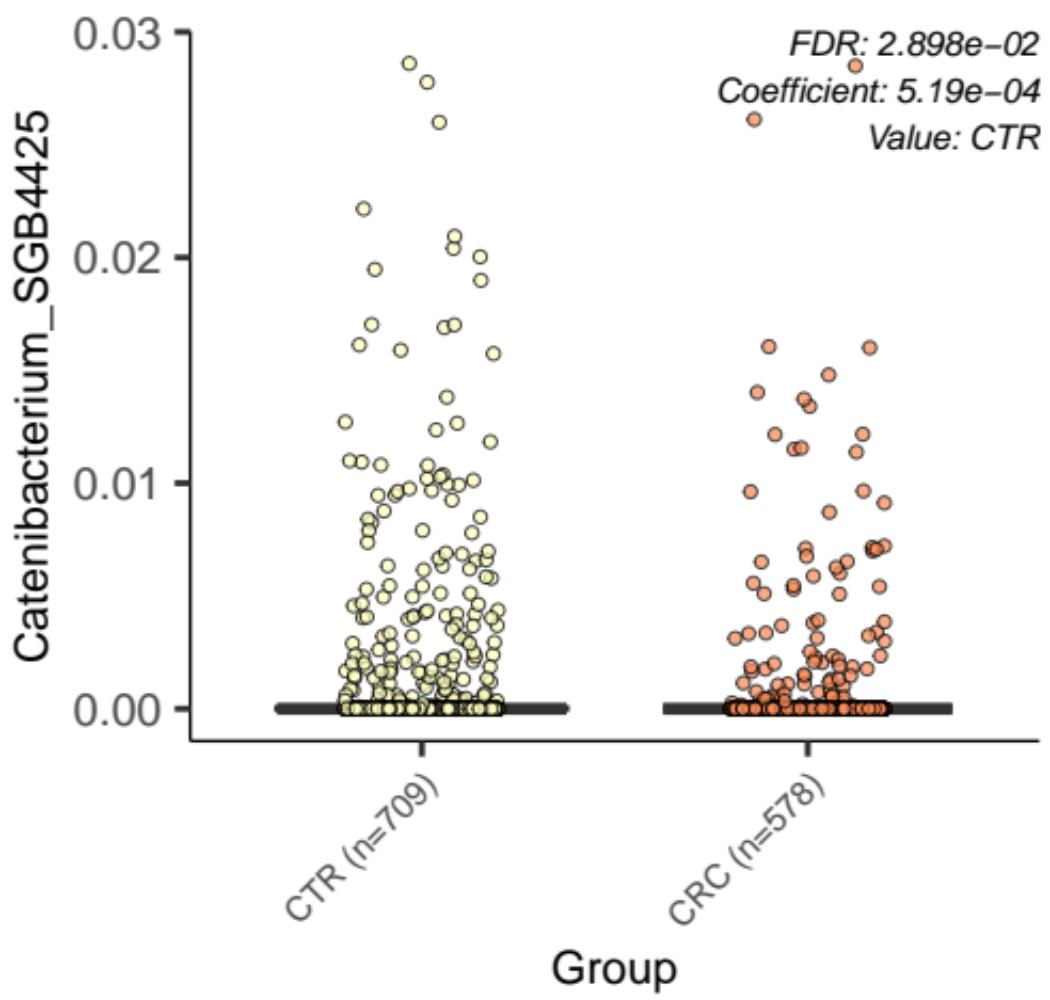


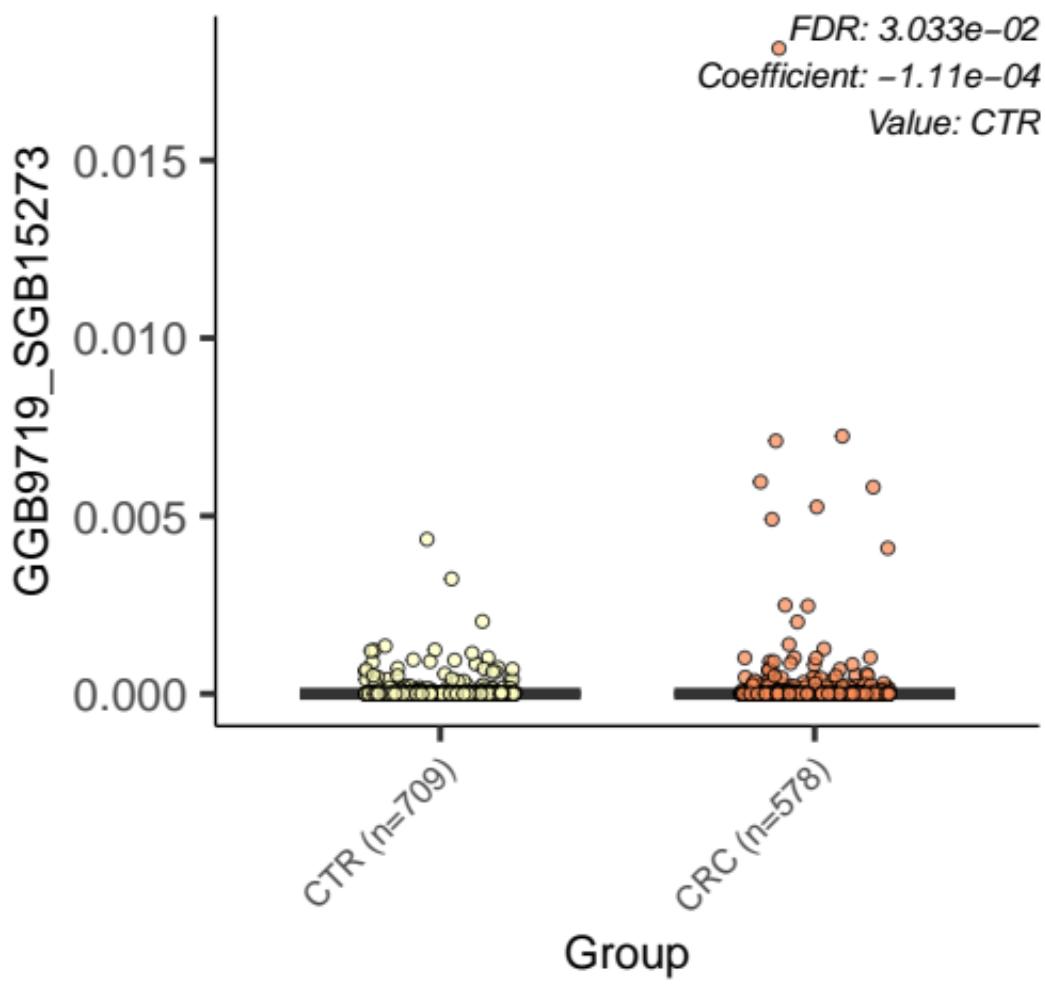


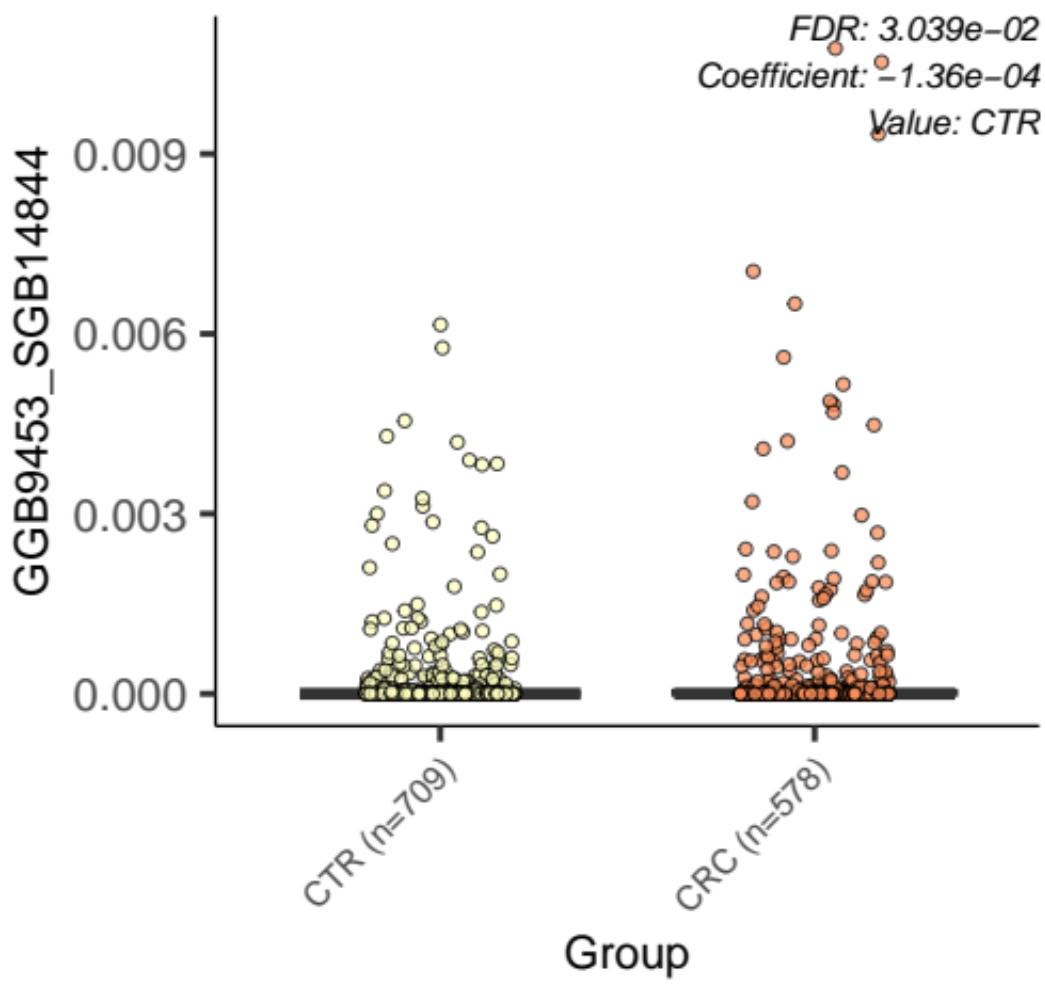


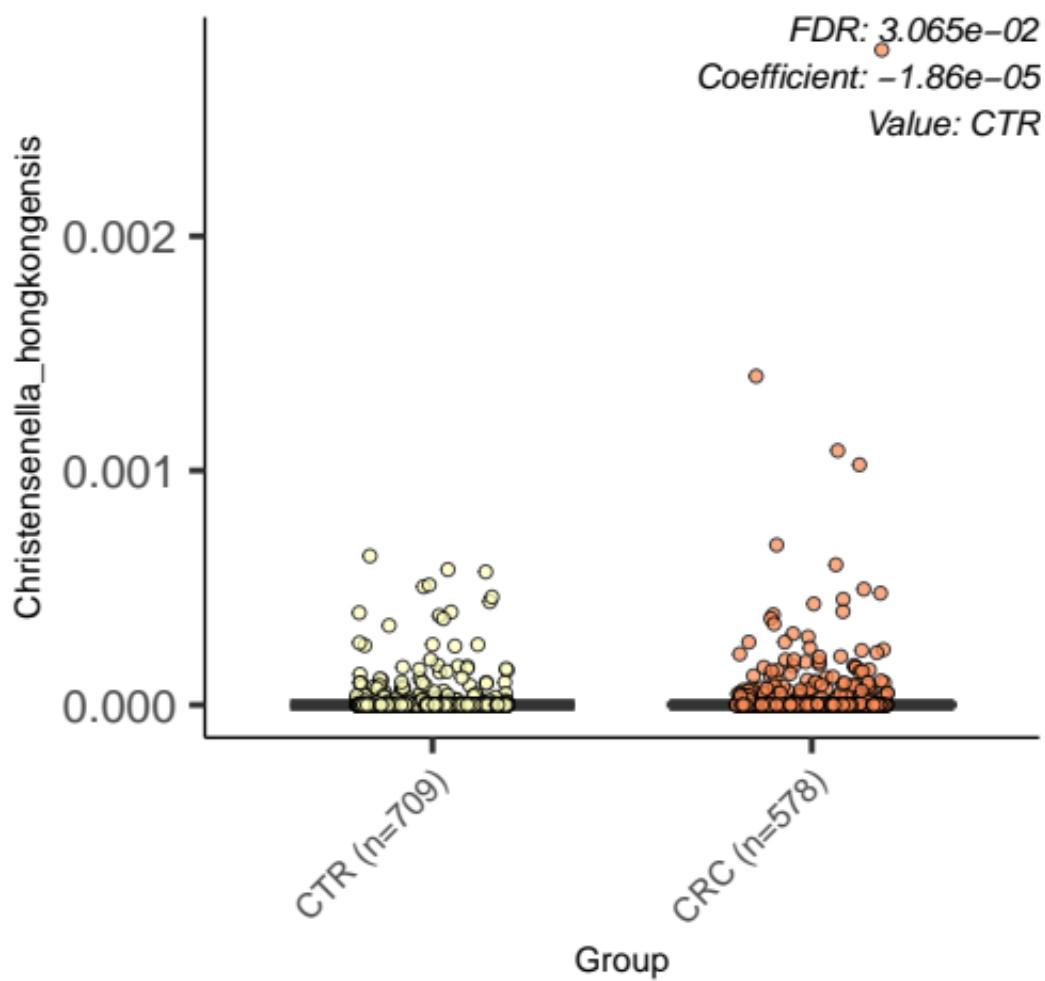


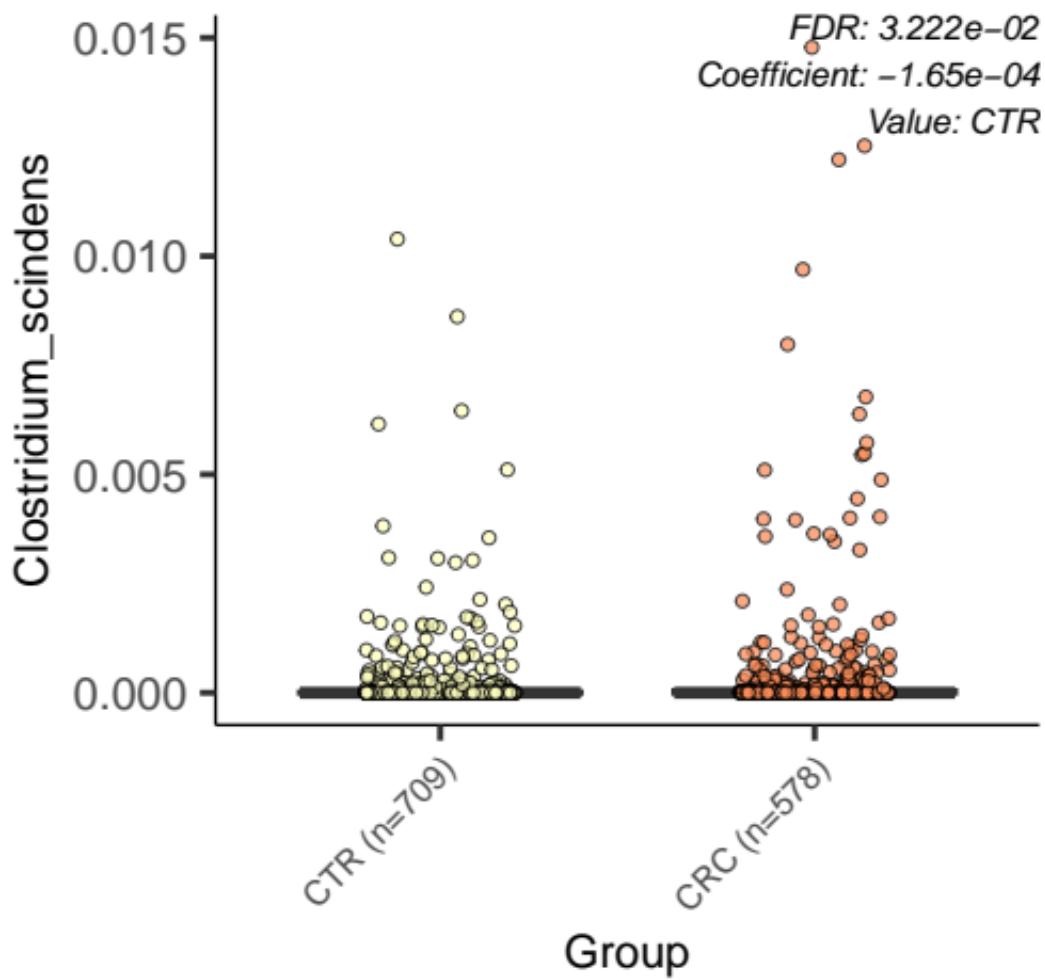


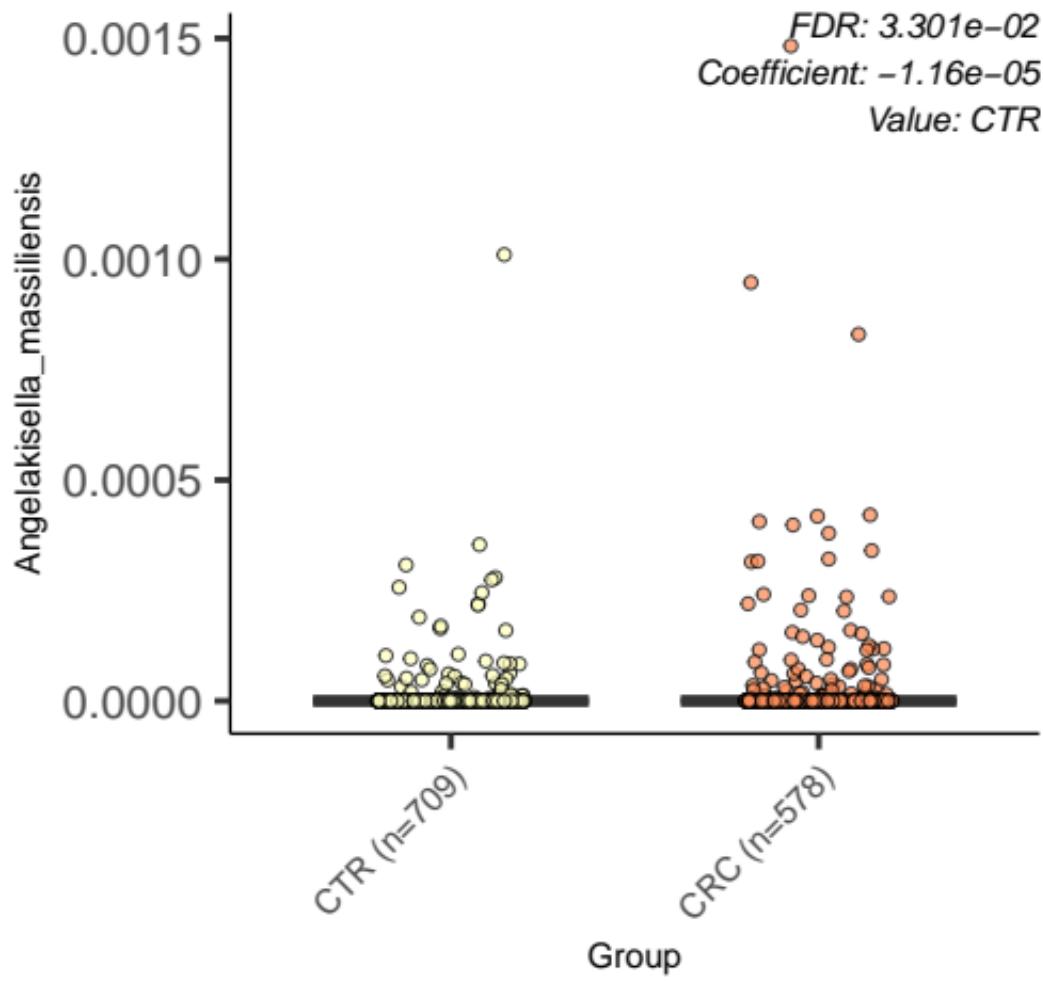


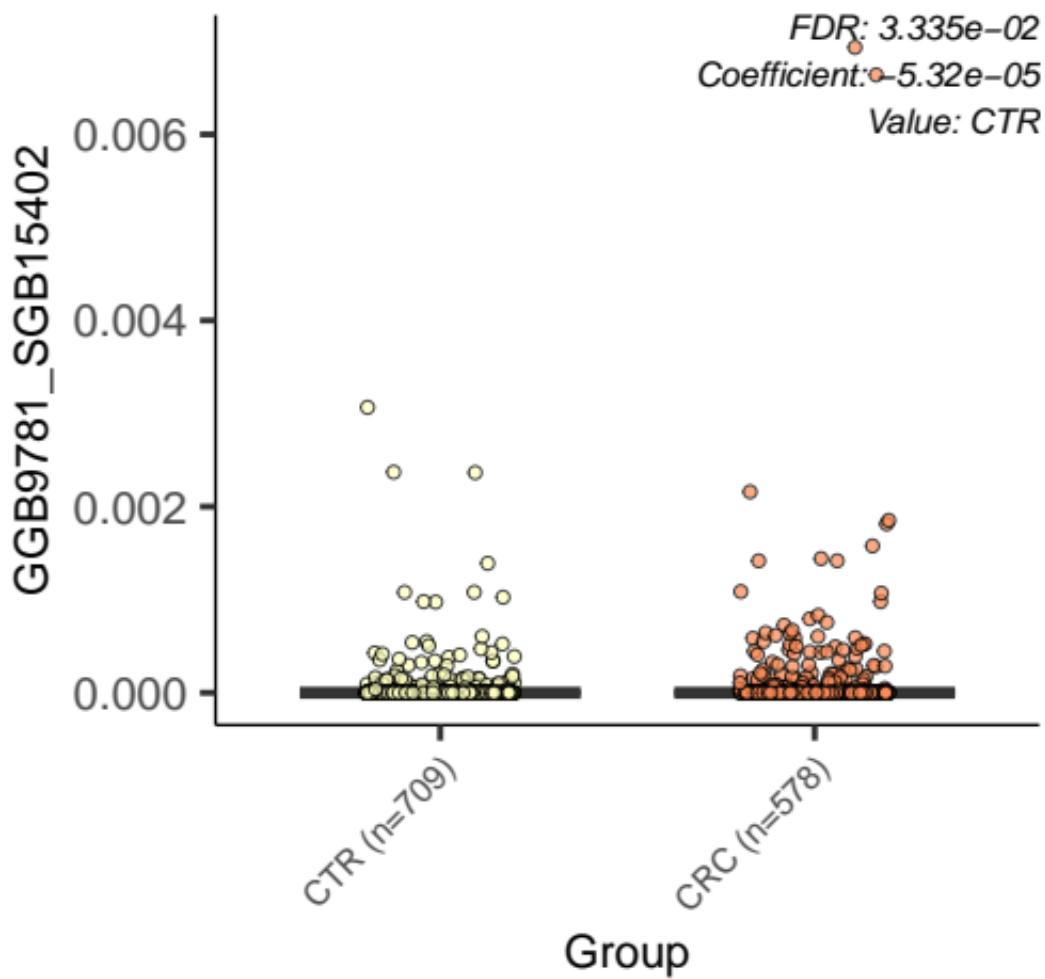


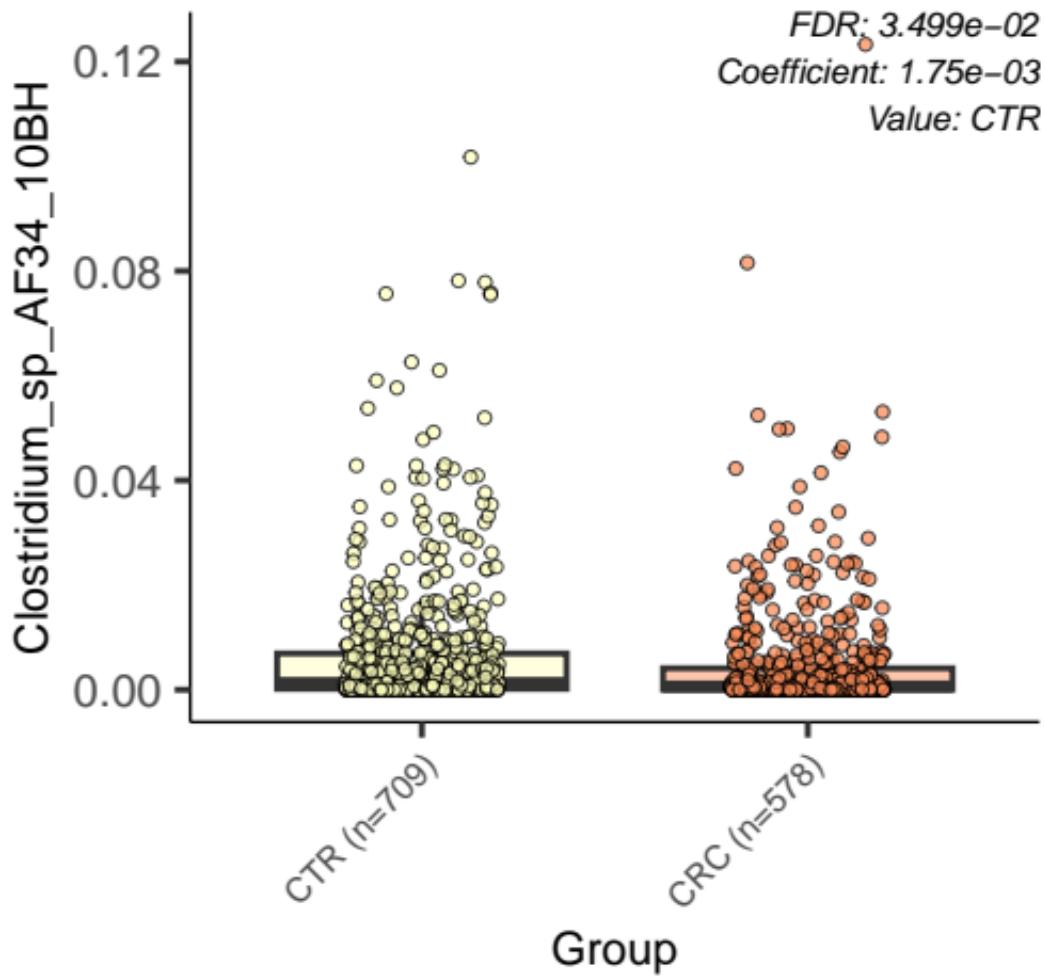


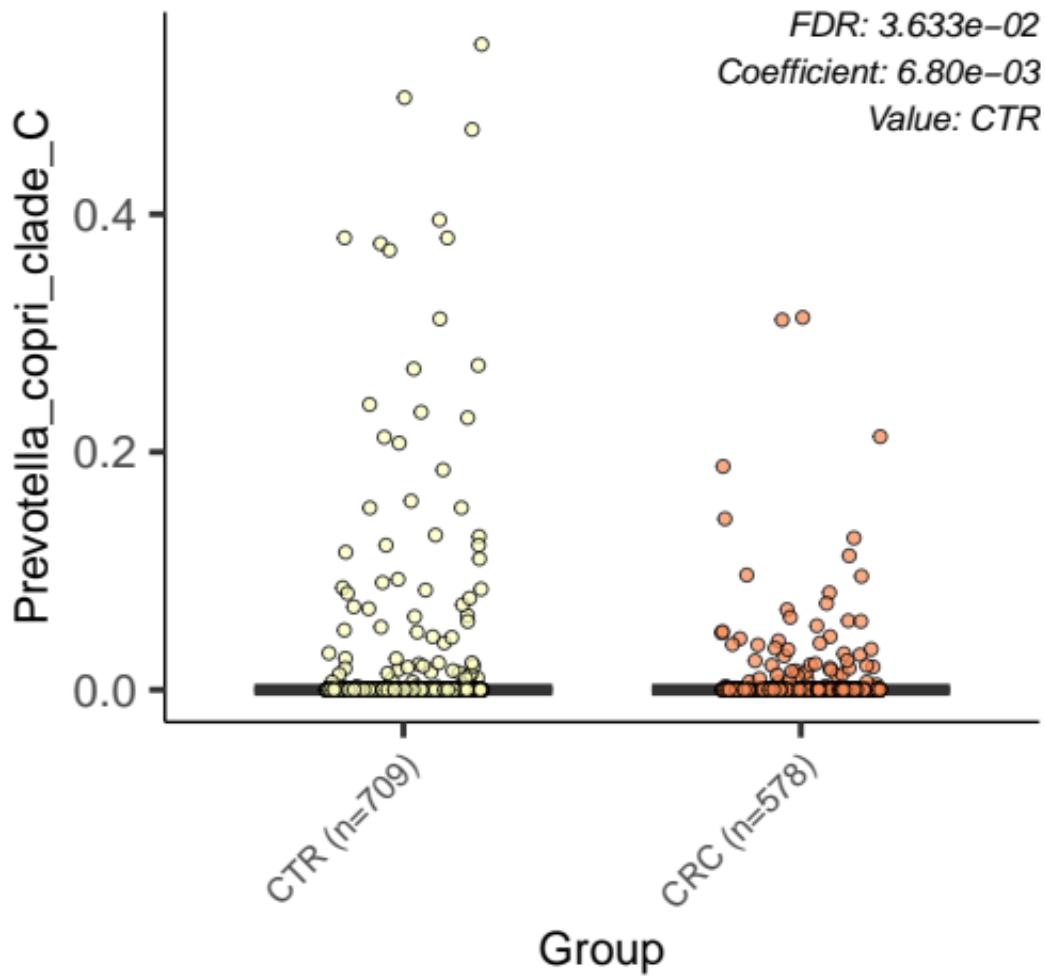


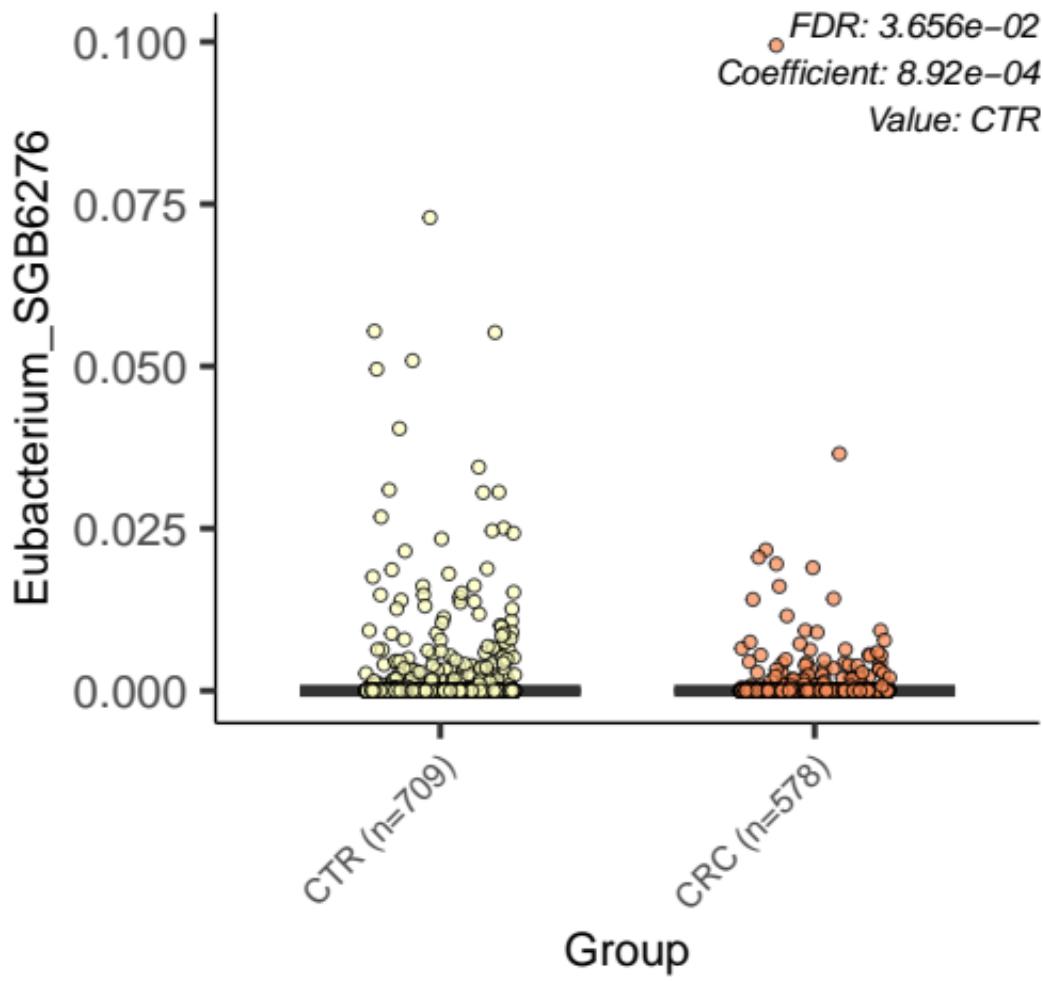


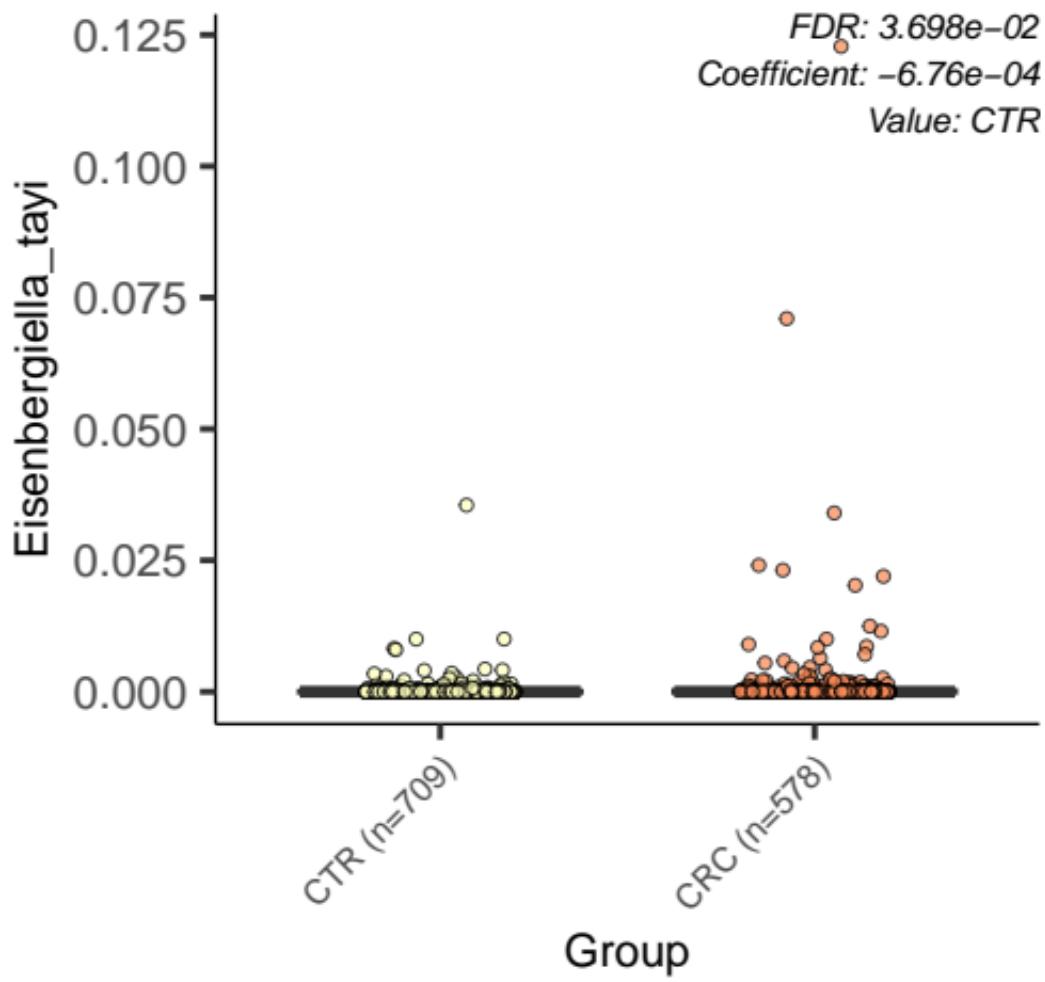




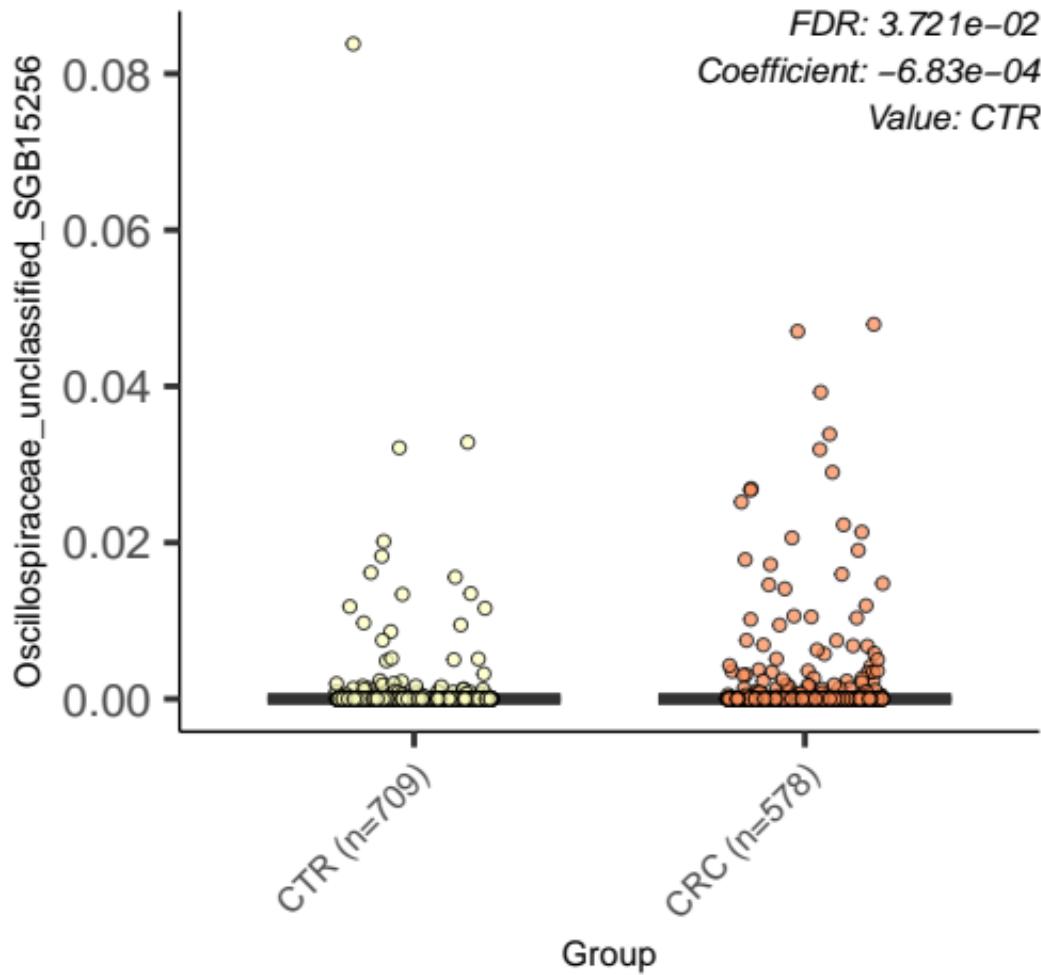


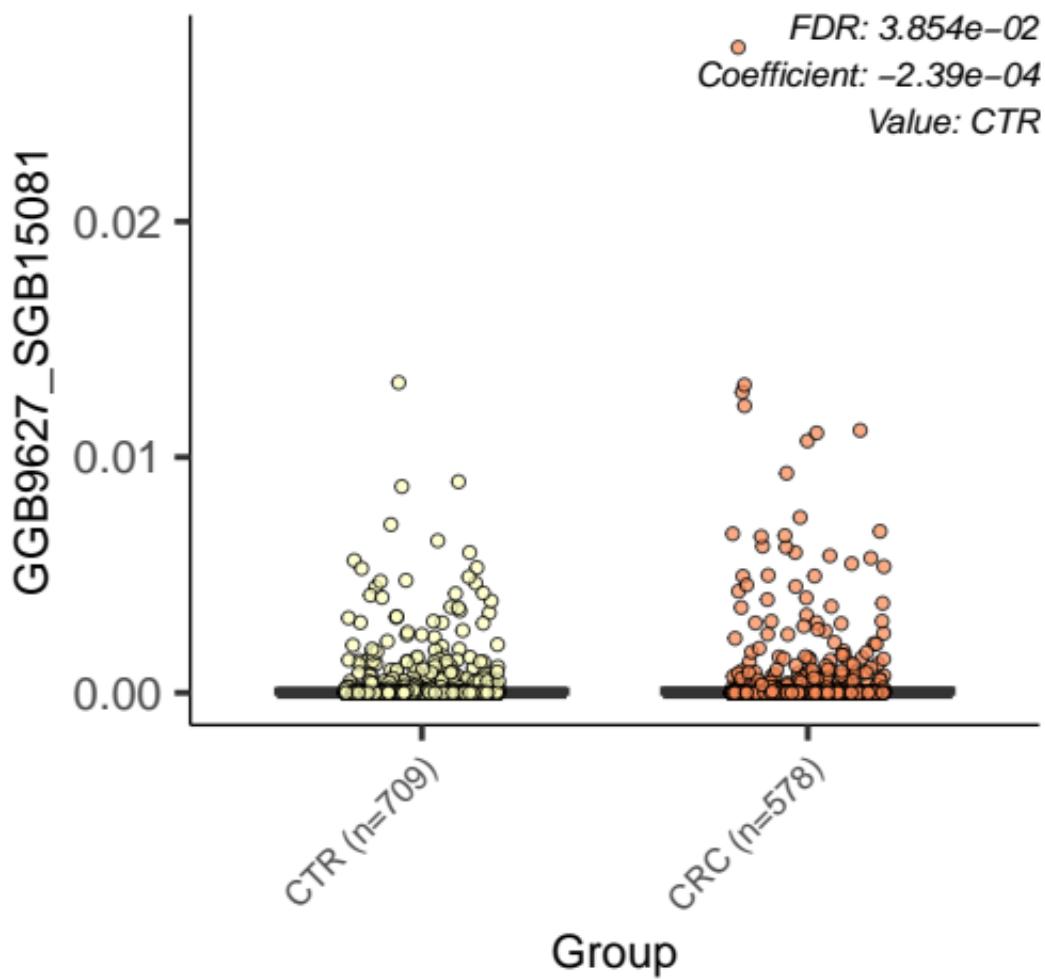


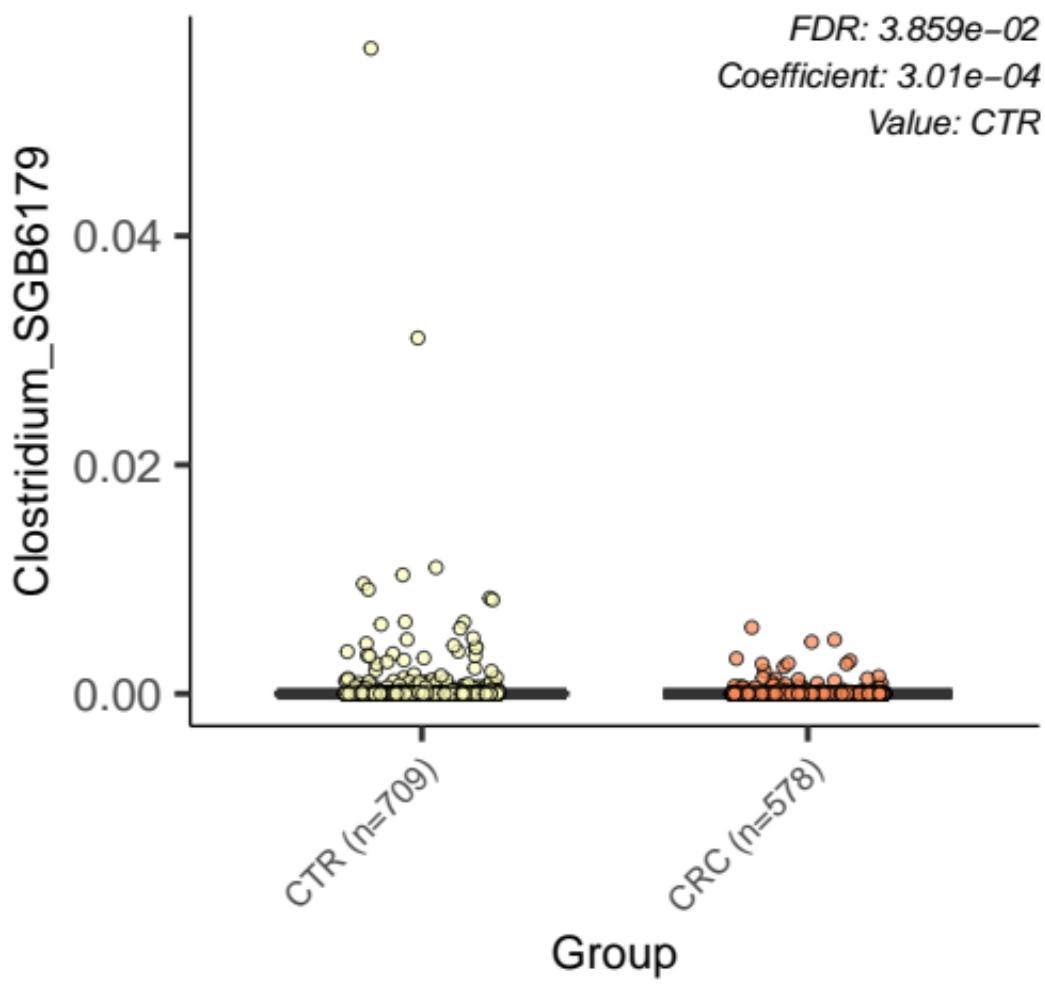


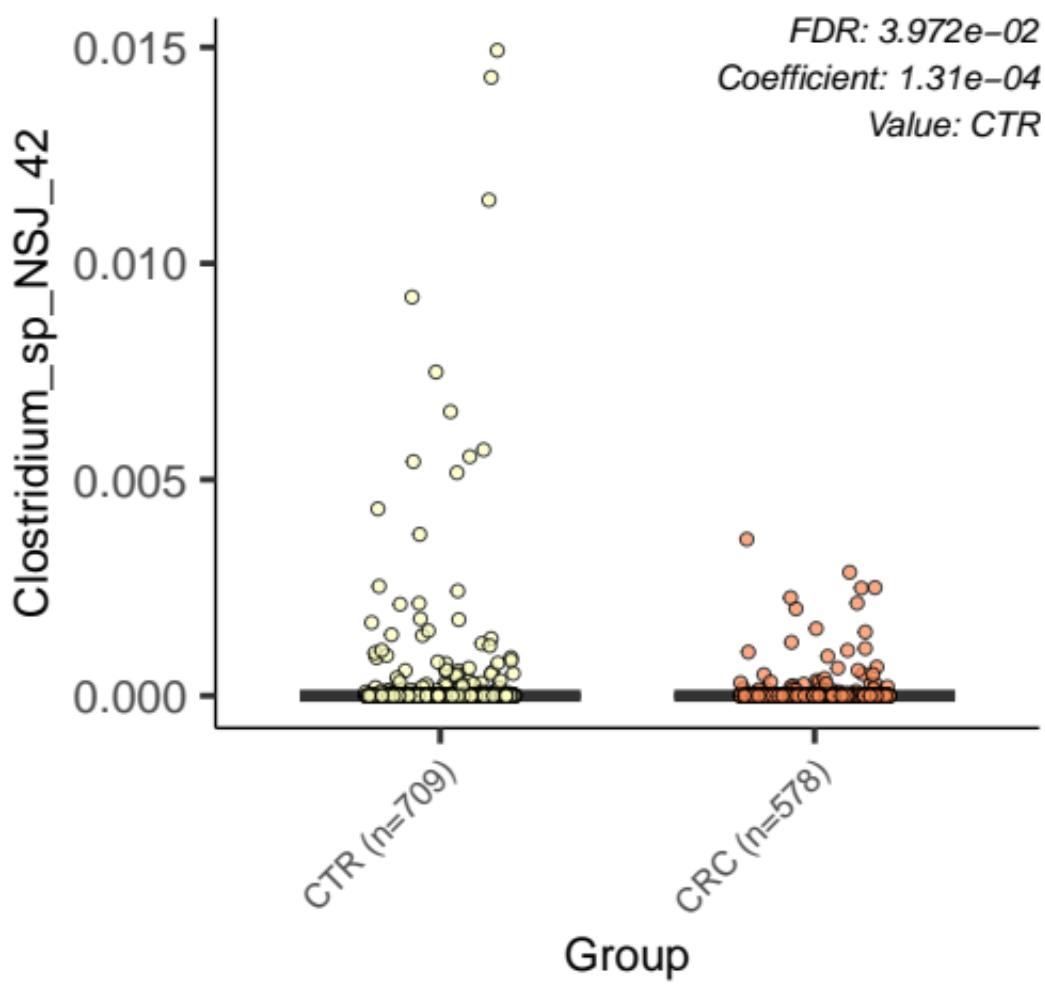


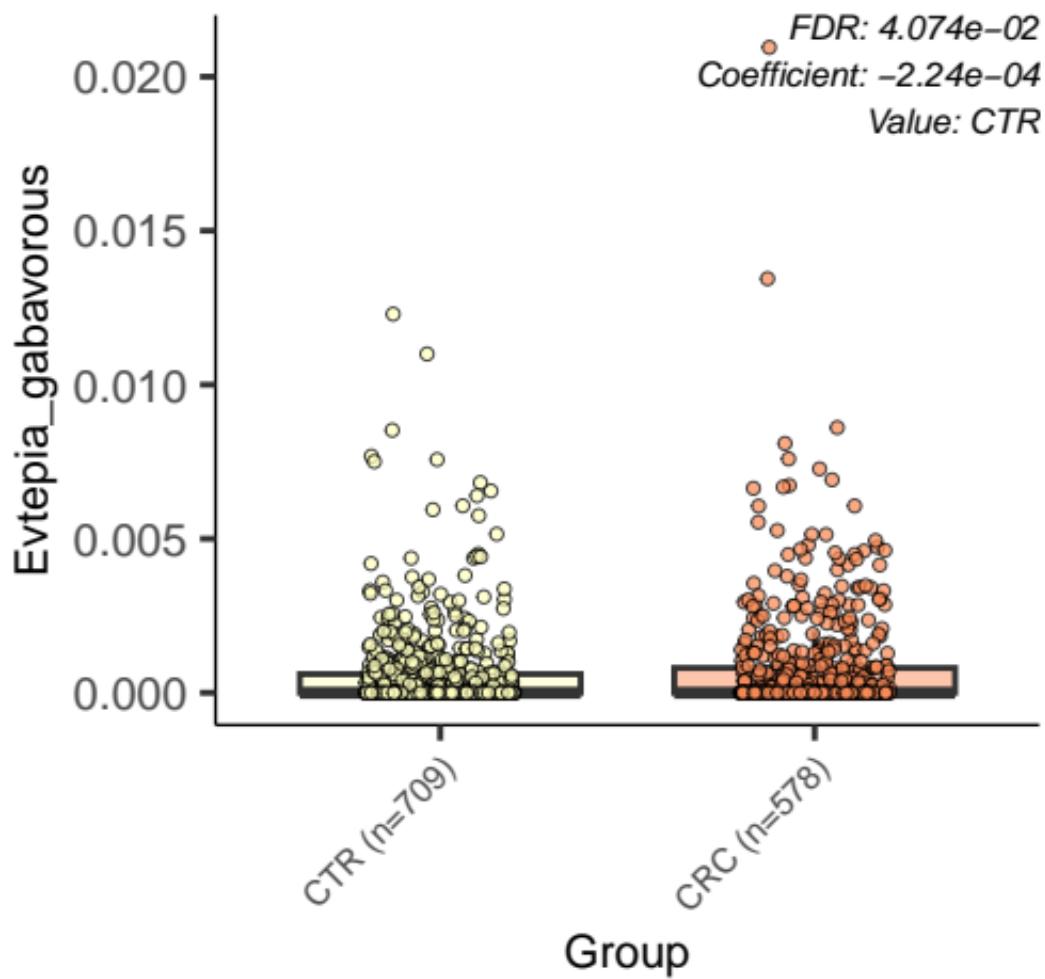
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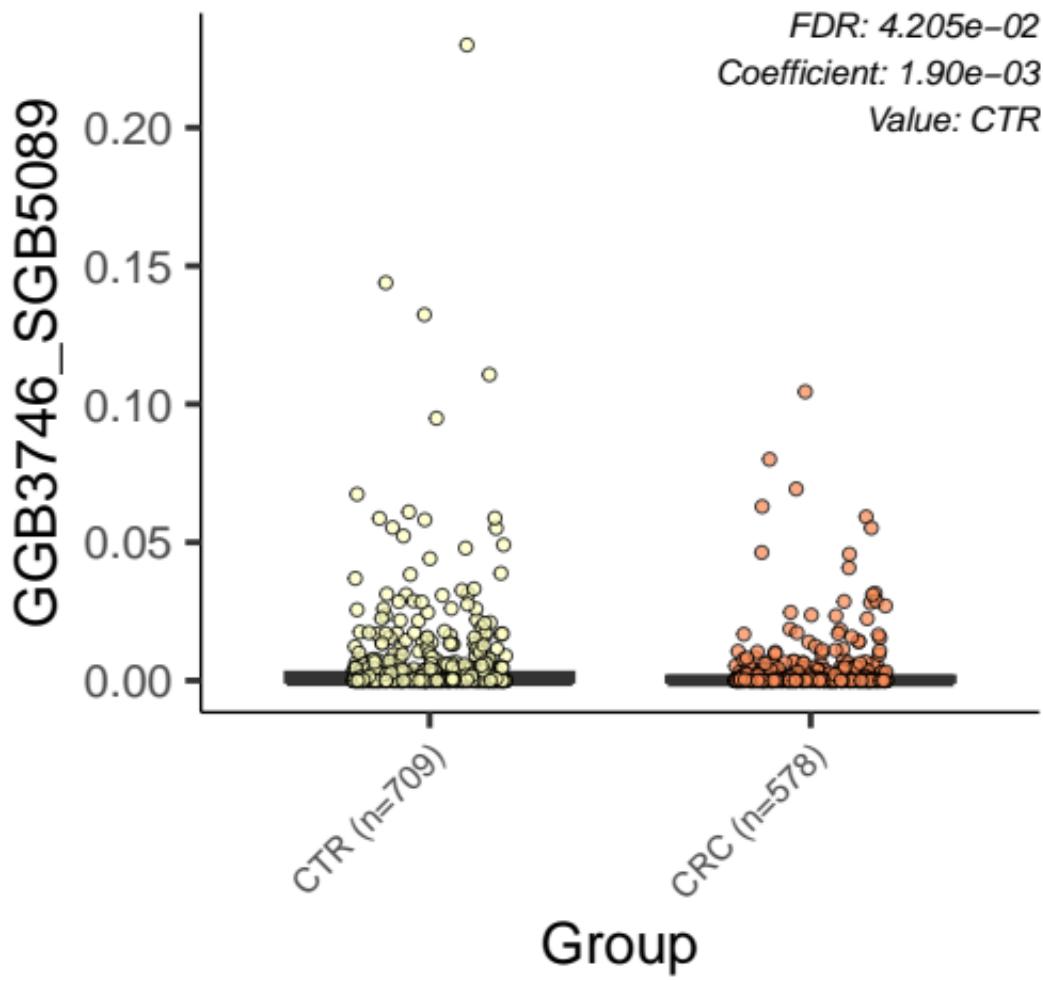


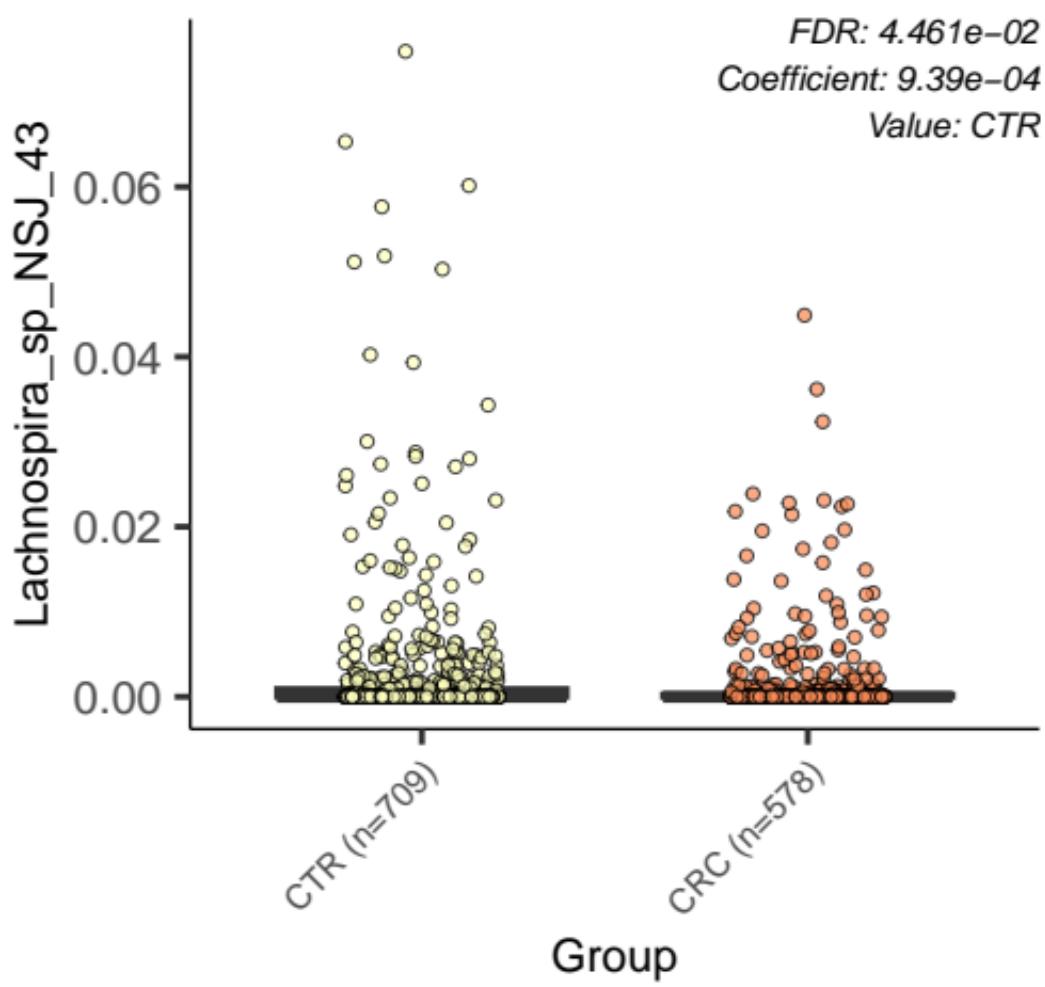


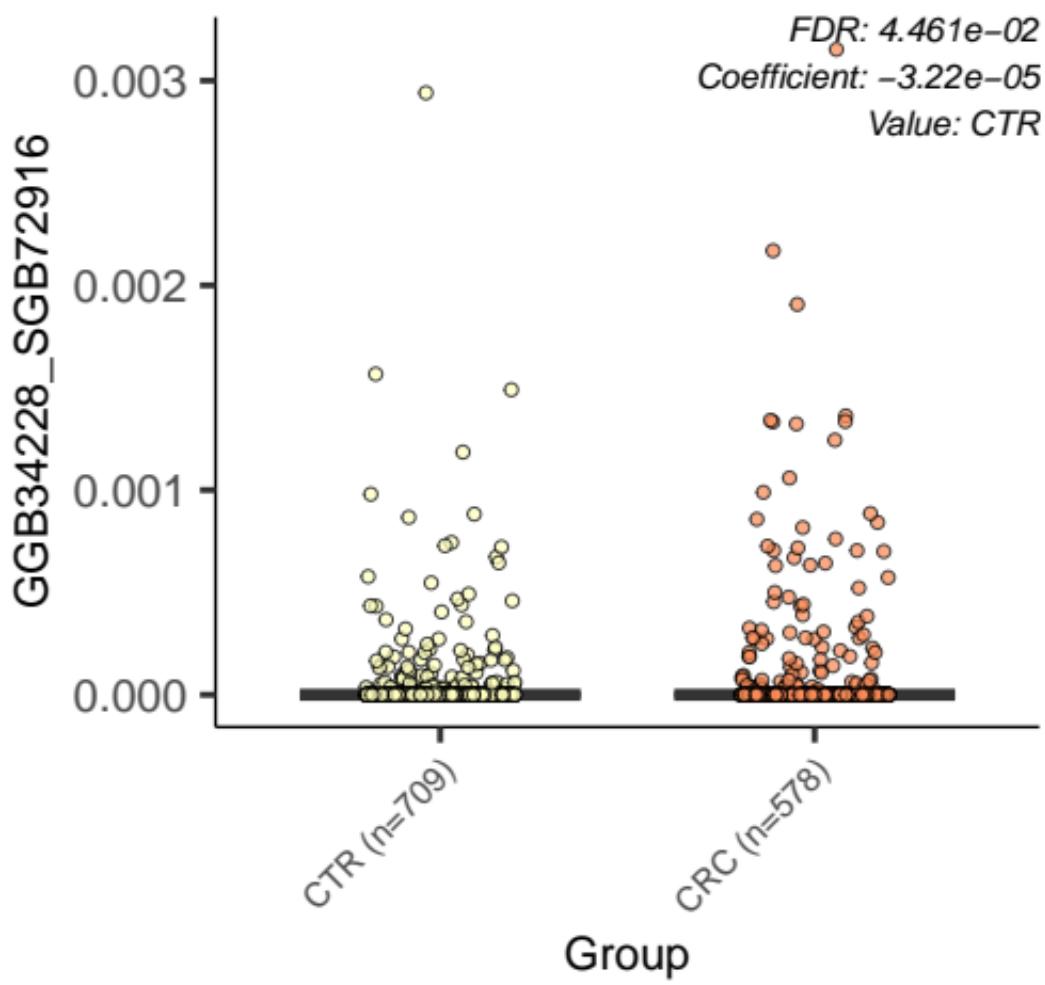


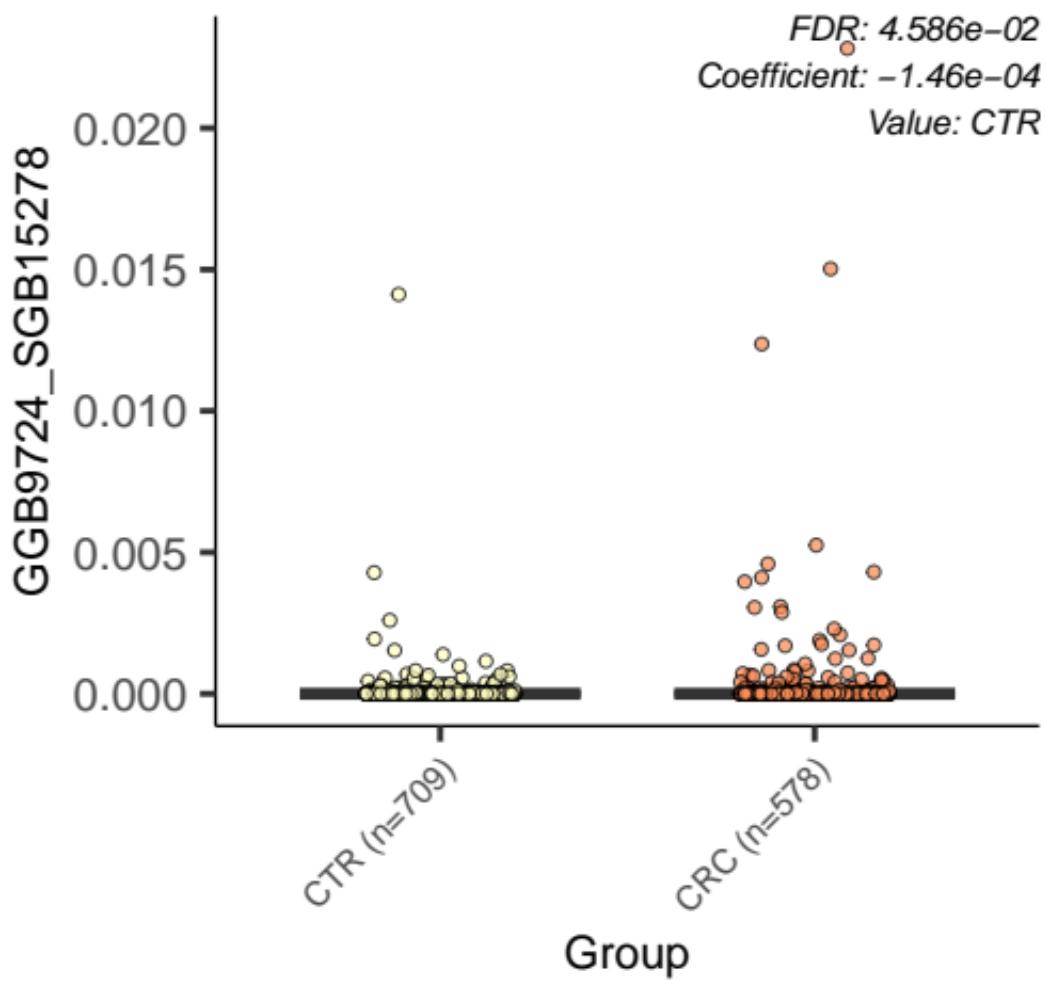


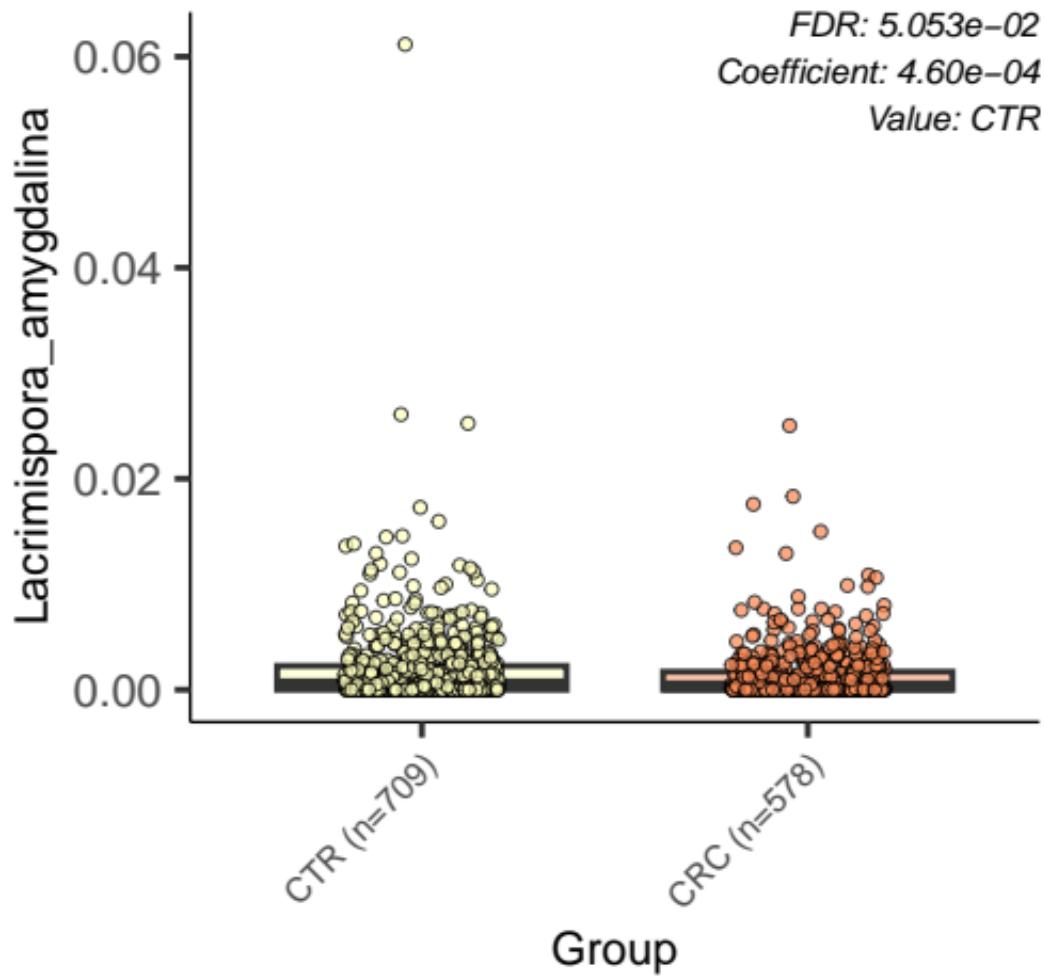


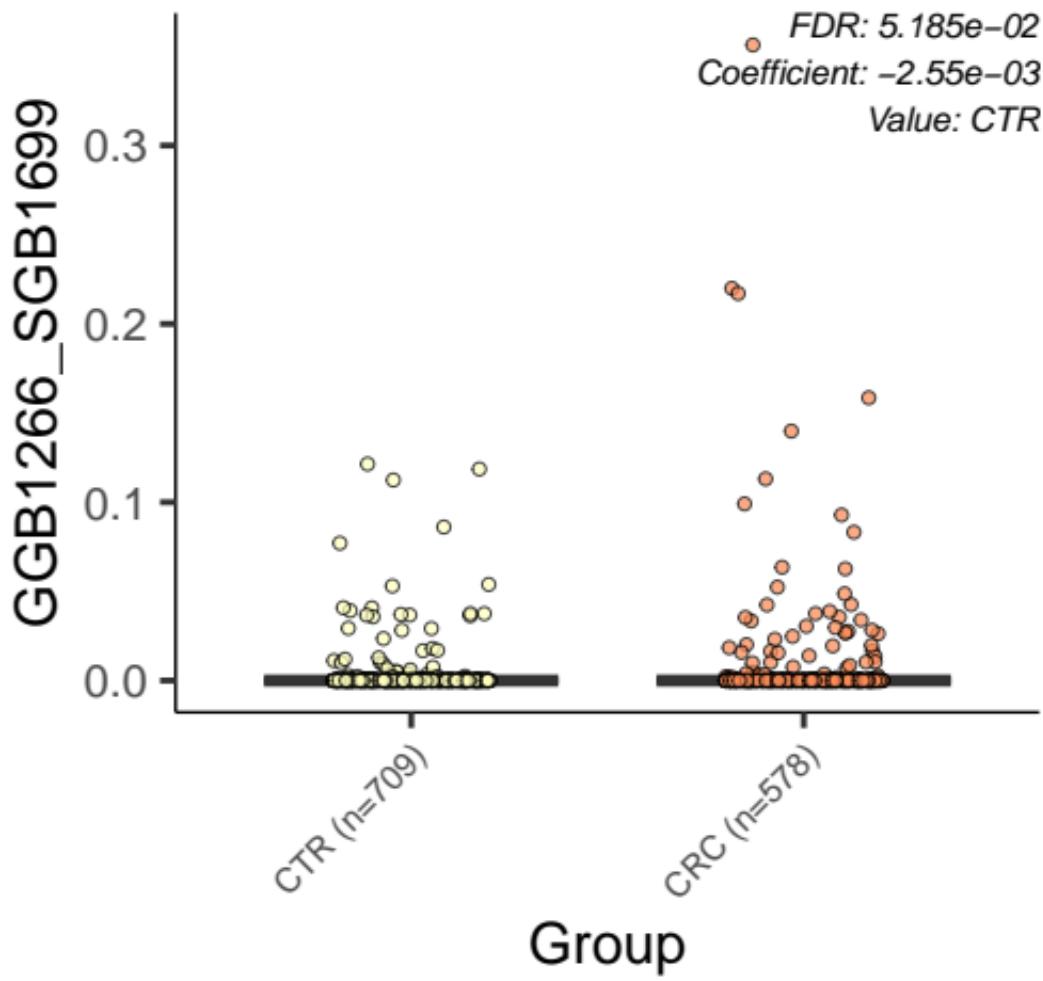


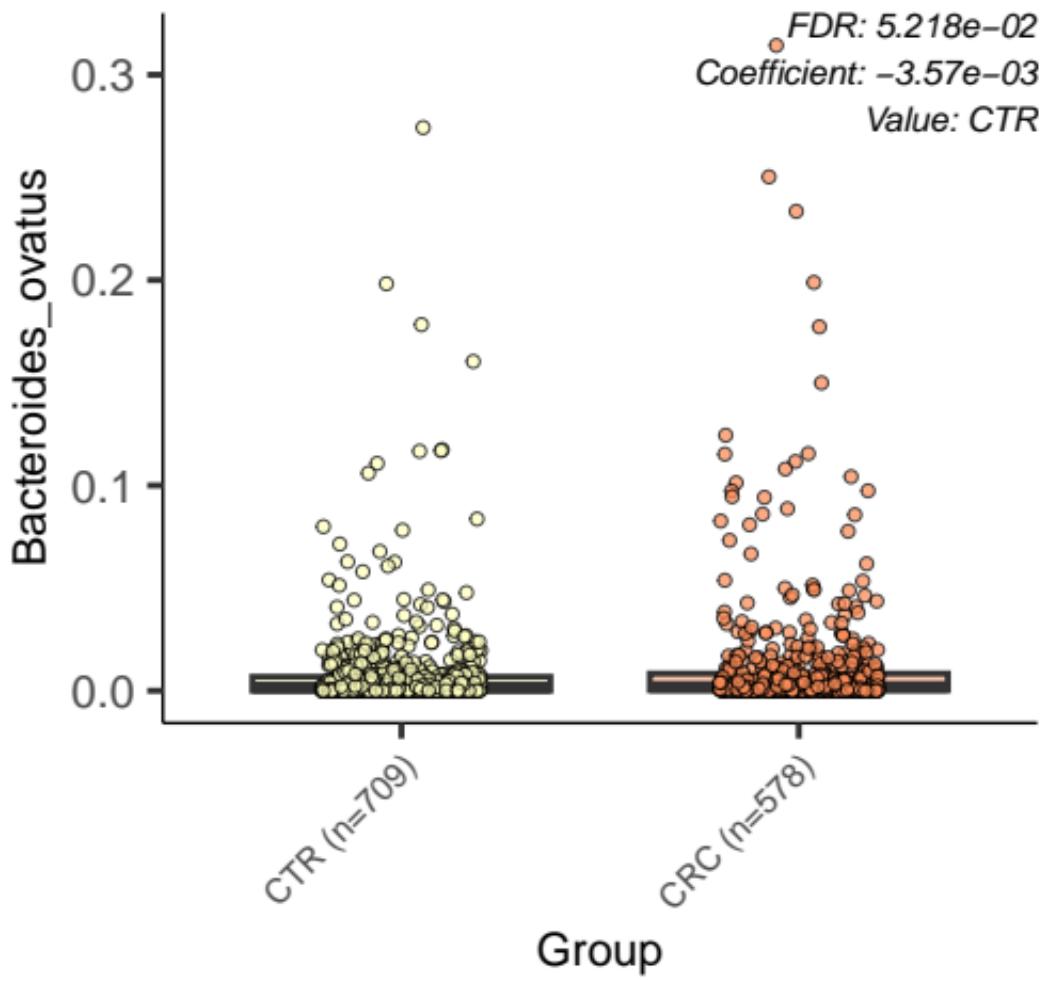


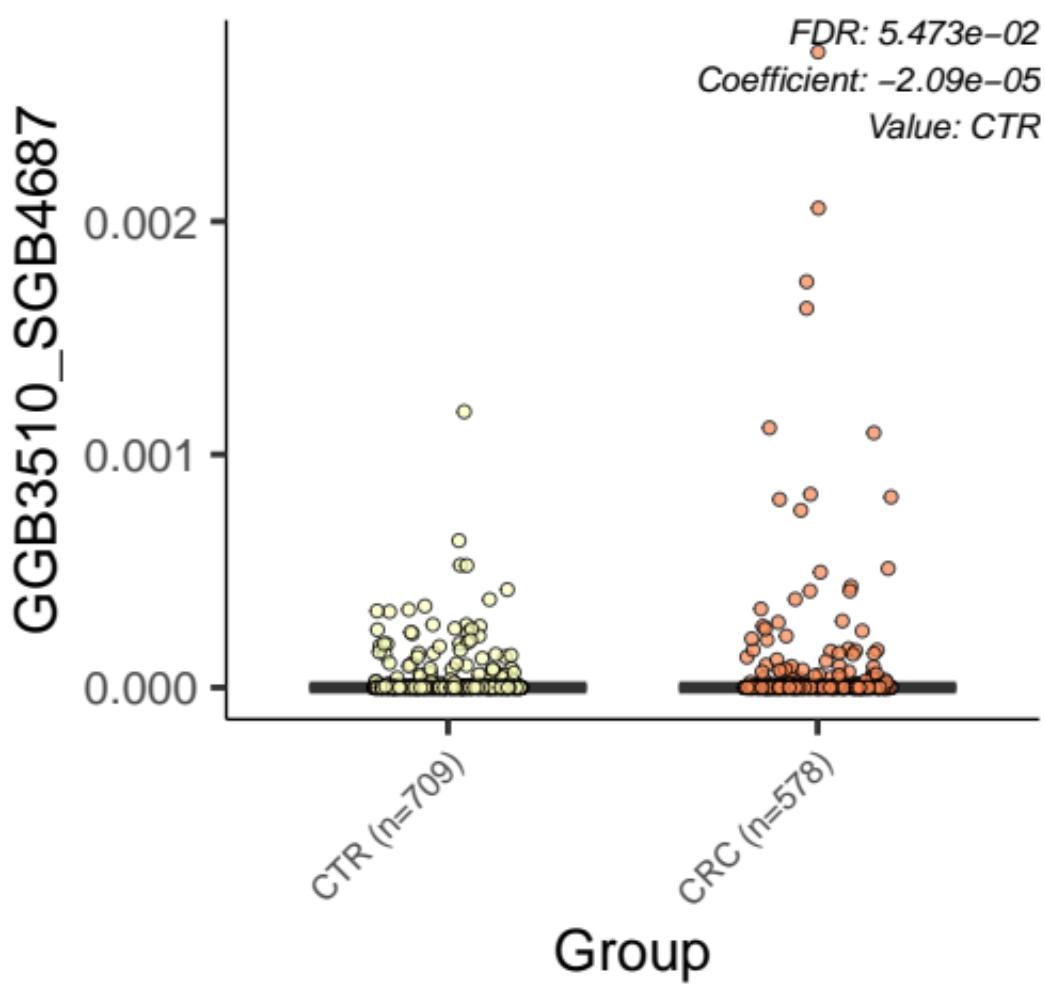


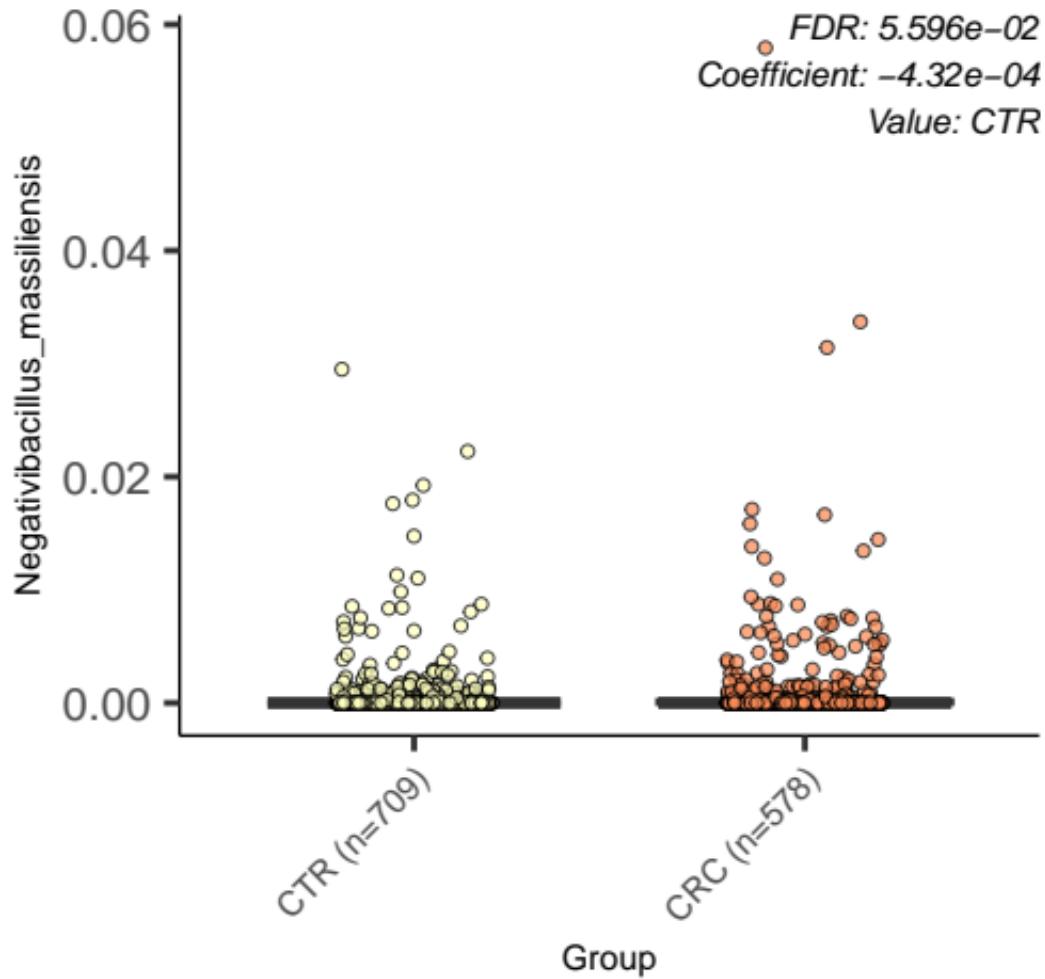


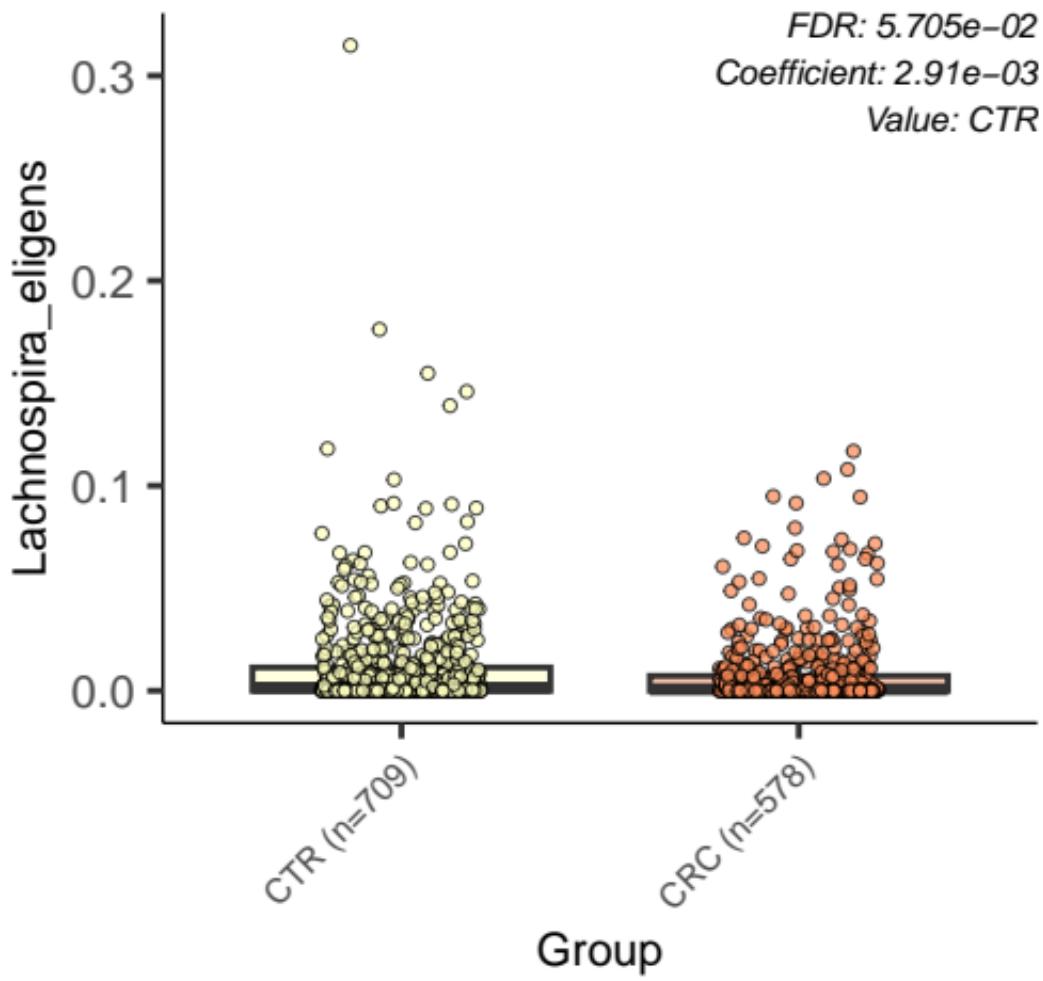


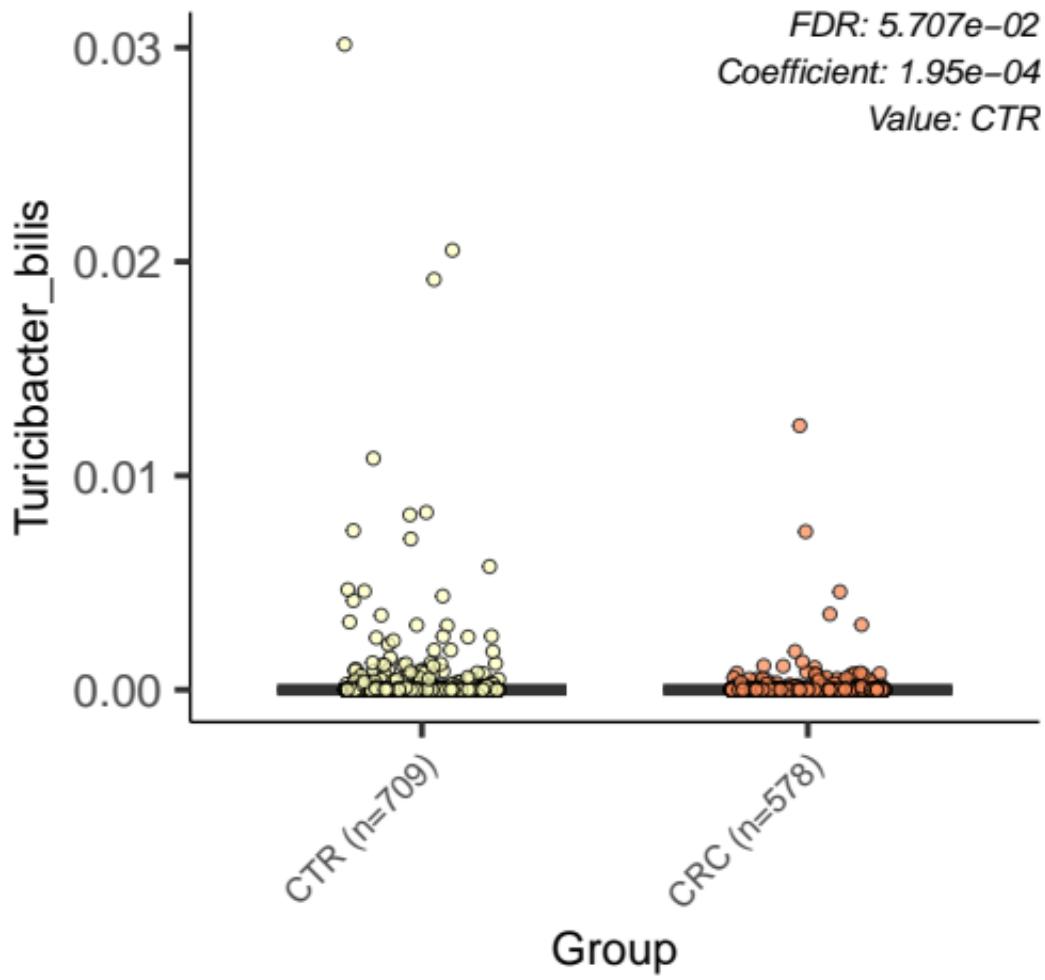


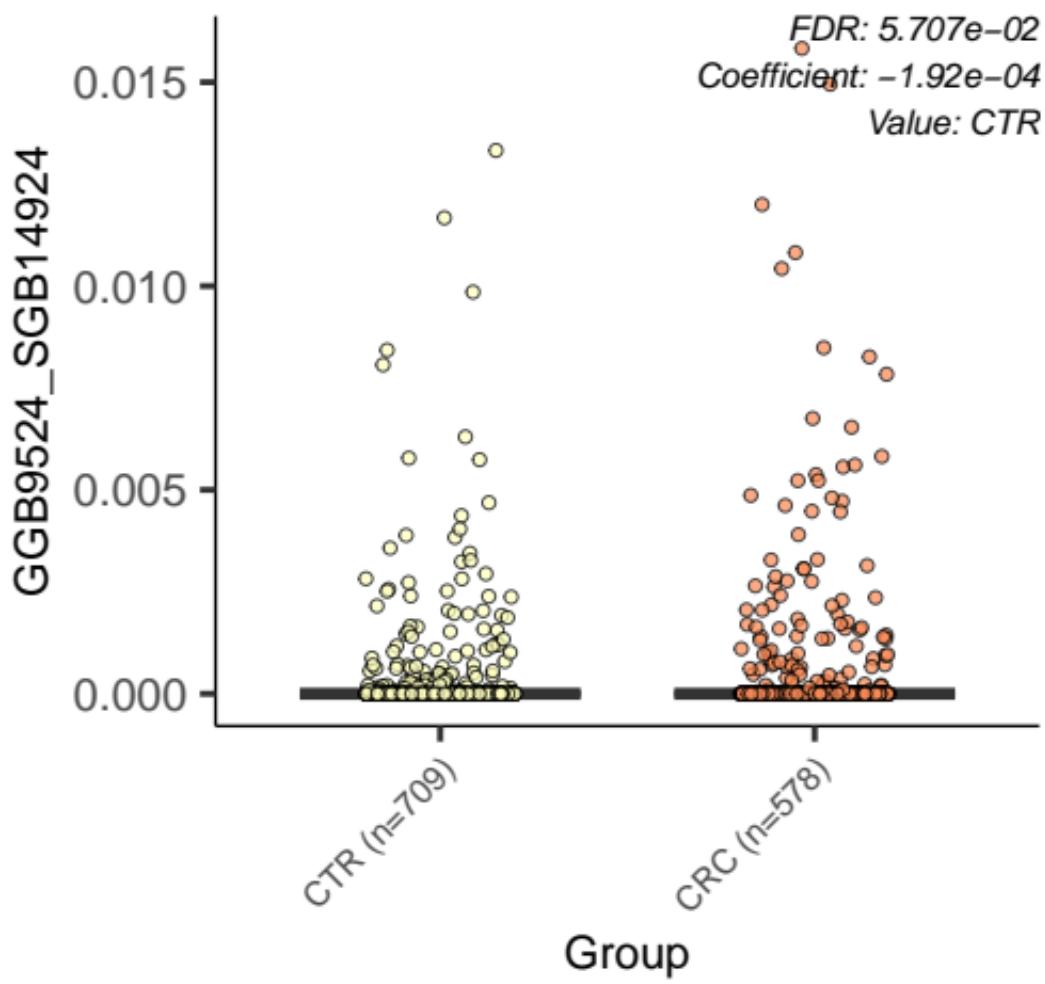


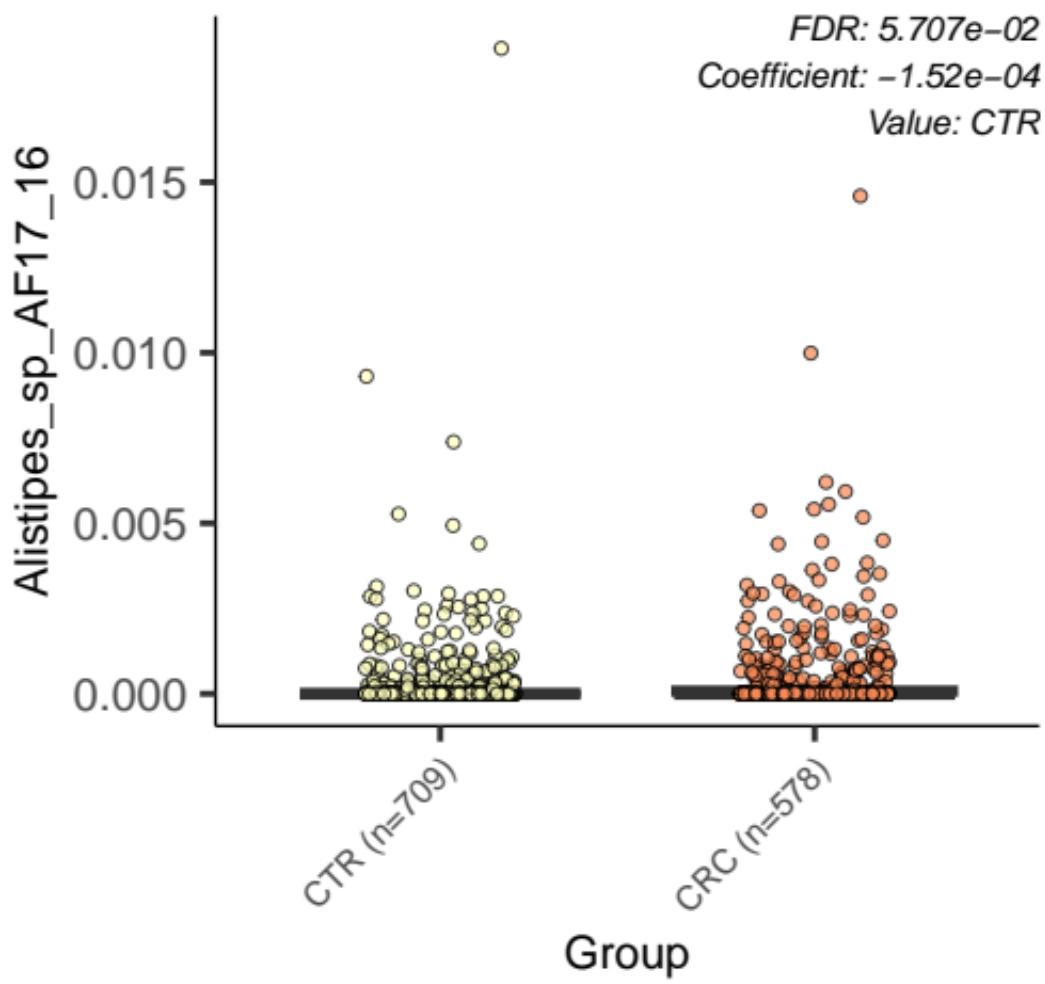


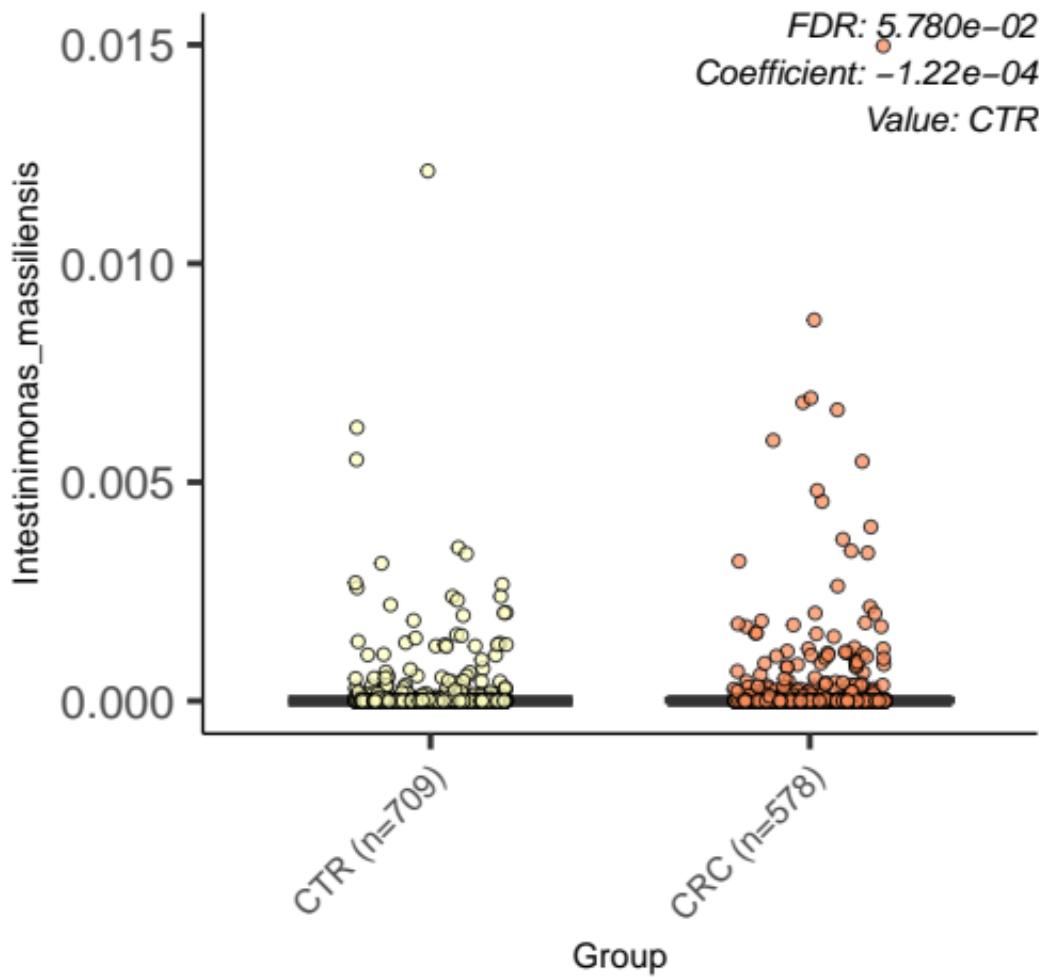


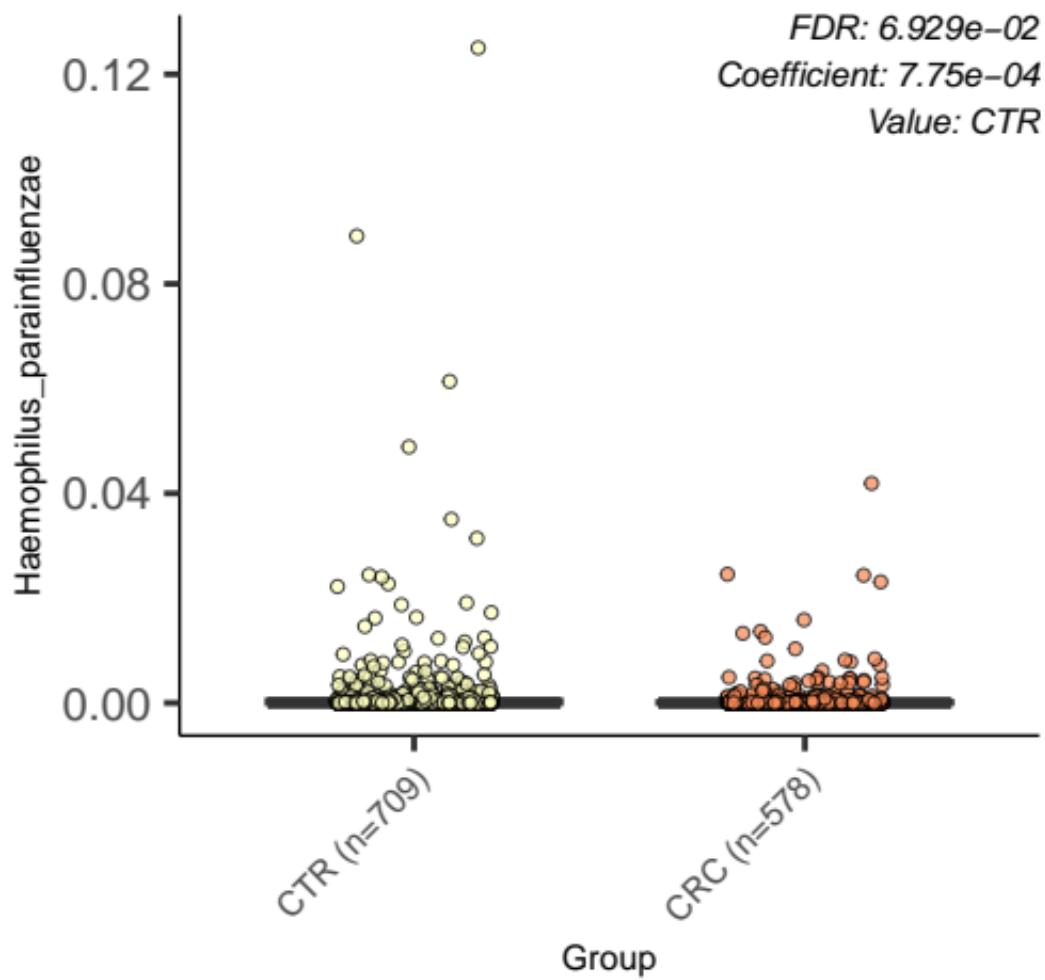


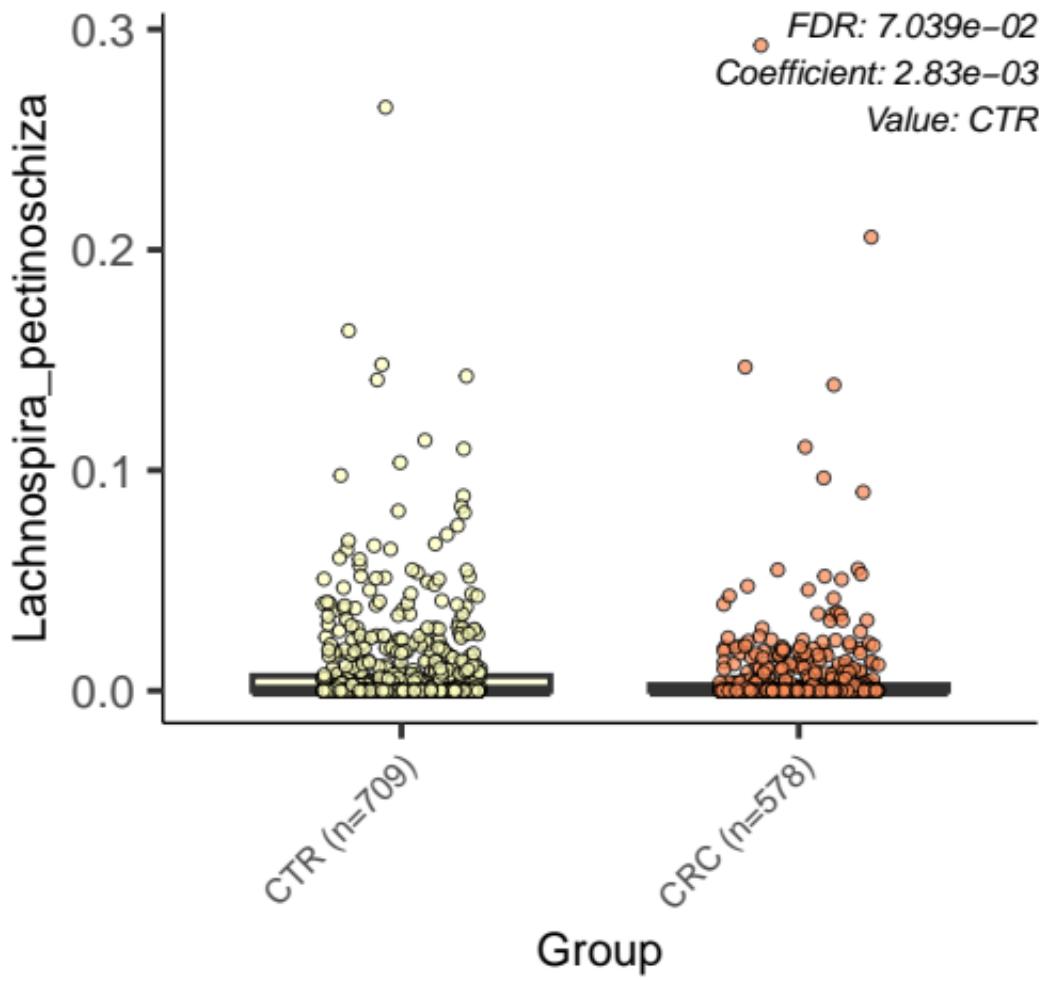


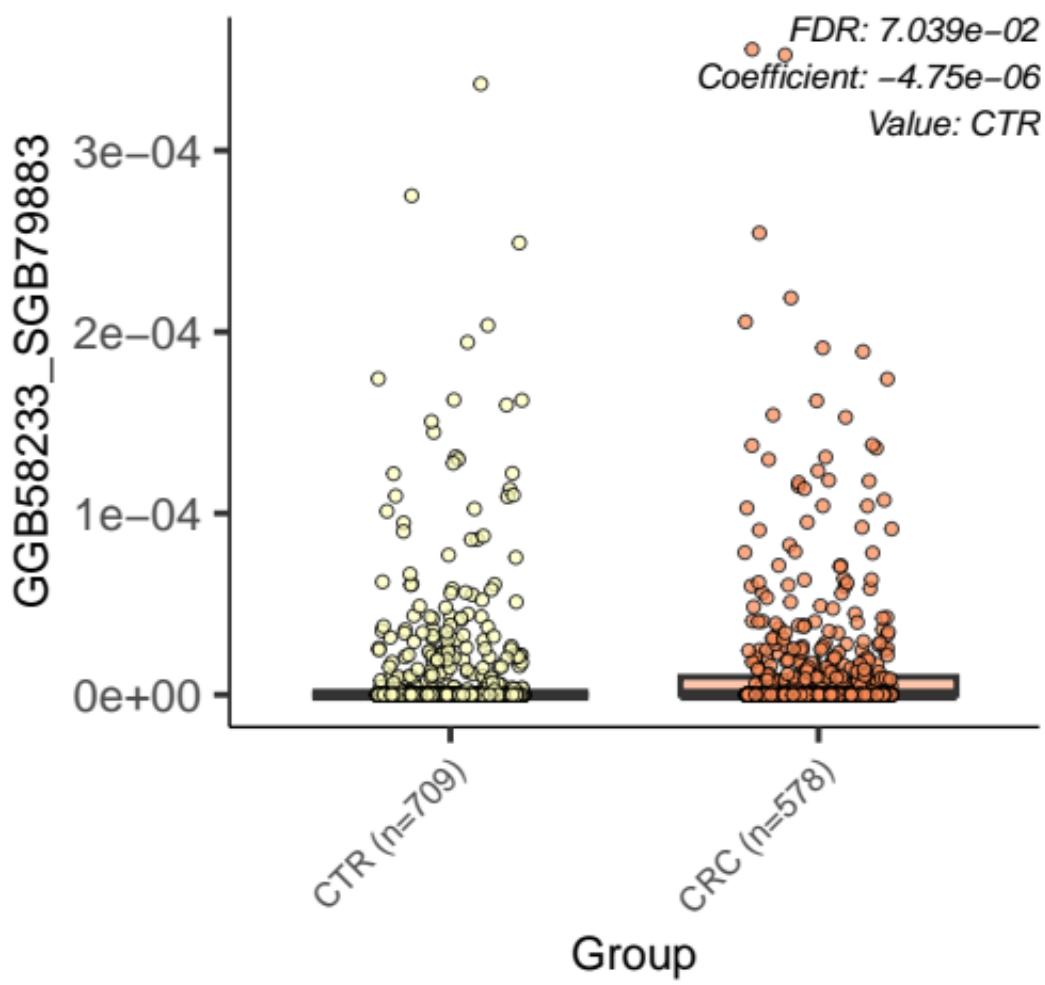


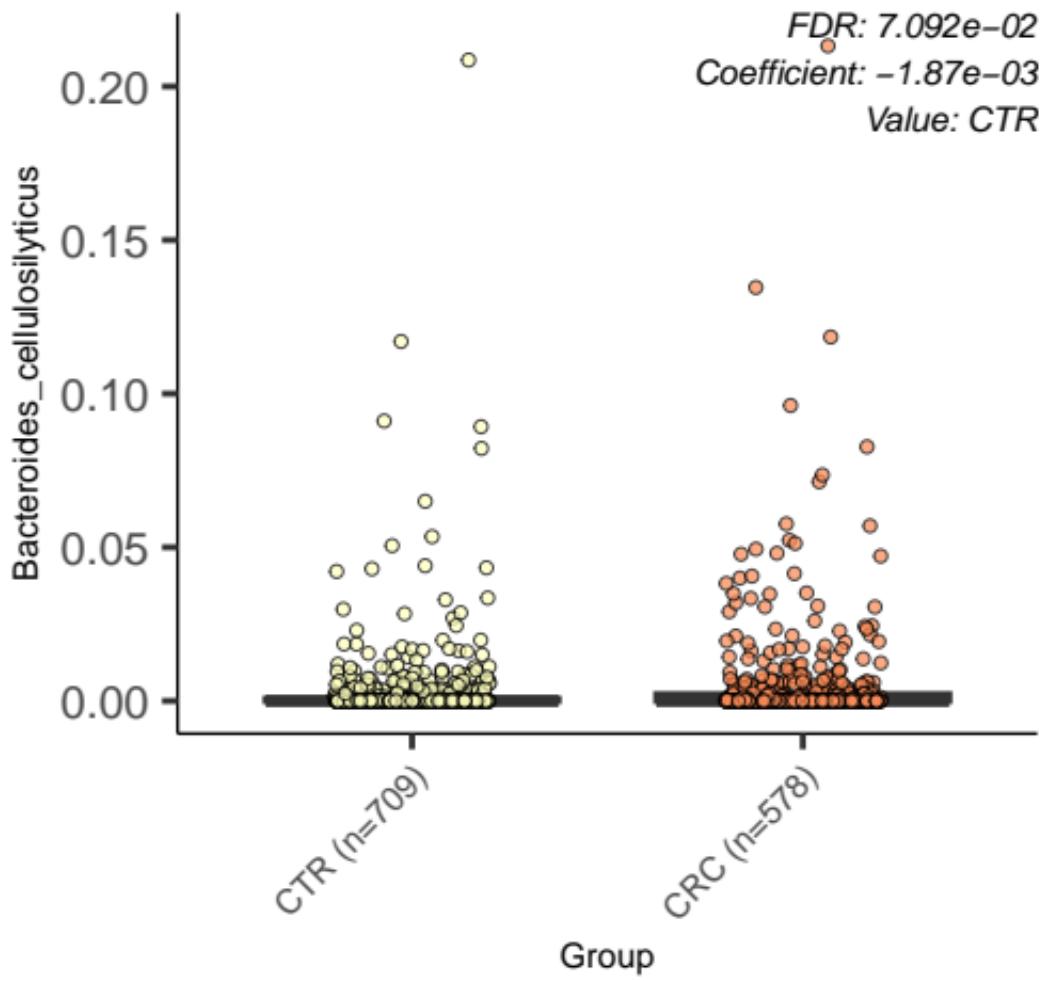


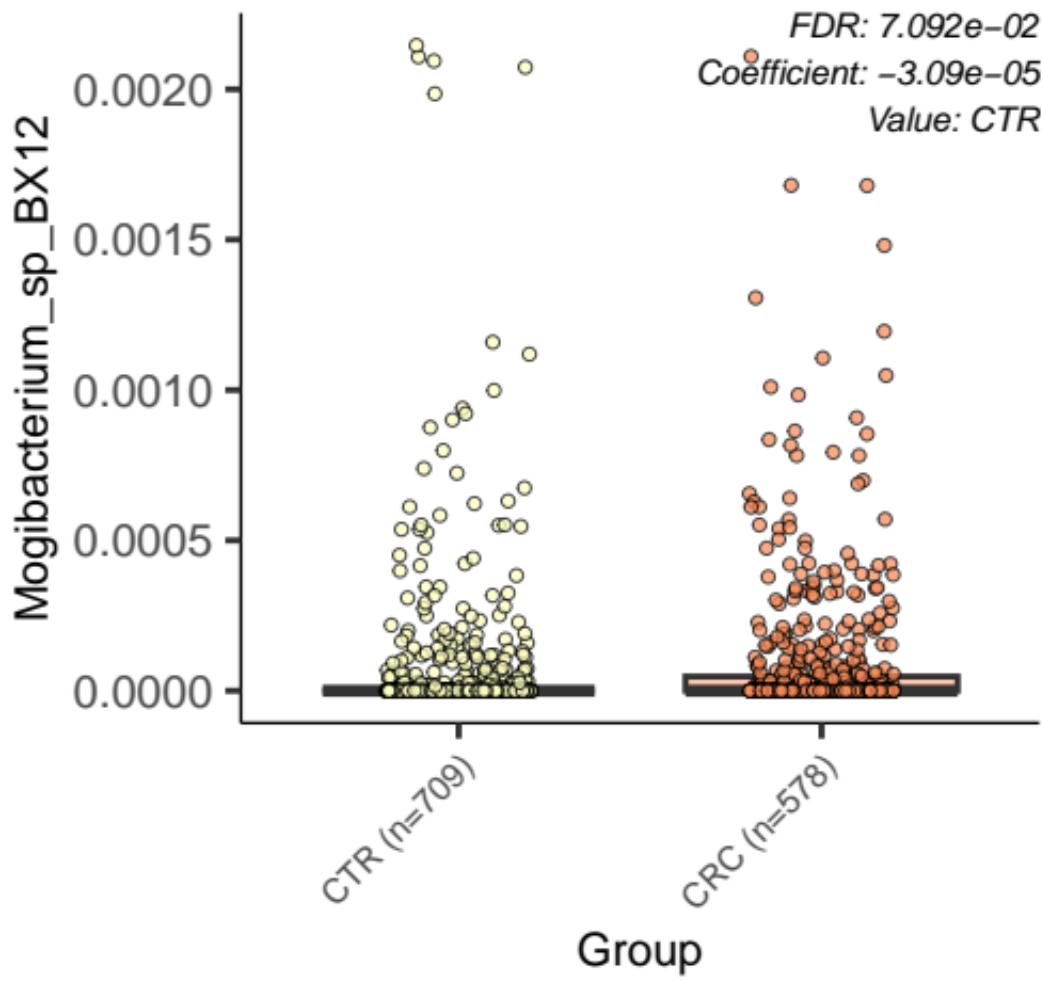


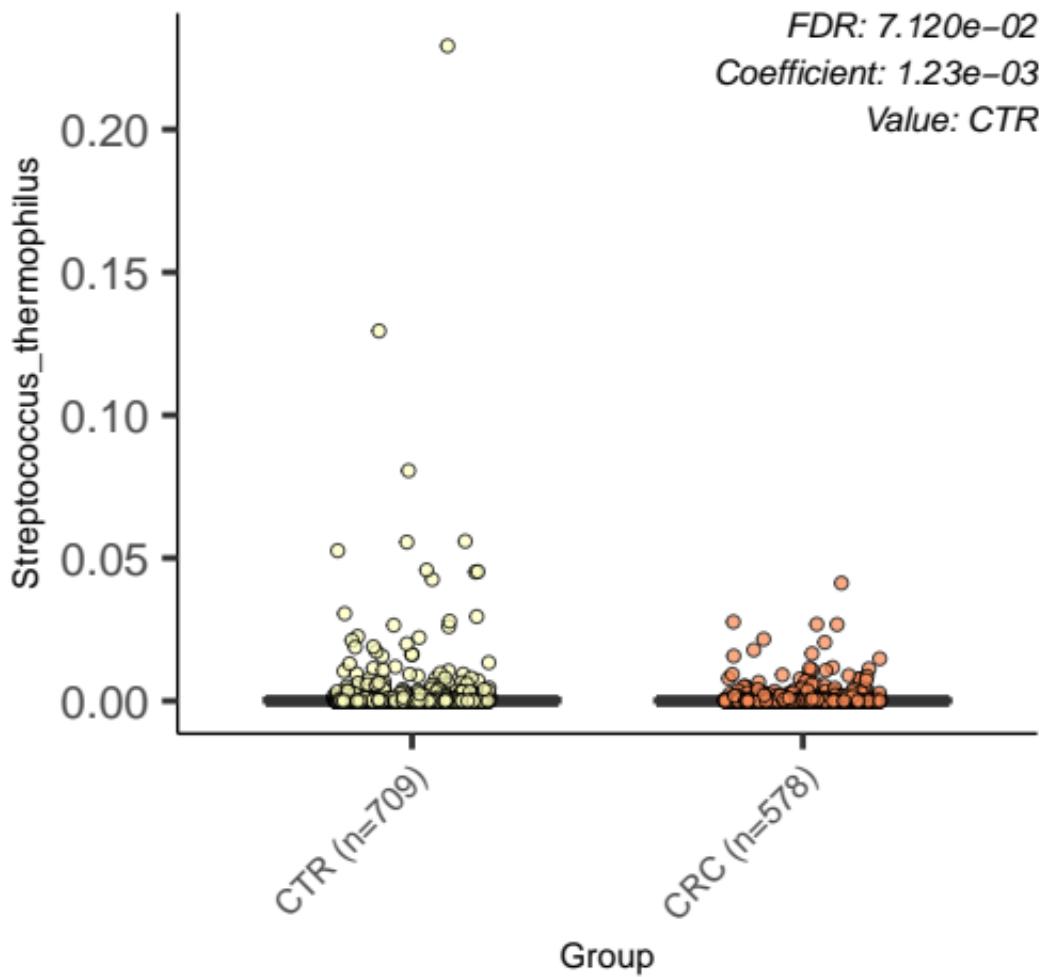


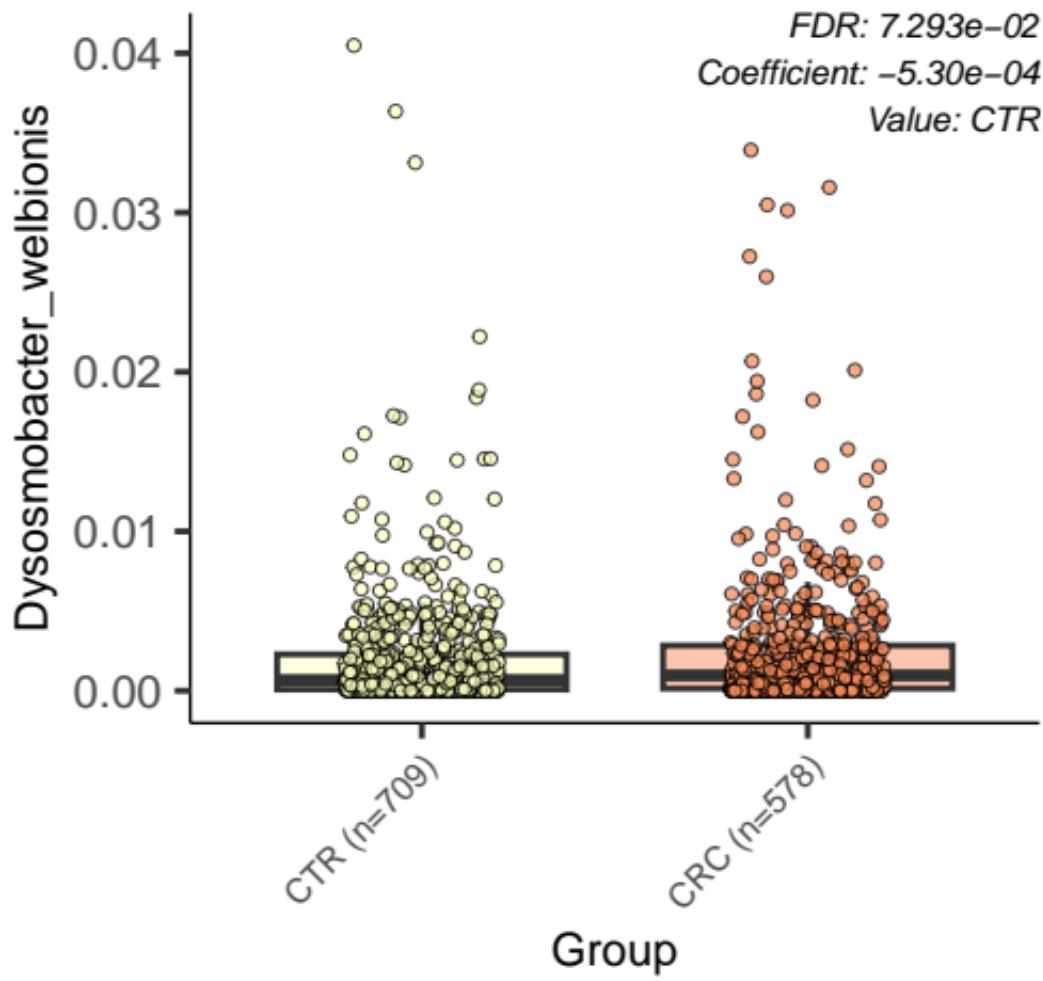


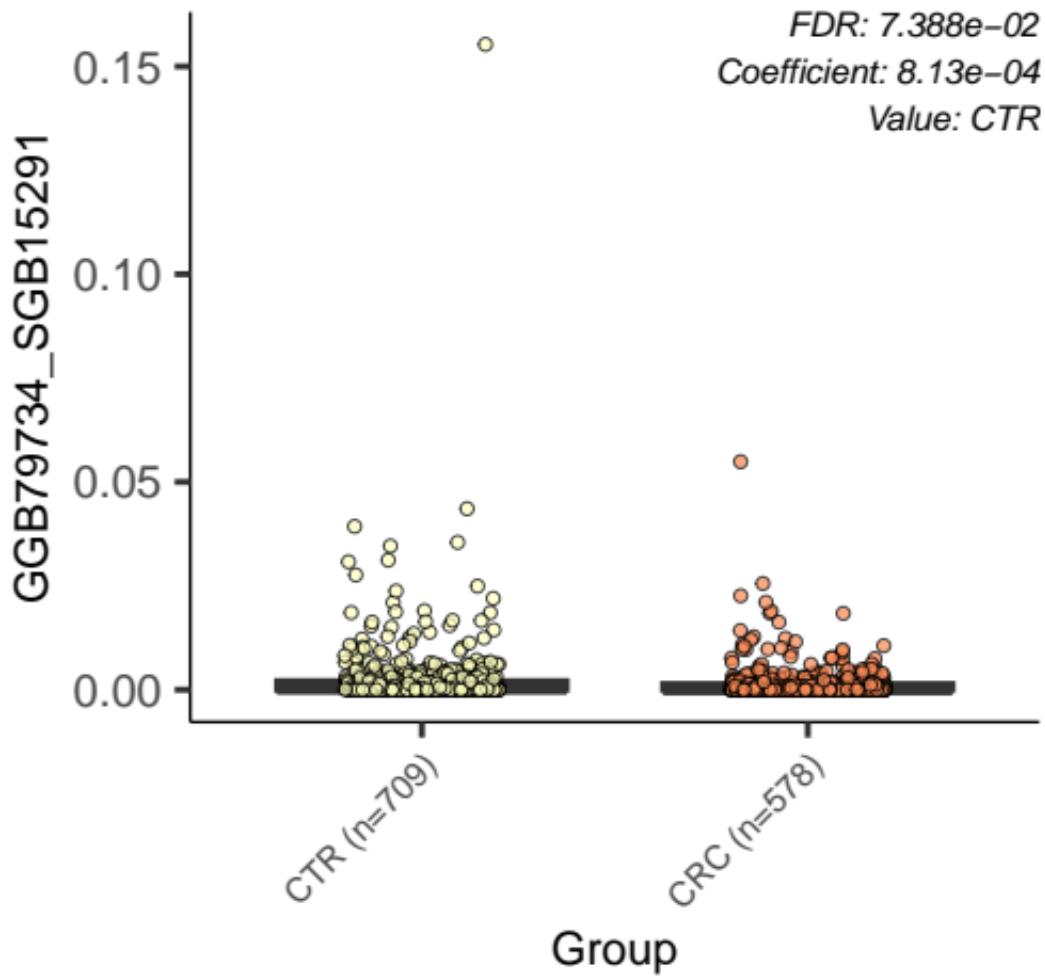


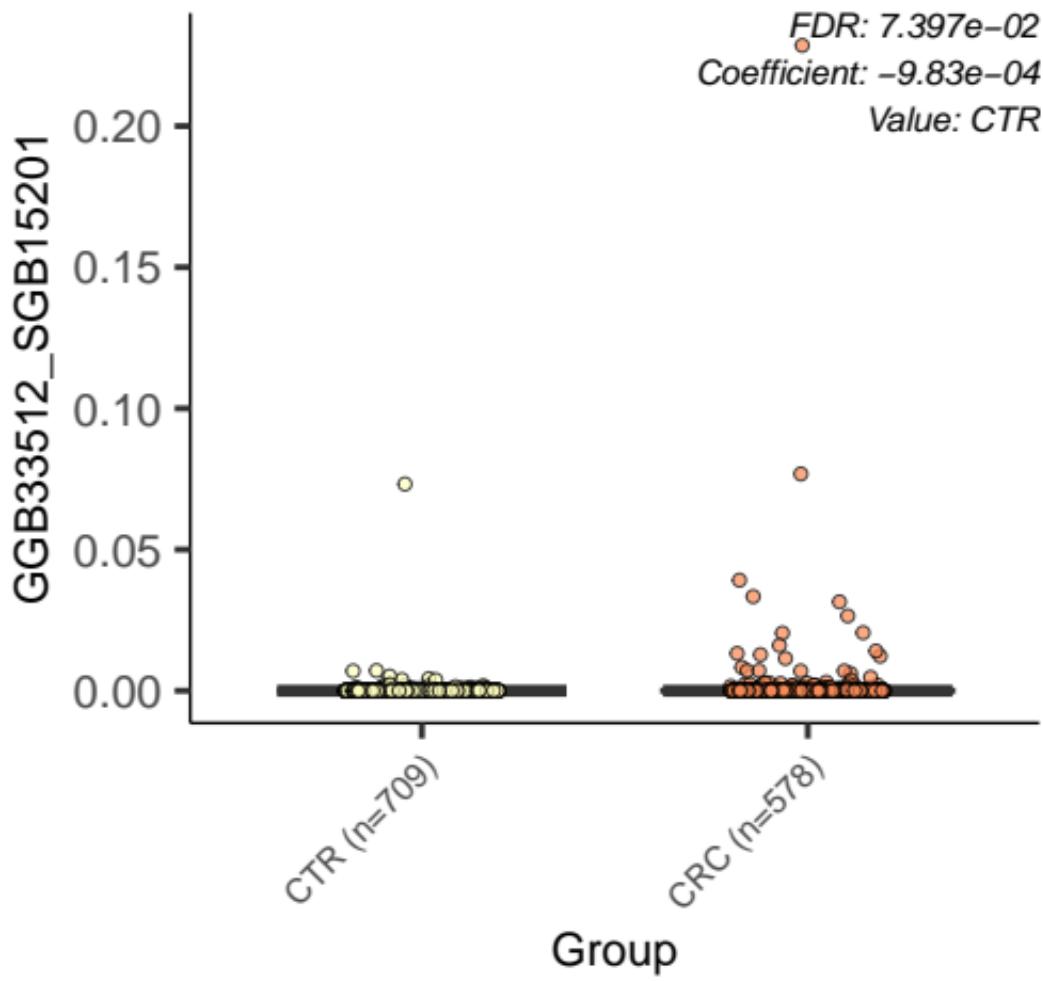


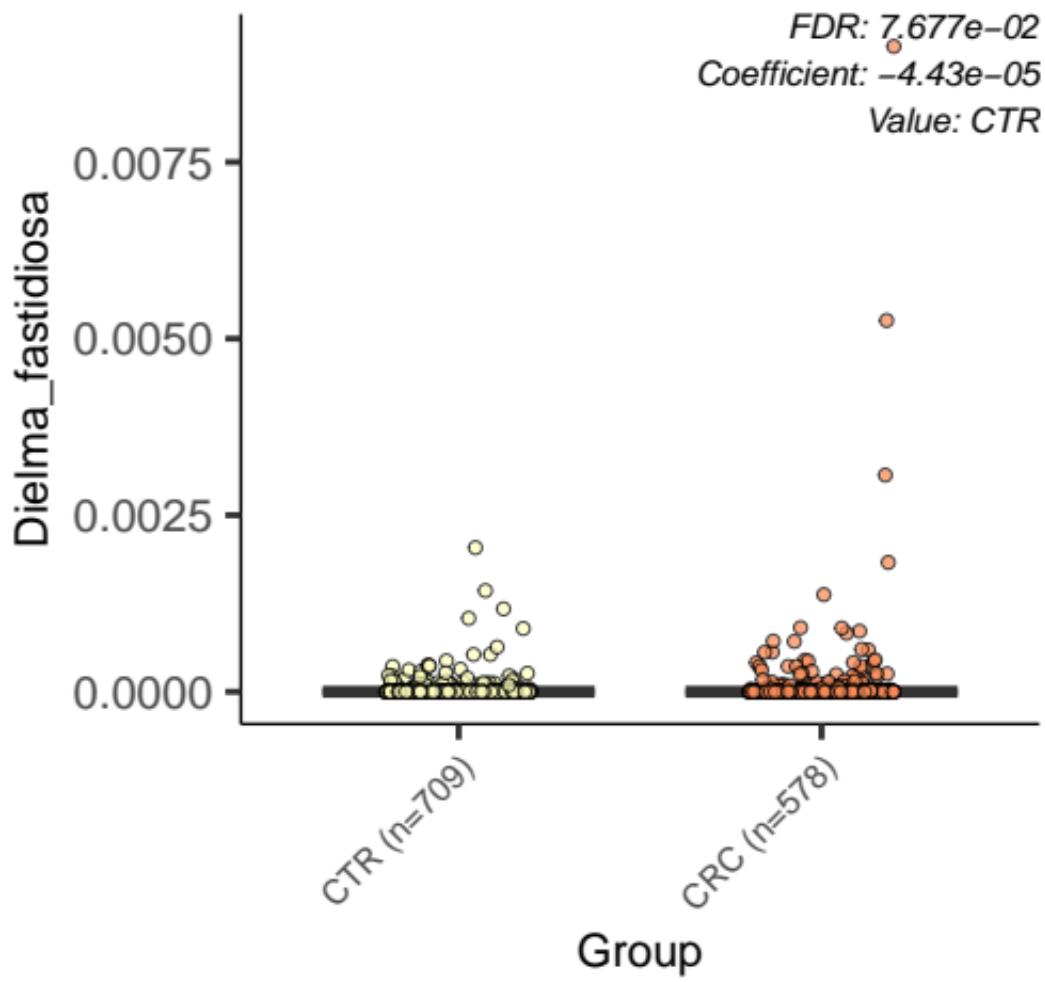


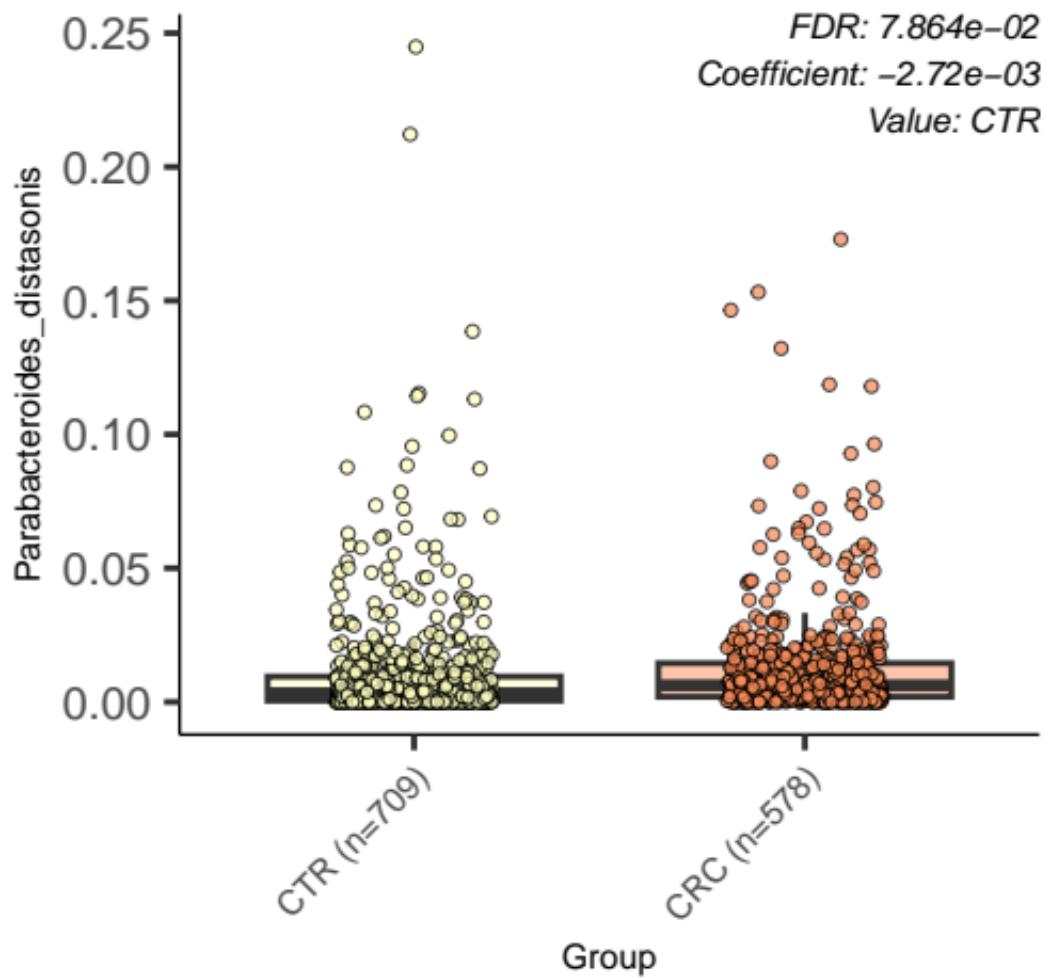


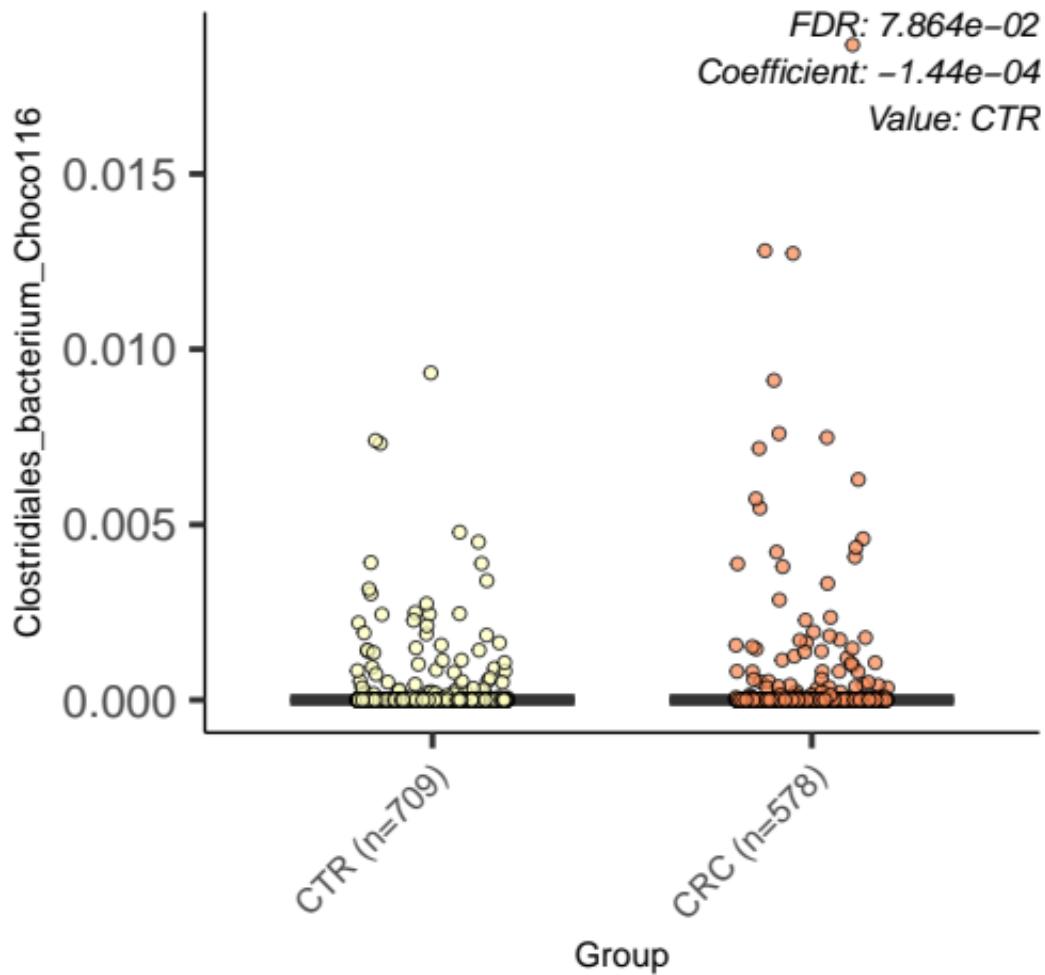


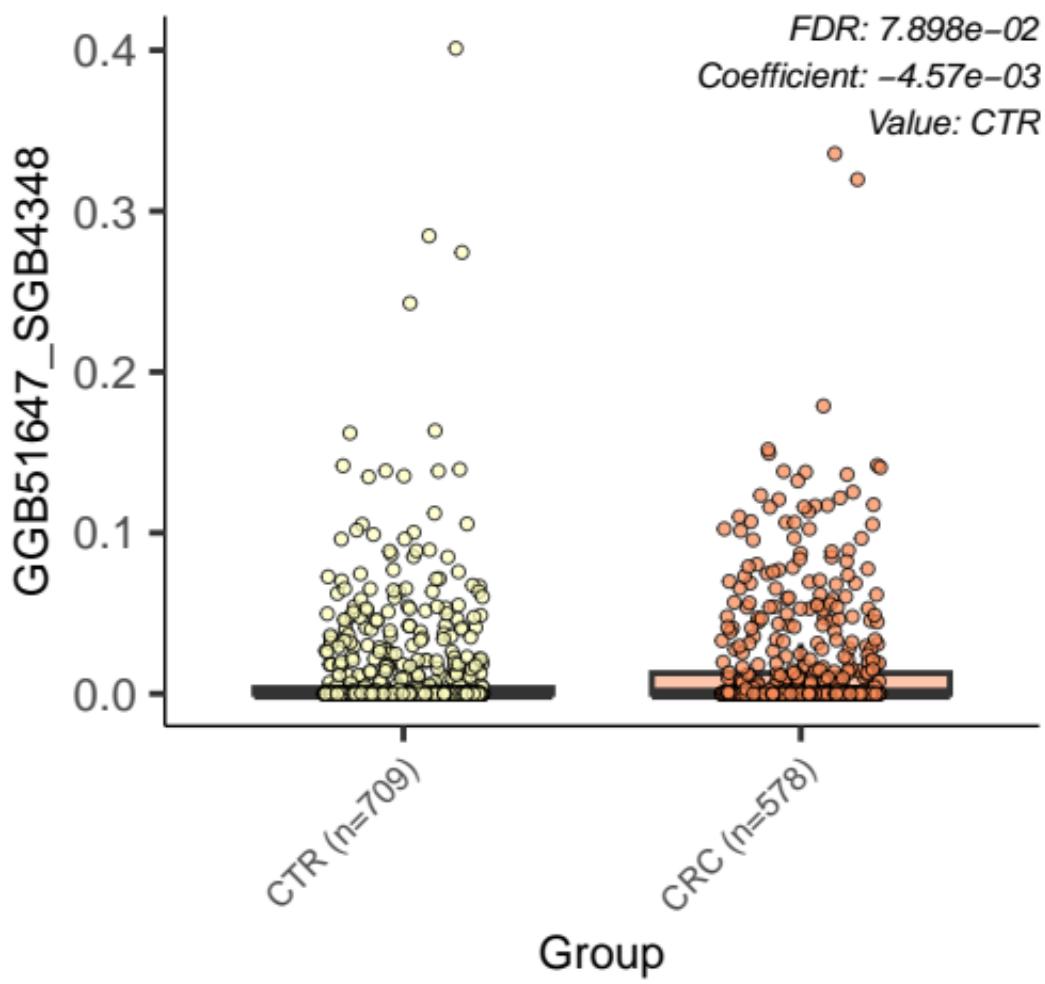


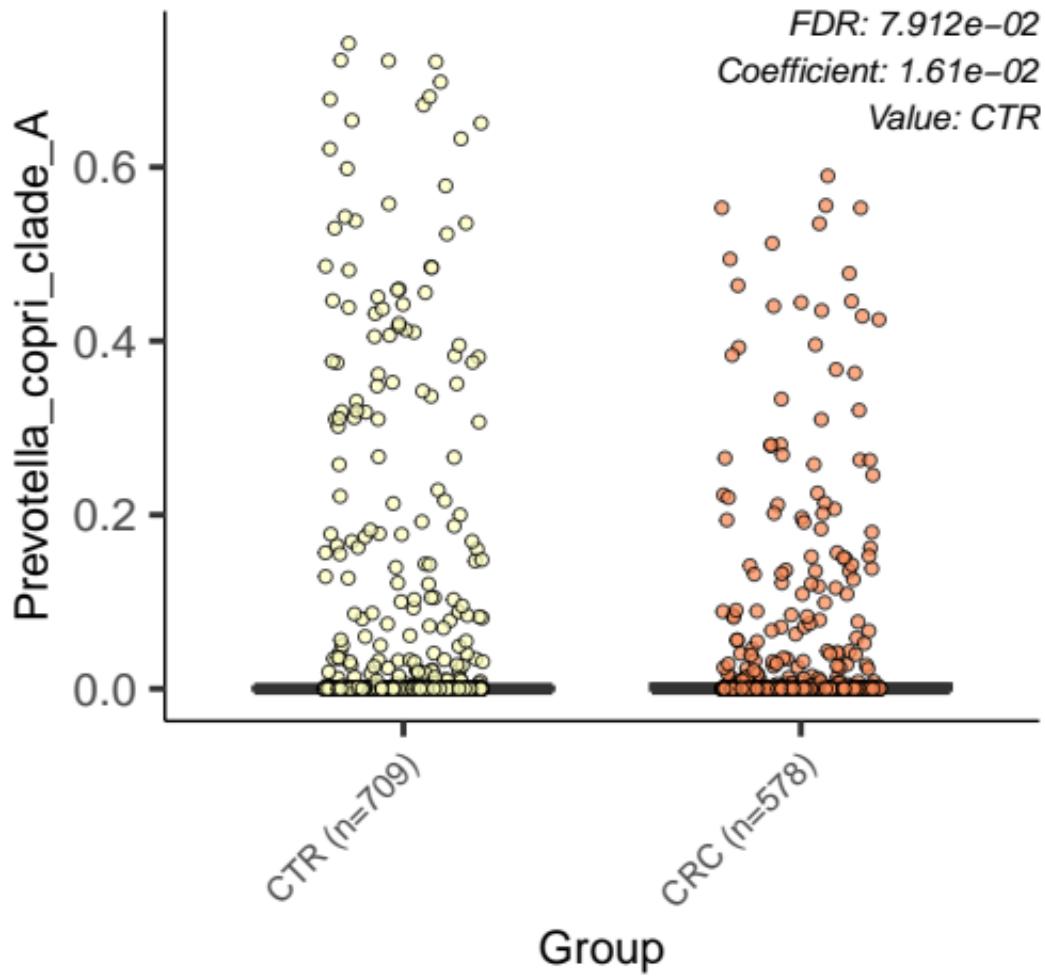


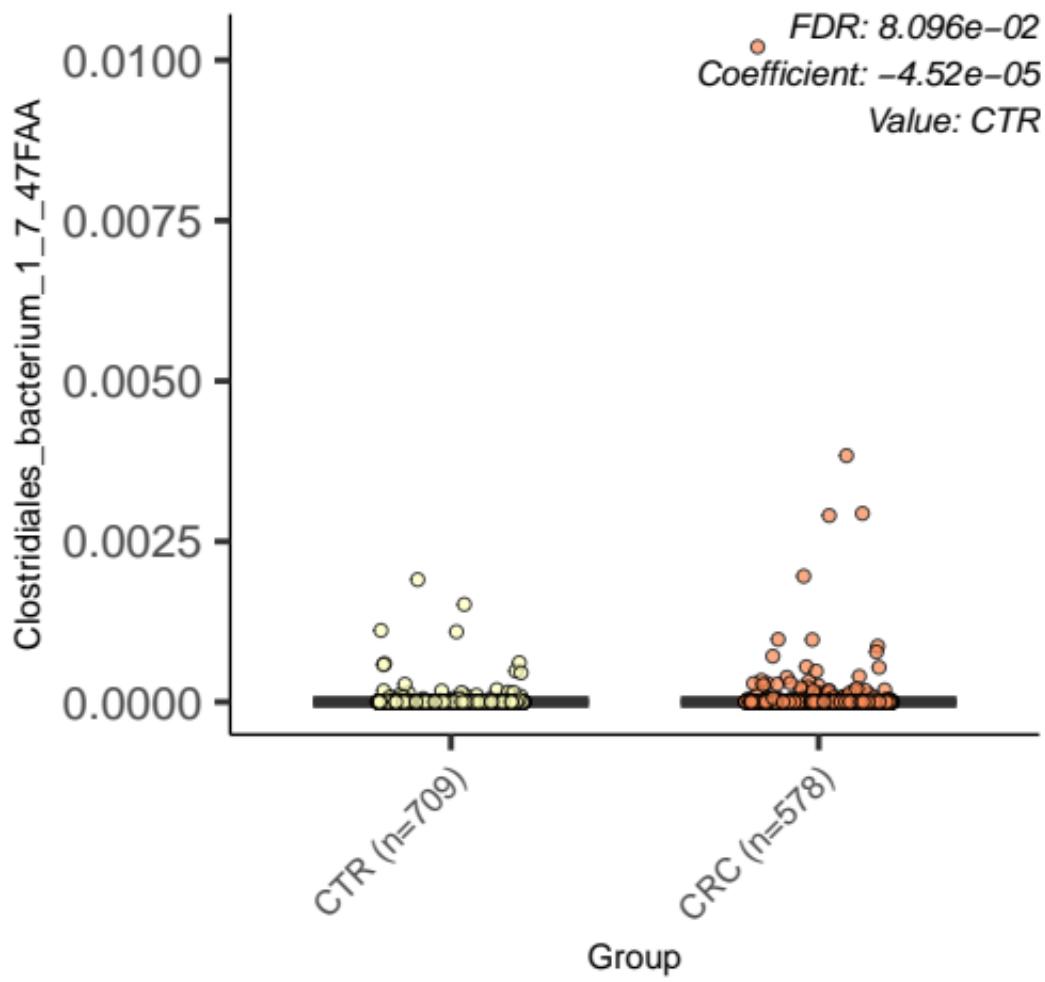


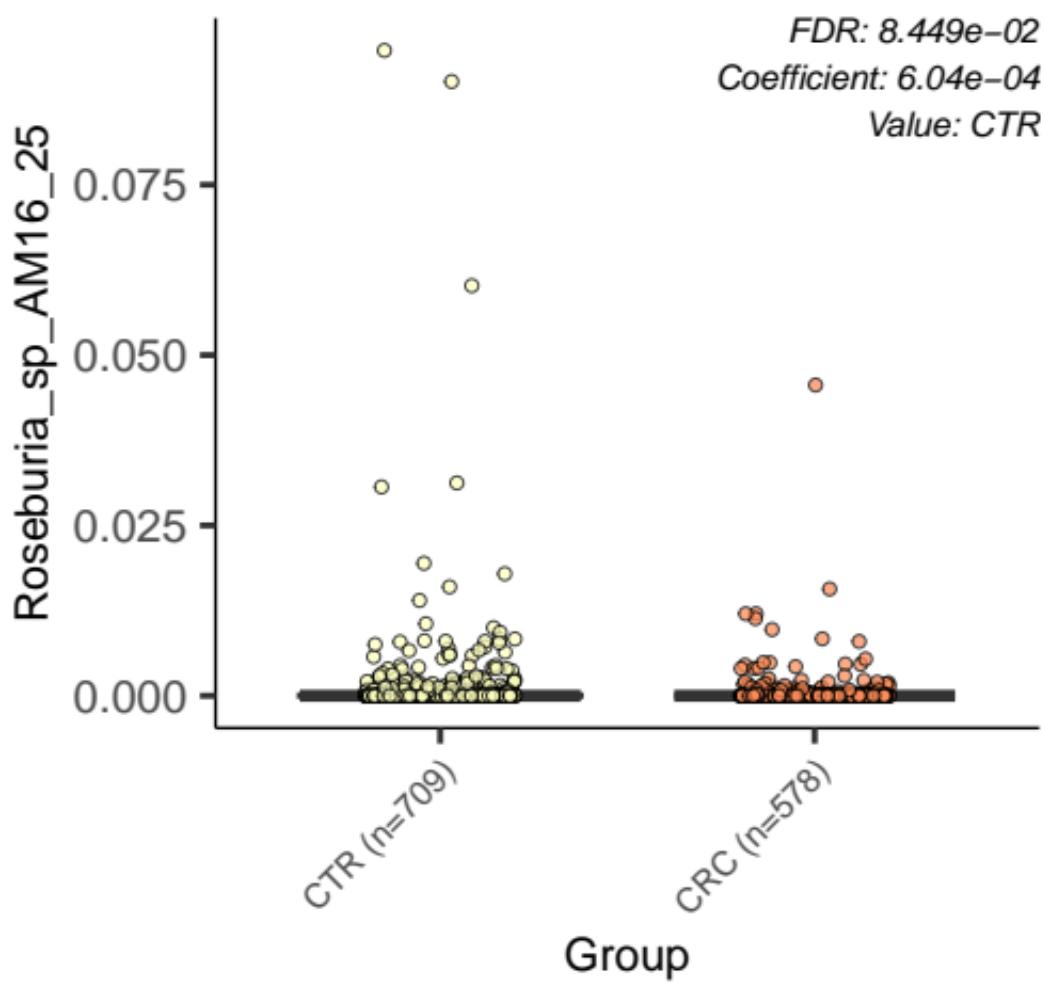


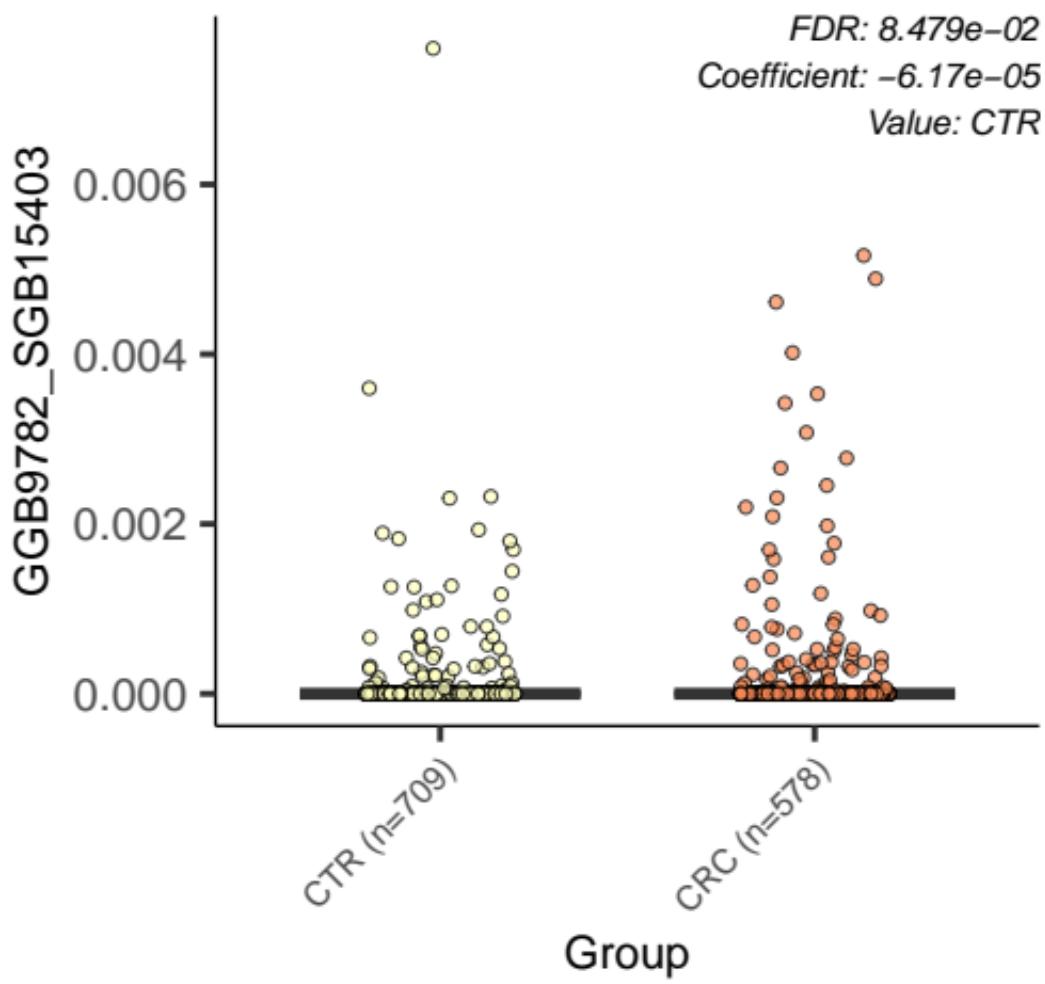


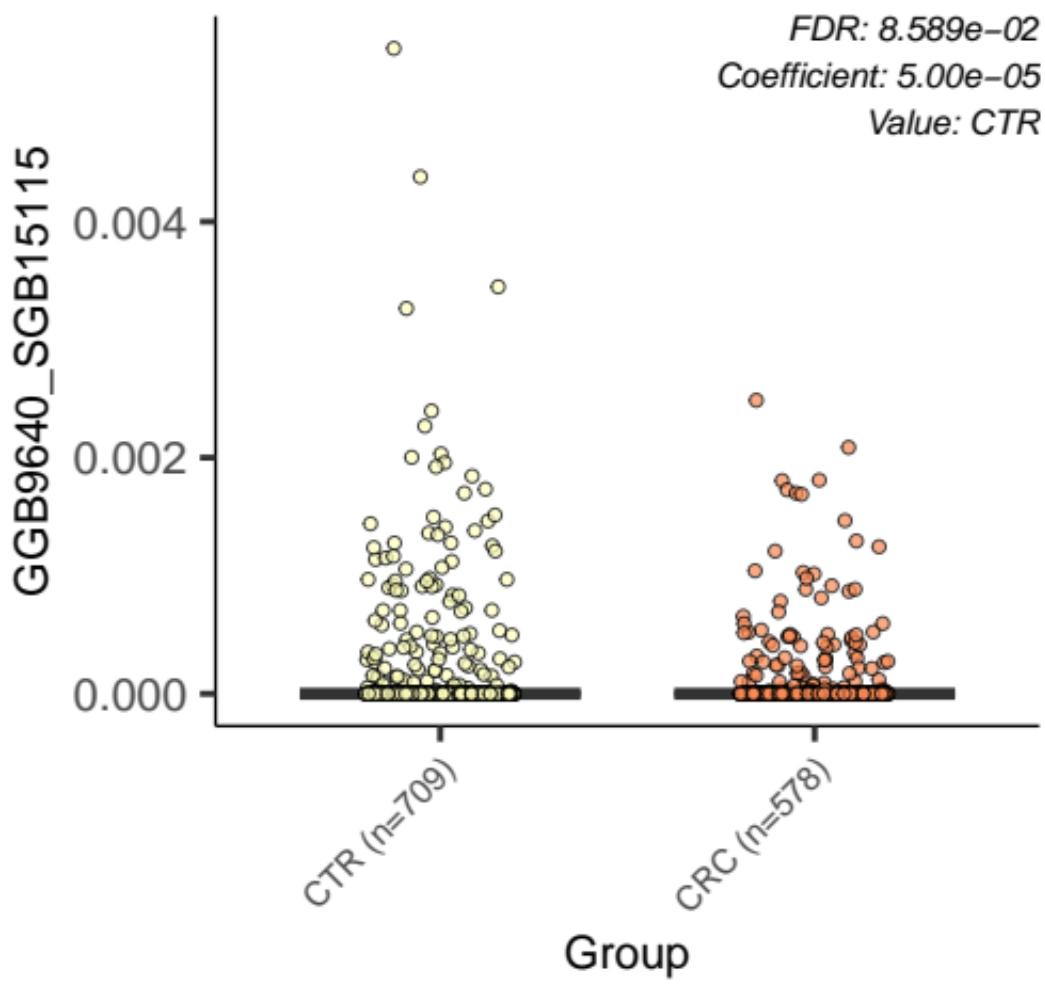


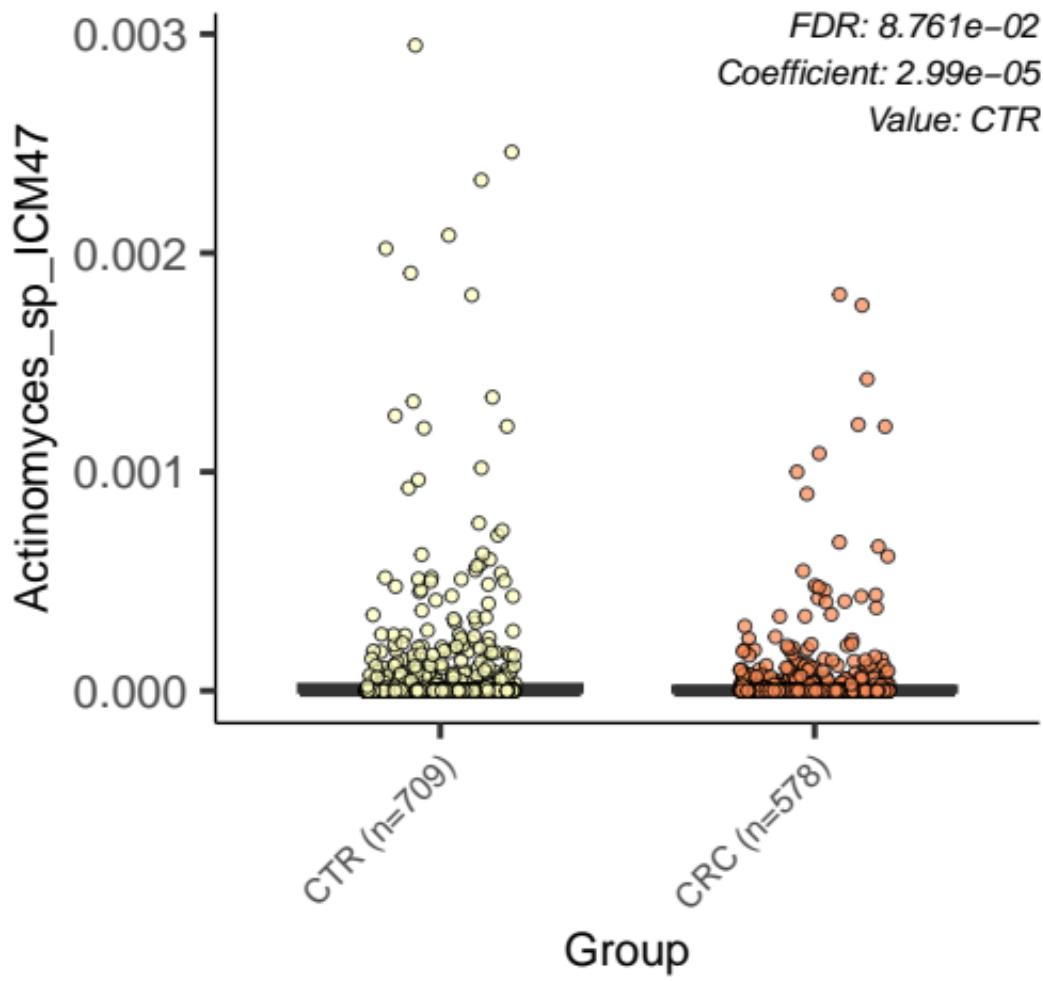












GGB2987_SGB3969

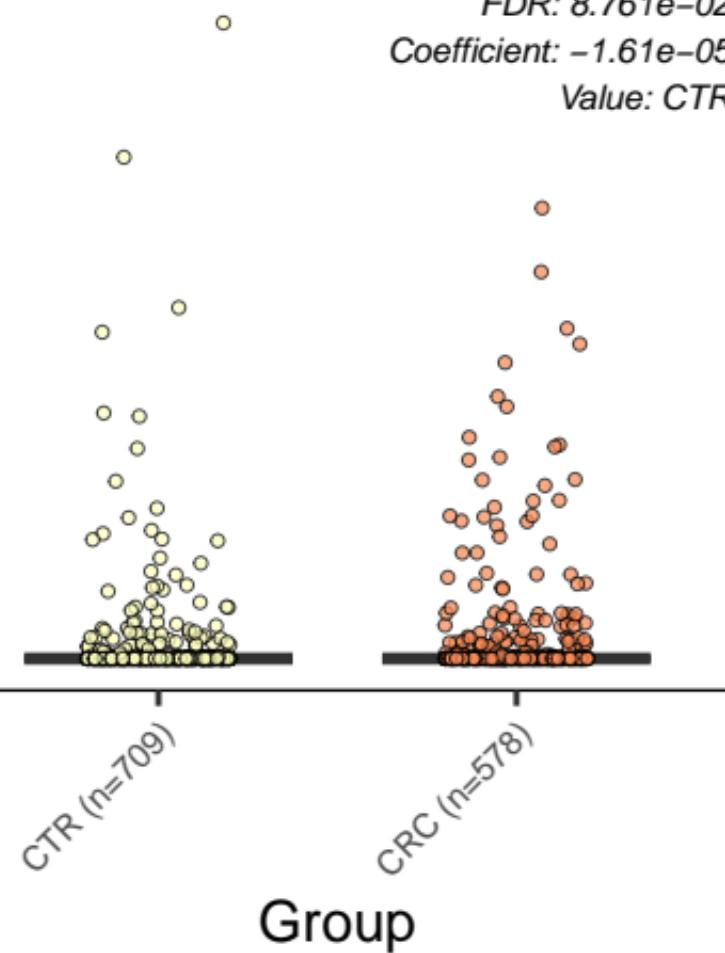
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Value: CTR

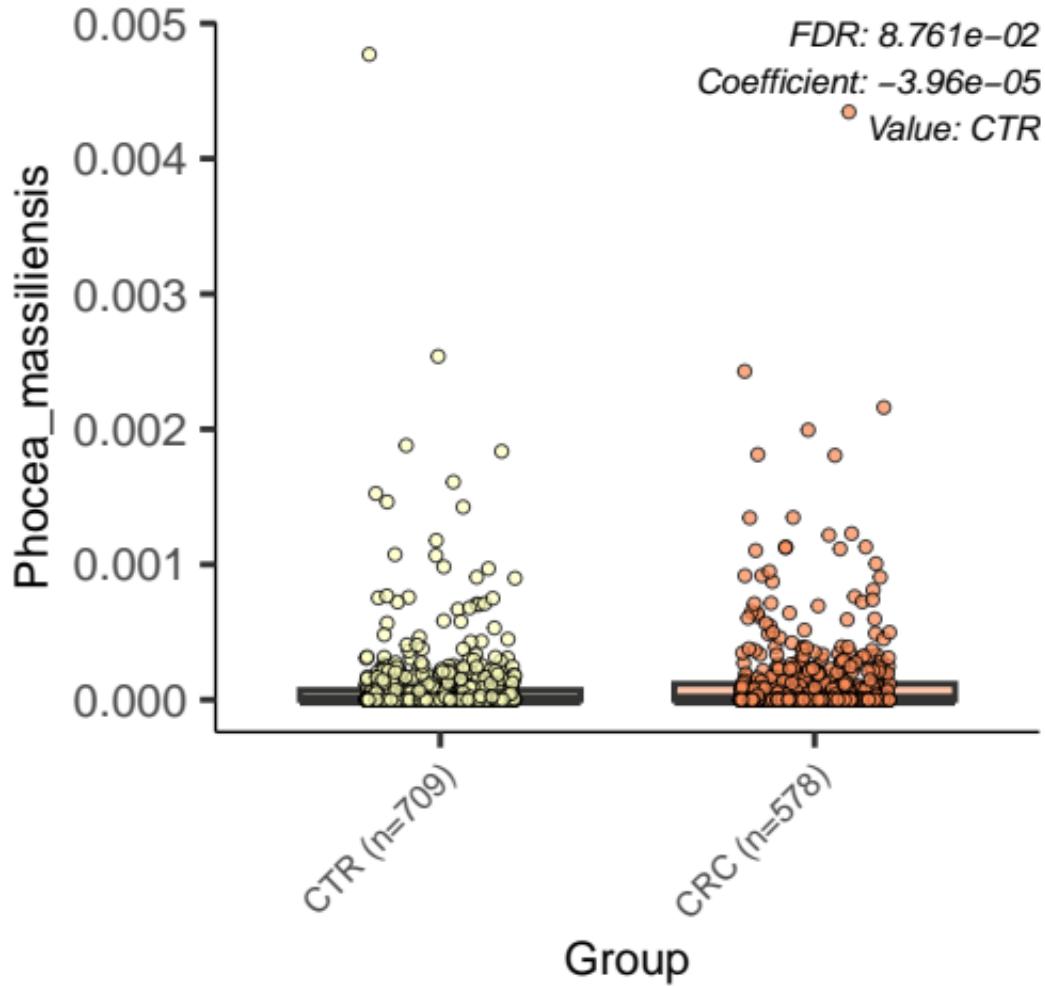
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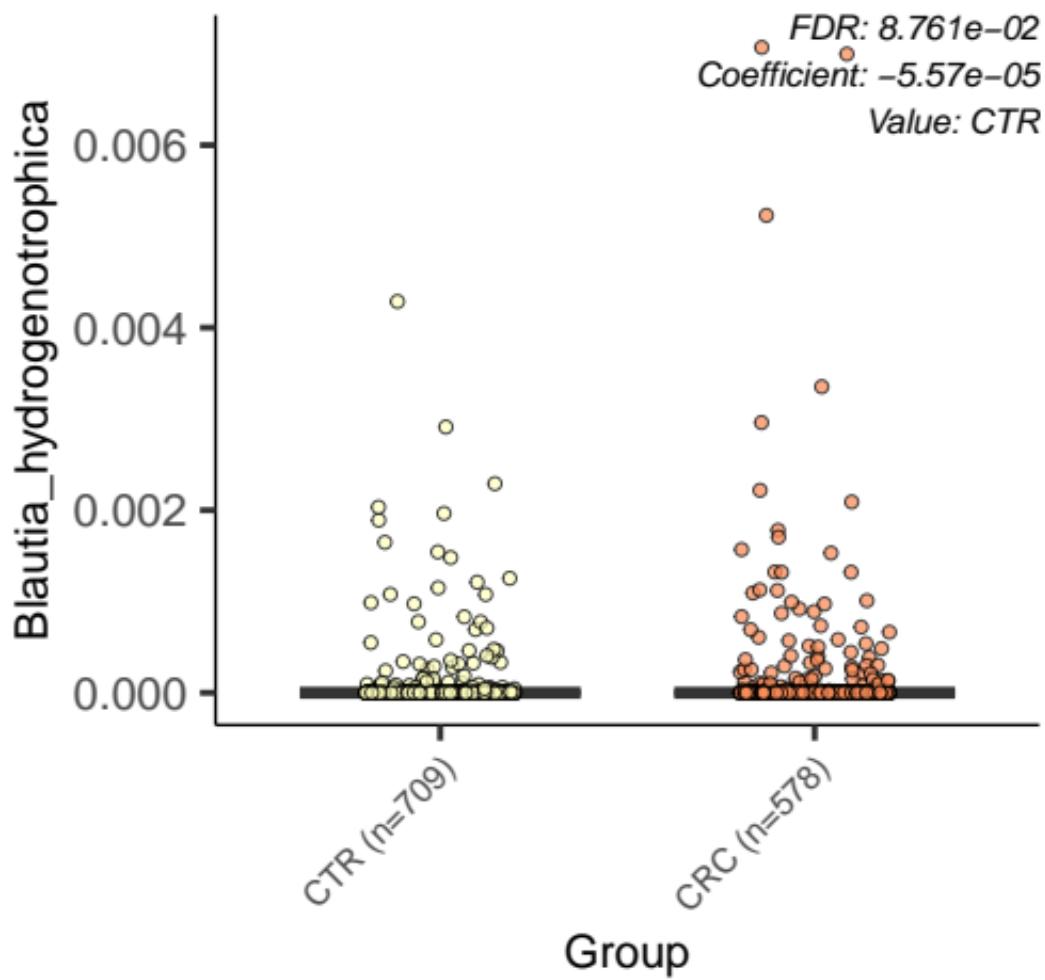
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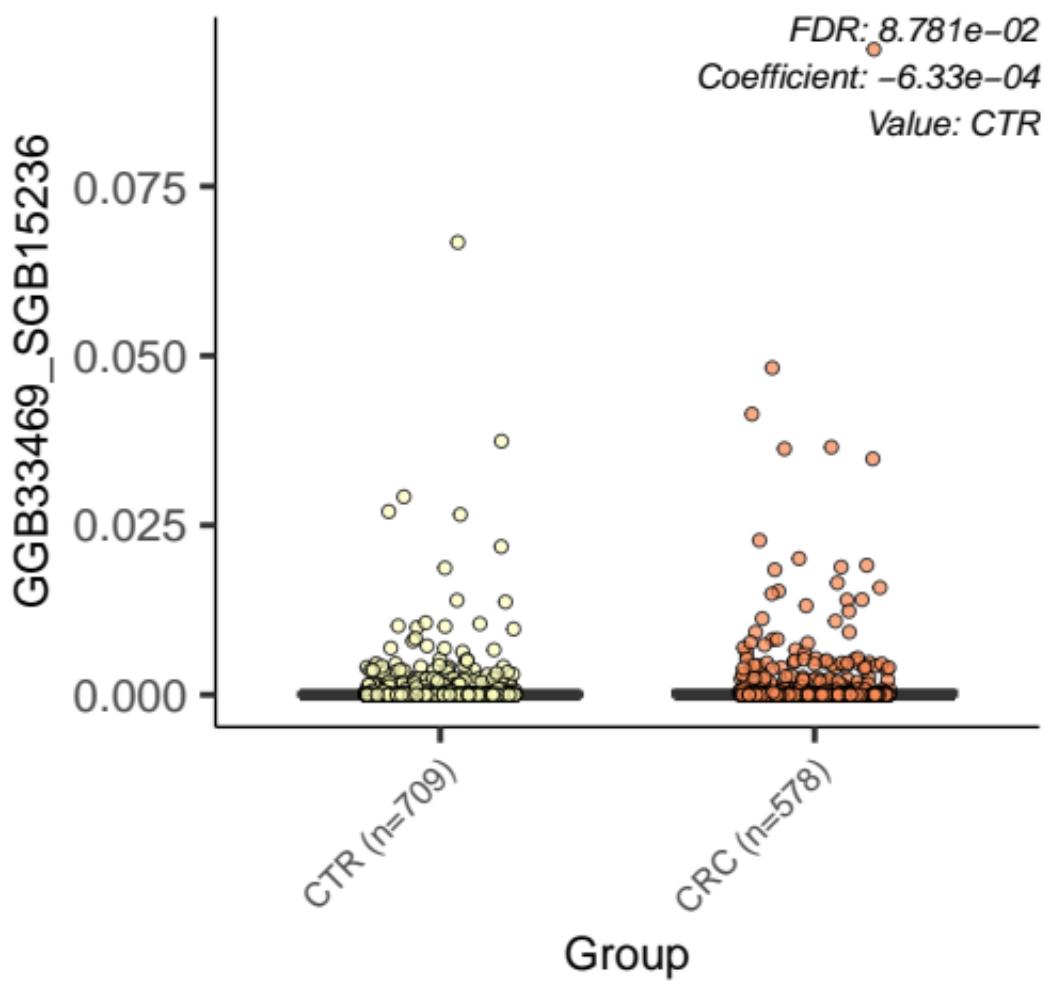
CRC (n=578)

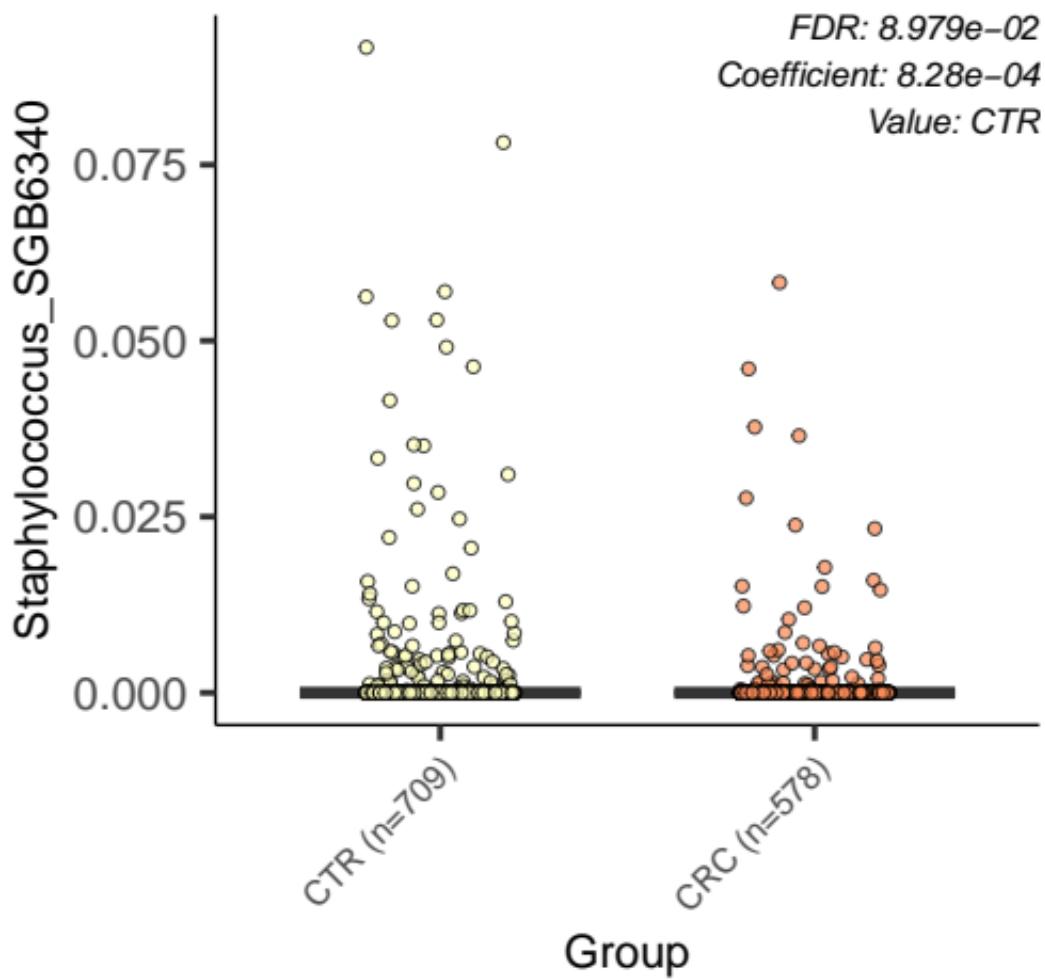
Group

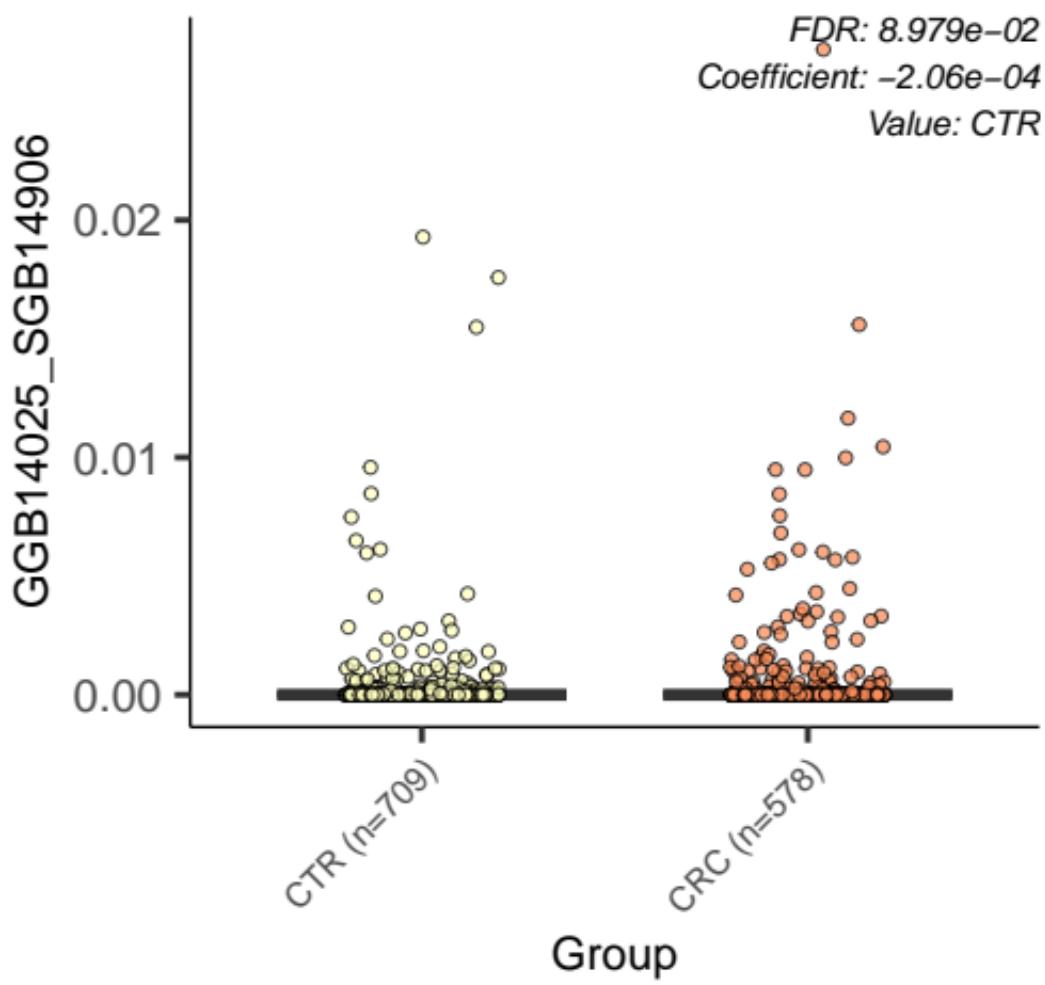












GGB4684_SGB6478

FDR: 9.176e-02
Coefficient: -1.90e-03
Value: CTR

0.2

0.1

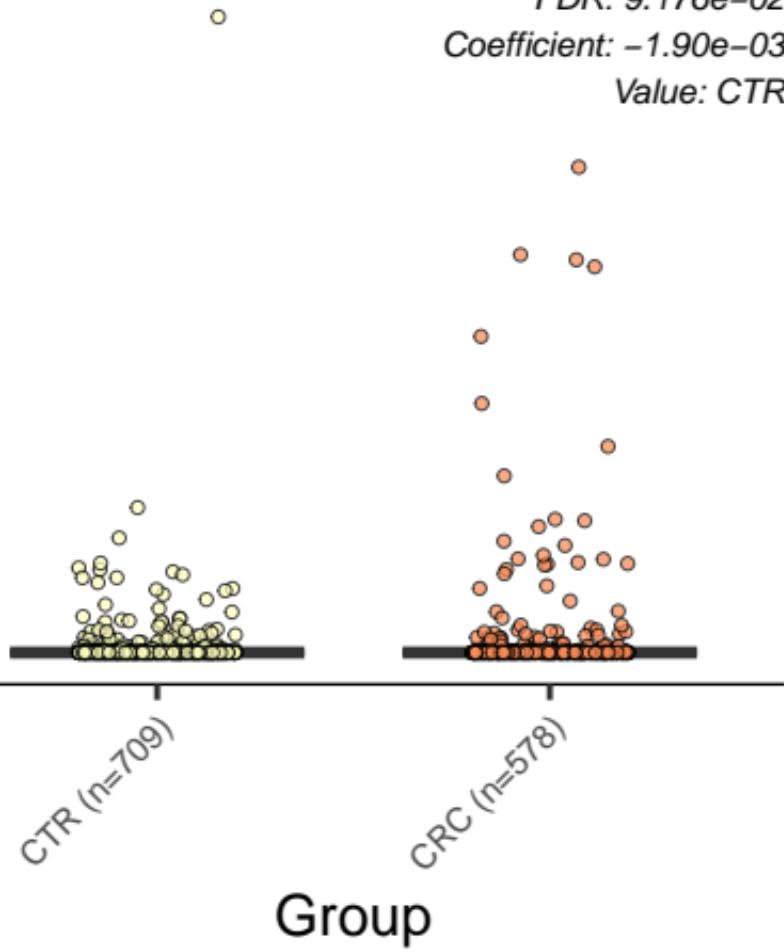
0.0

0.25

CTR (n=709)

CRC (n=578)

Group



GGB9712_SGB15244

FDR: 9.418e-02
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Value: CTR

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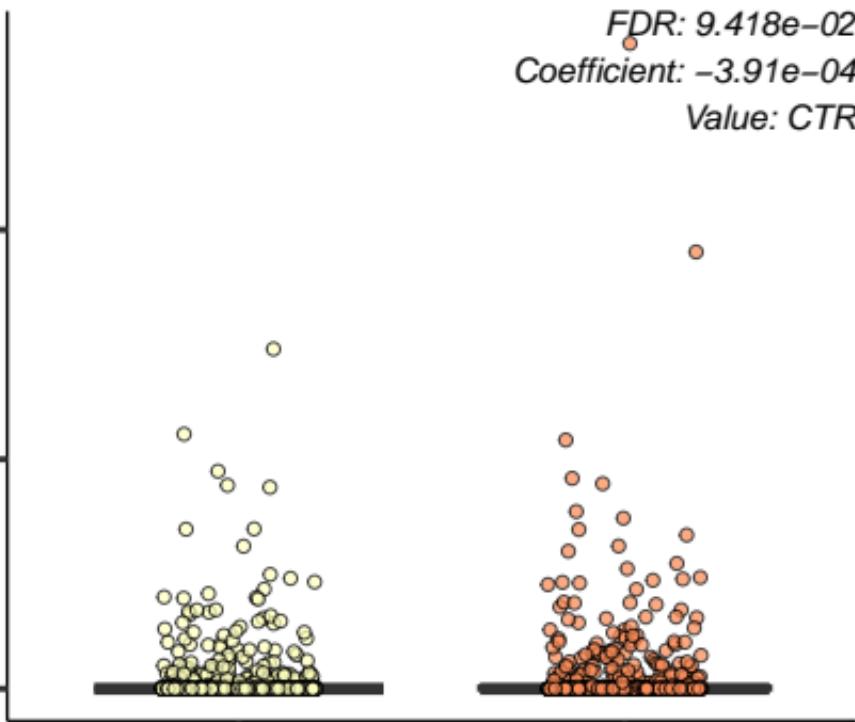
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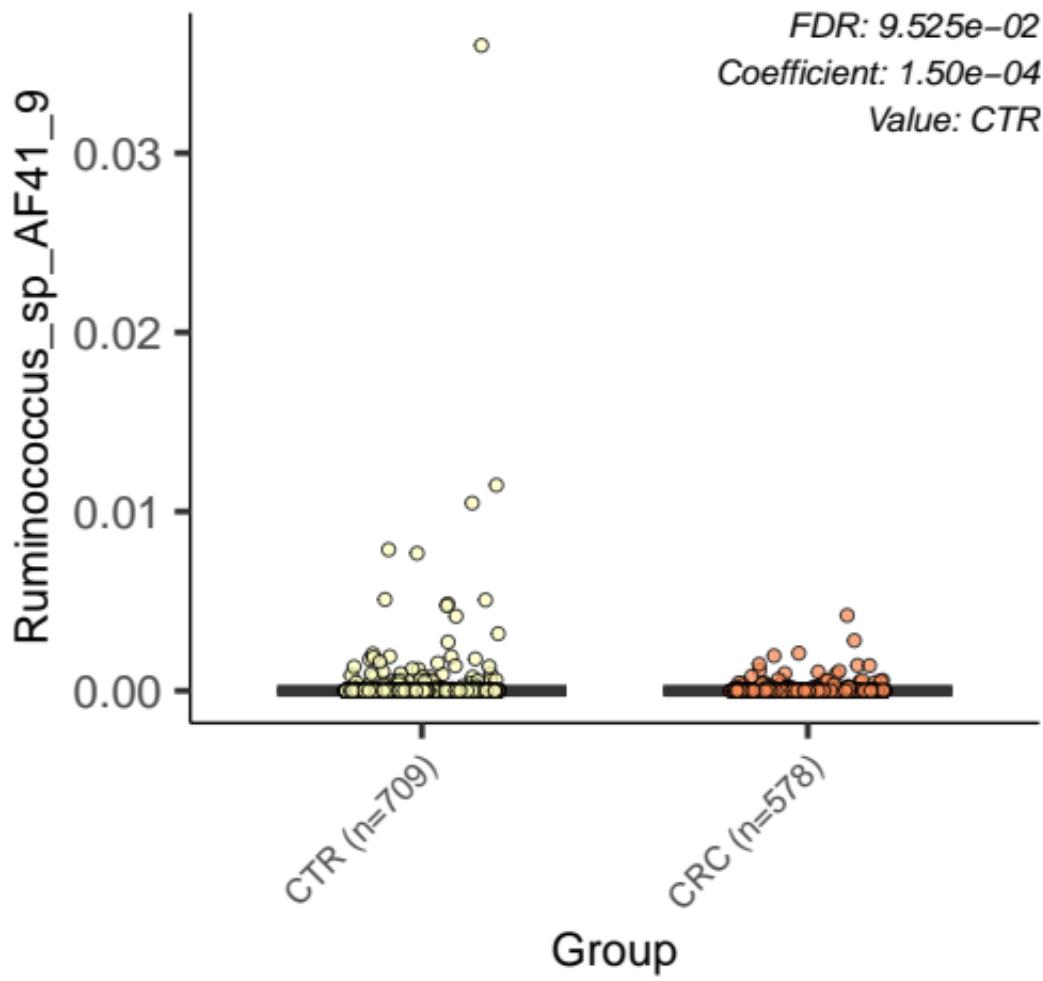
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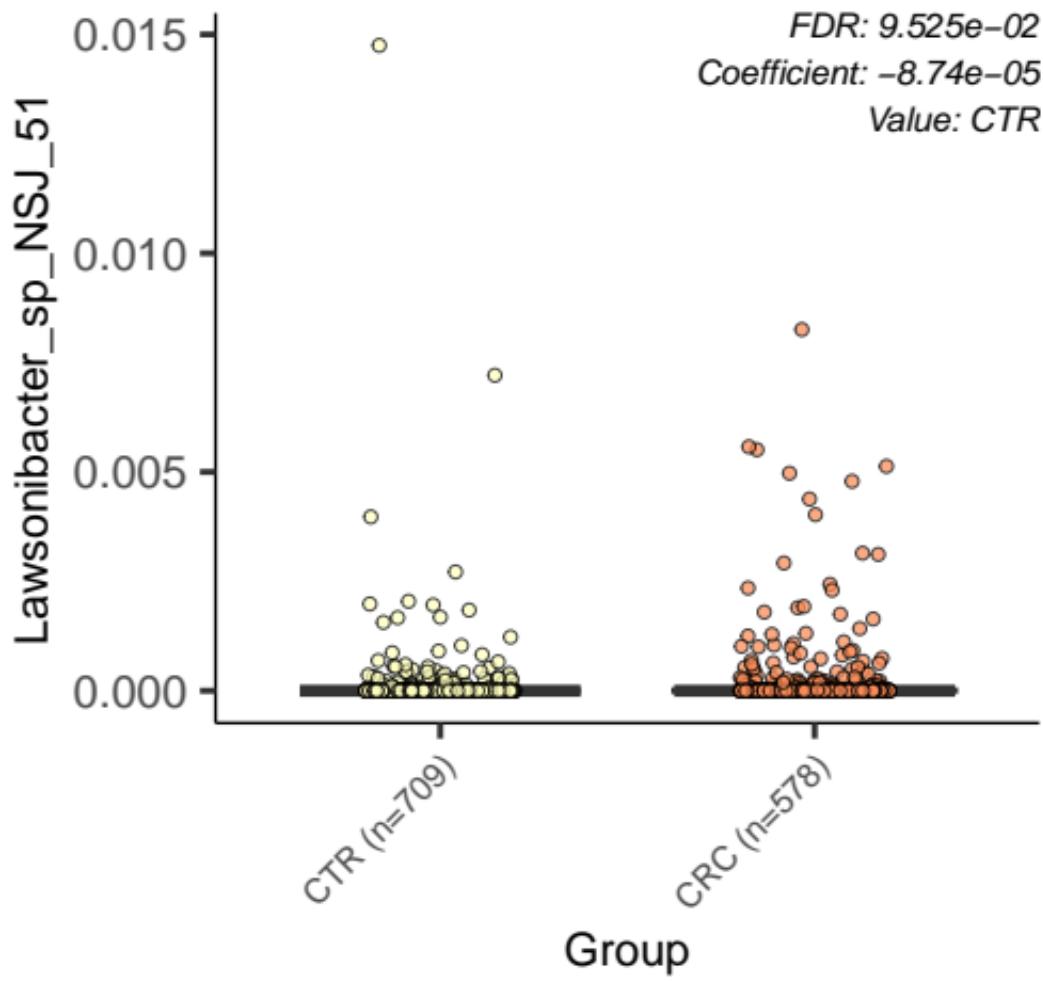
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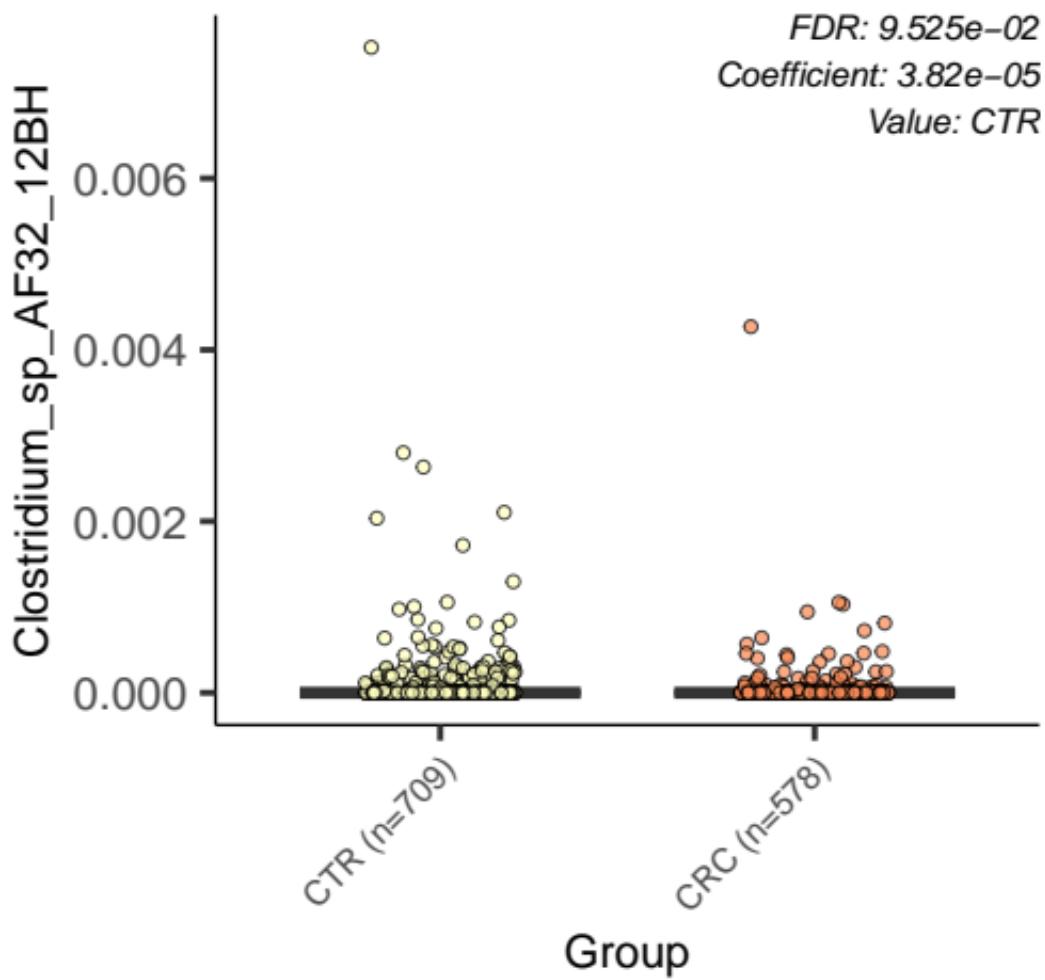
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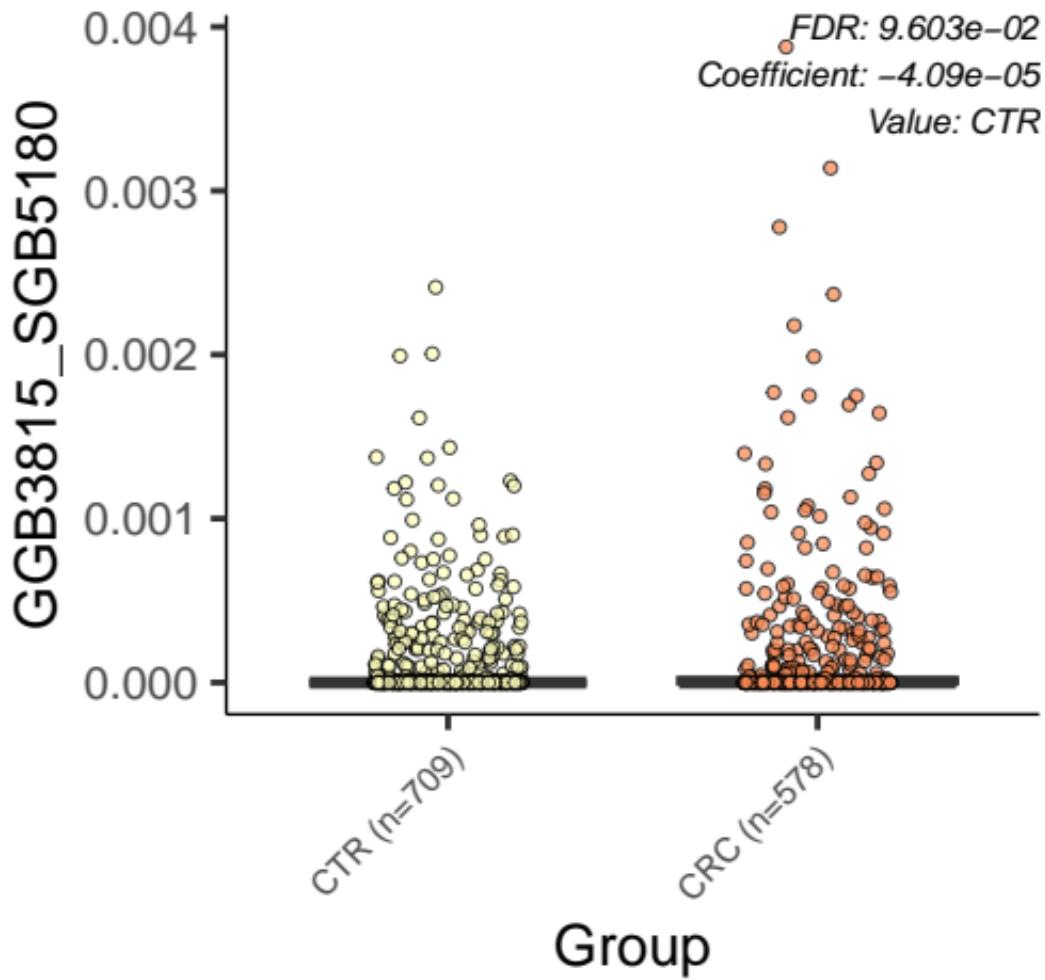
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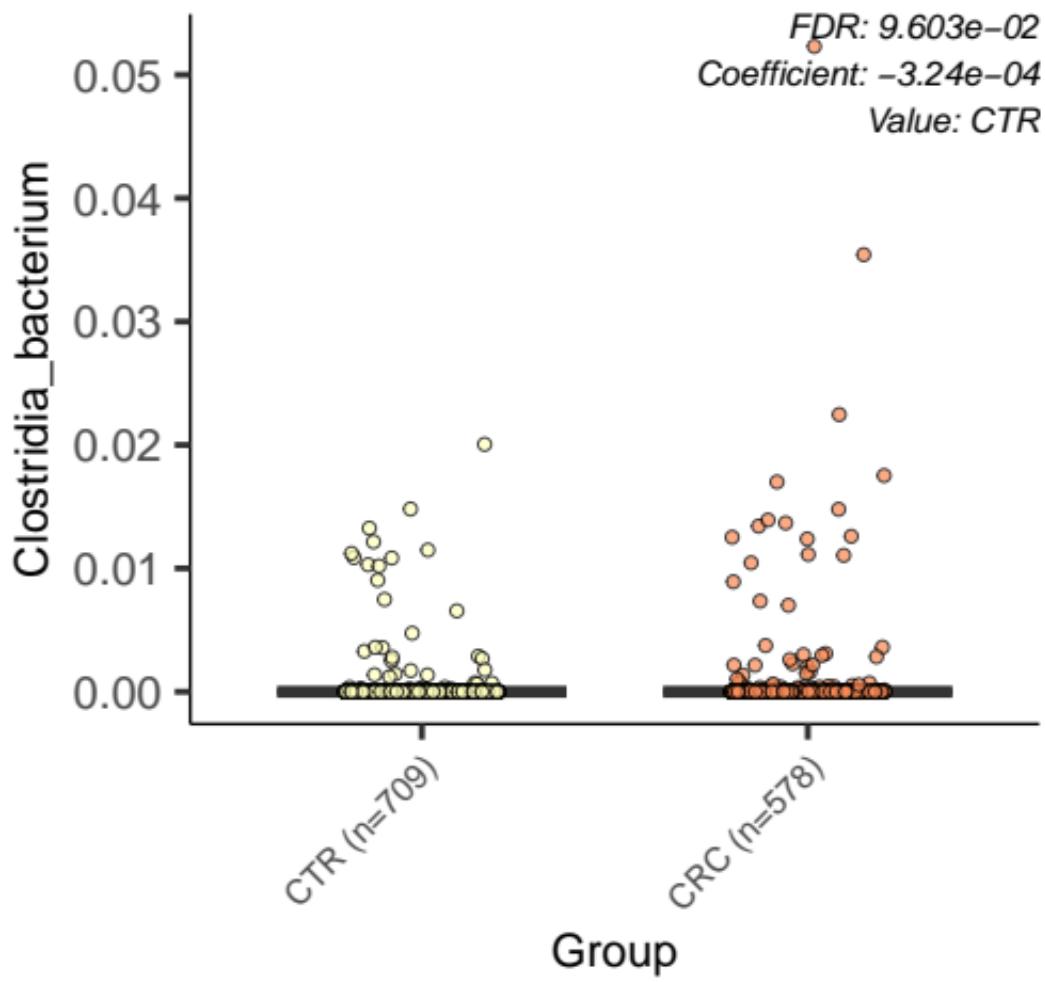


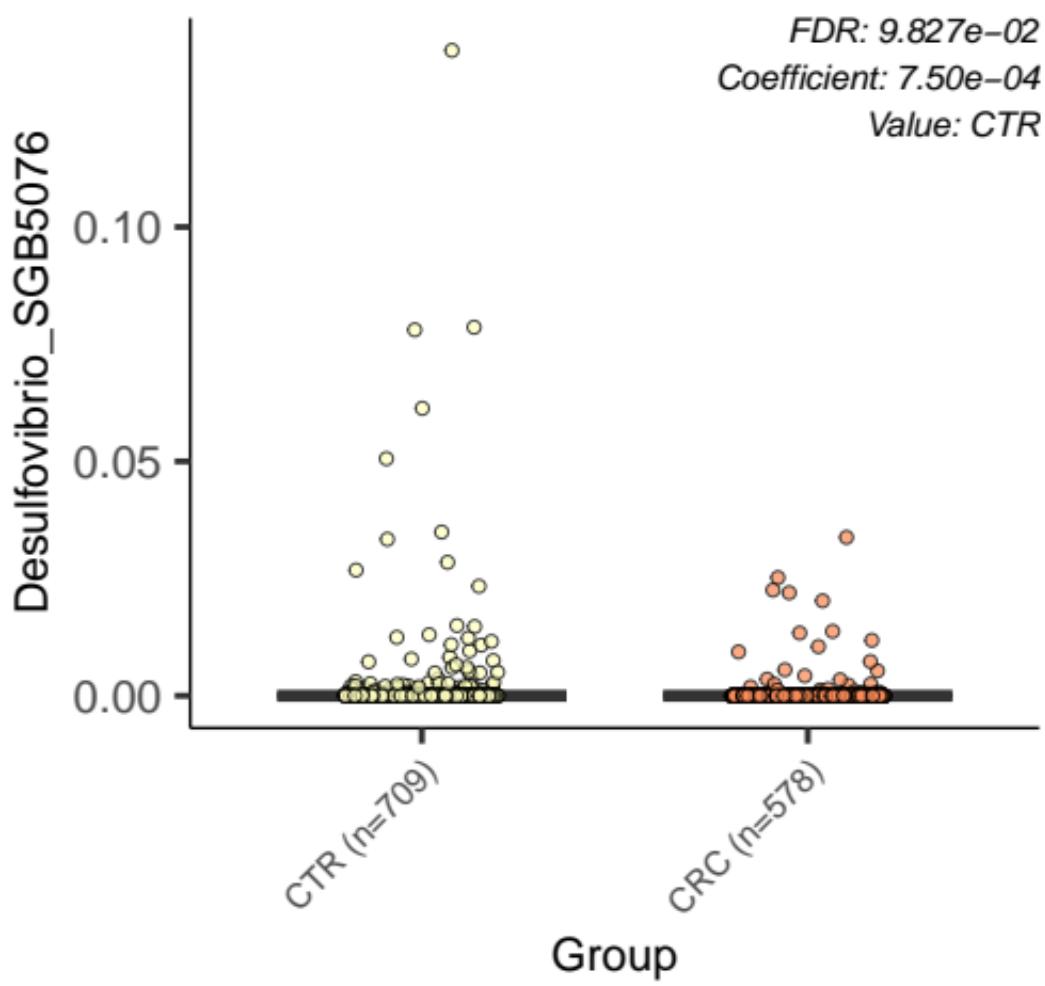












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Value: CTR

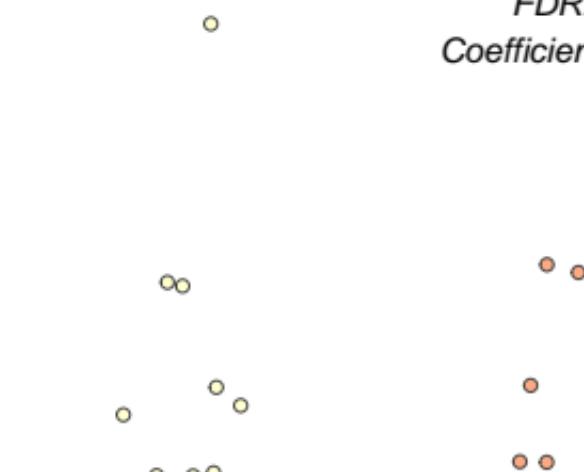
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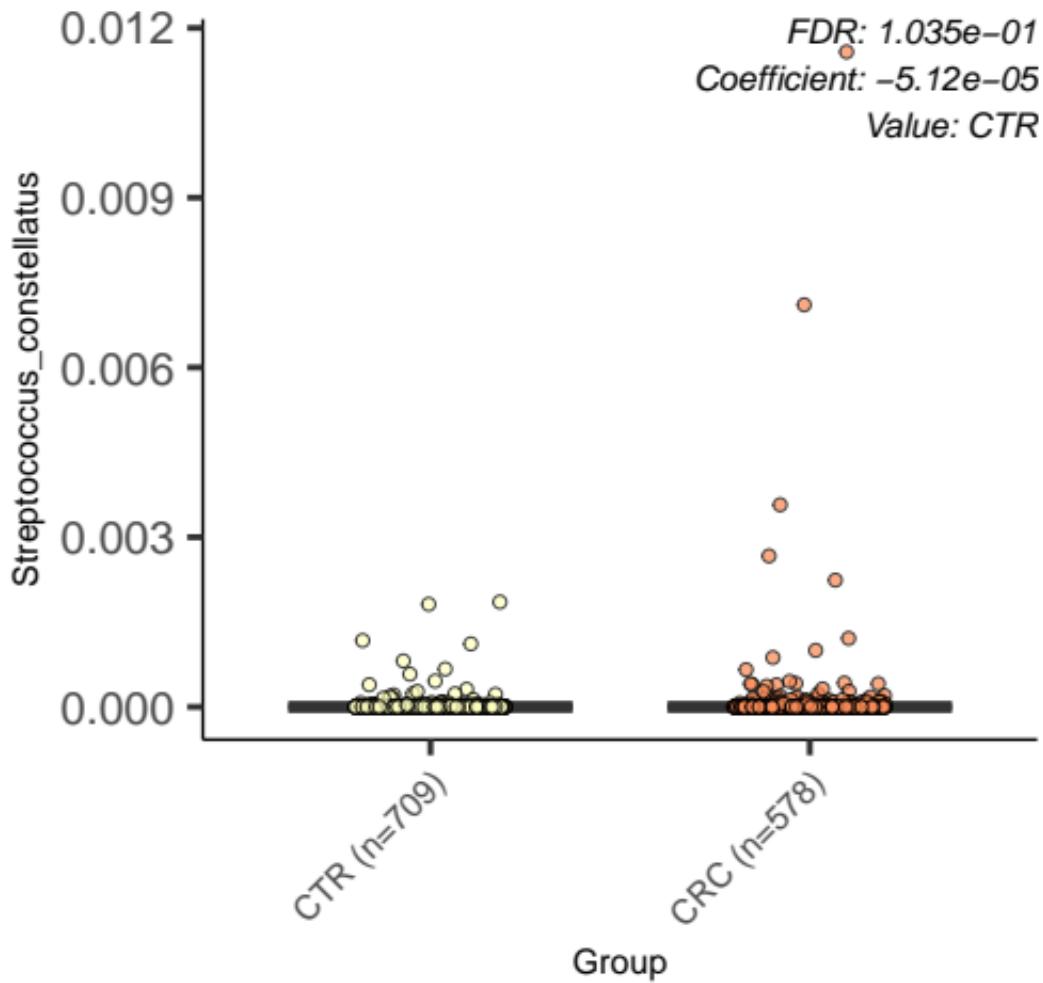
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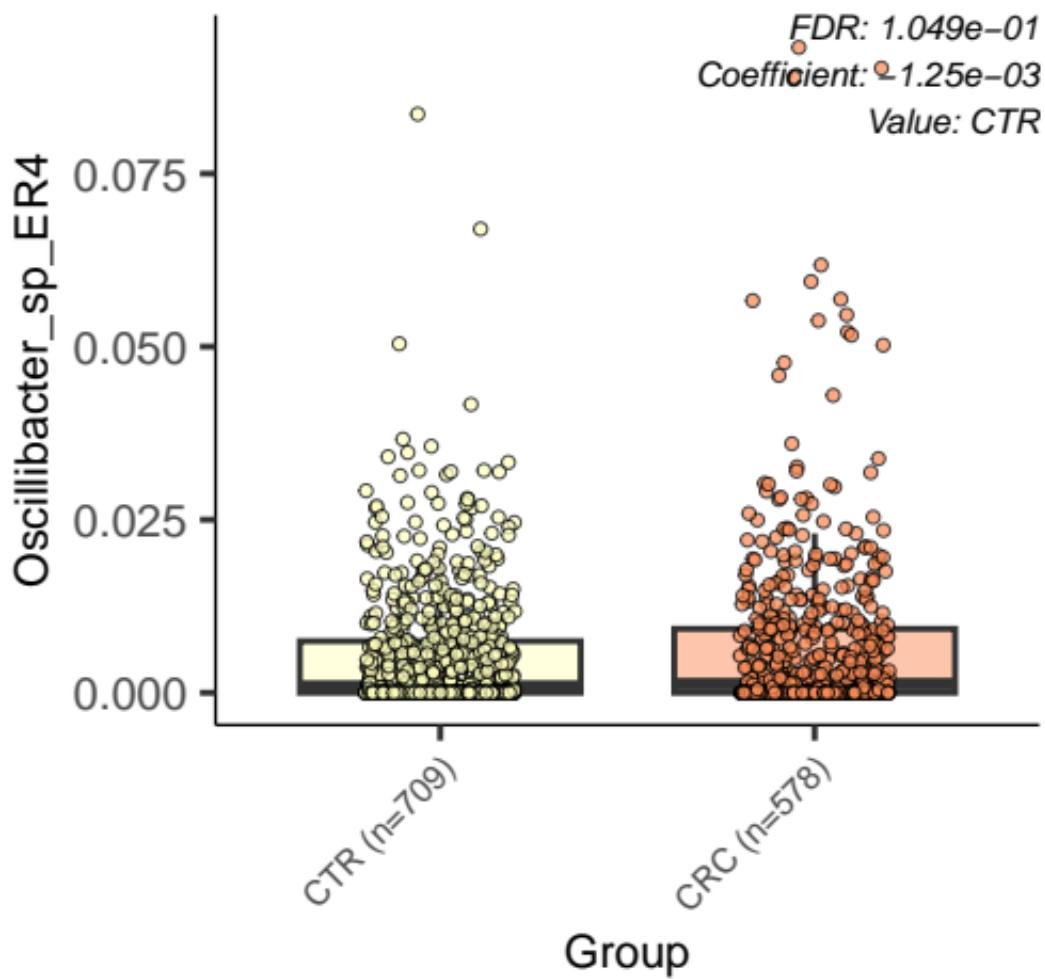
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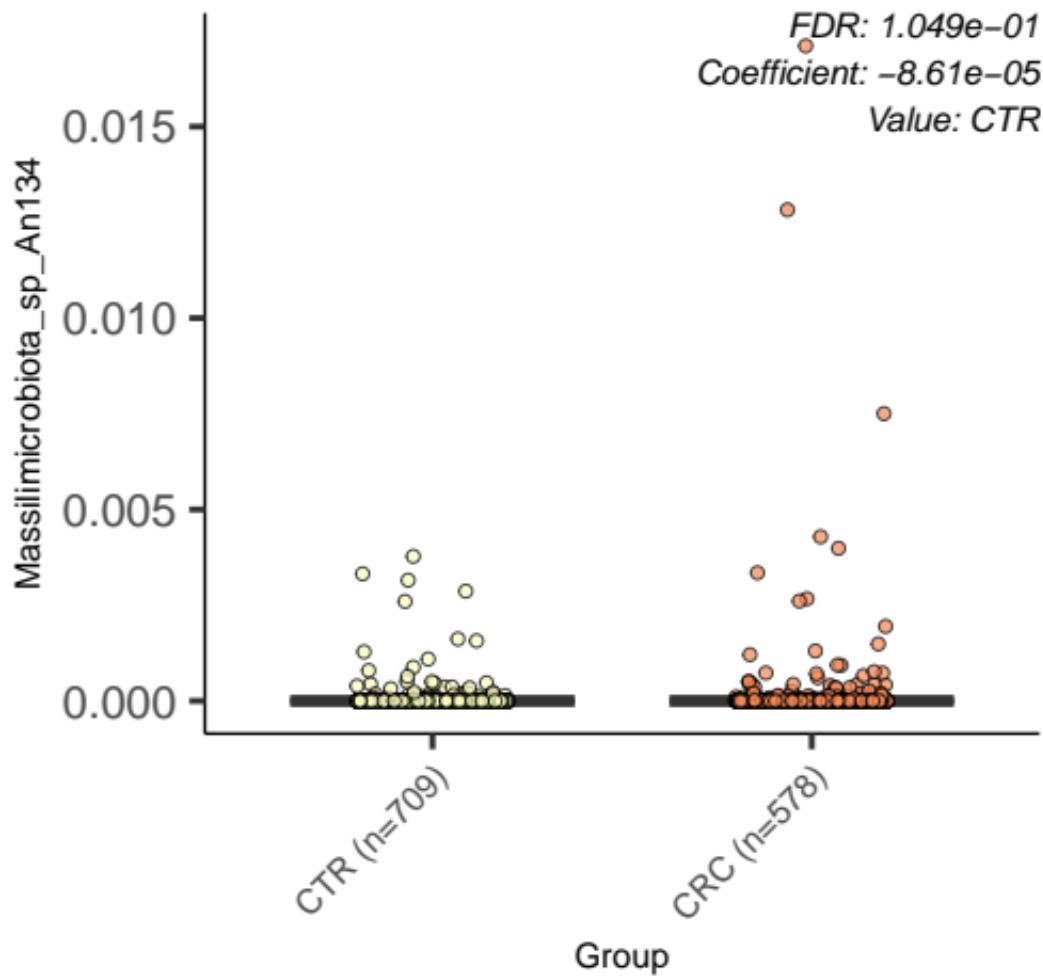
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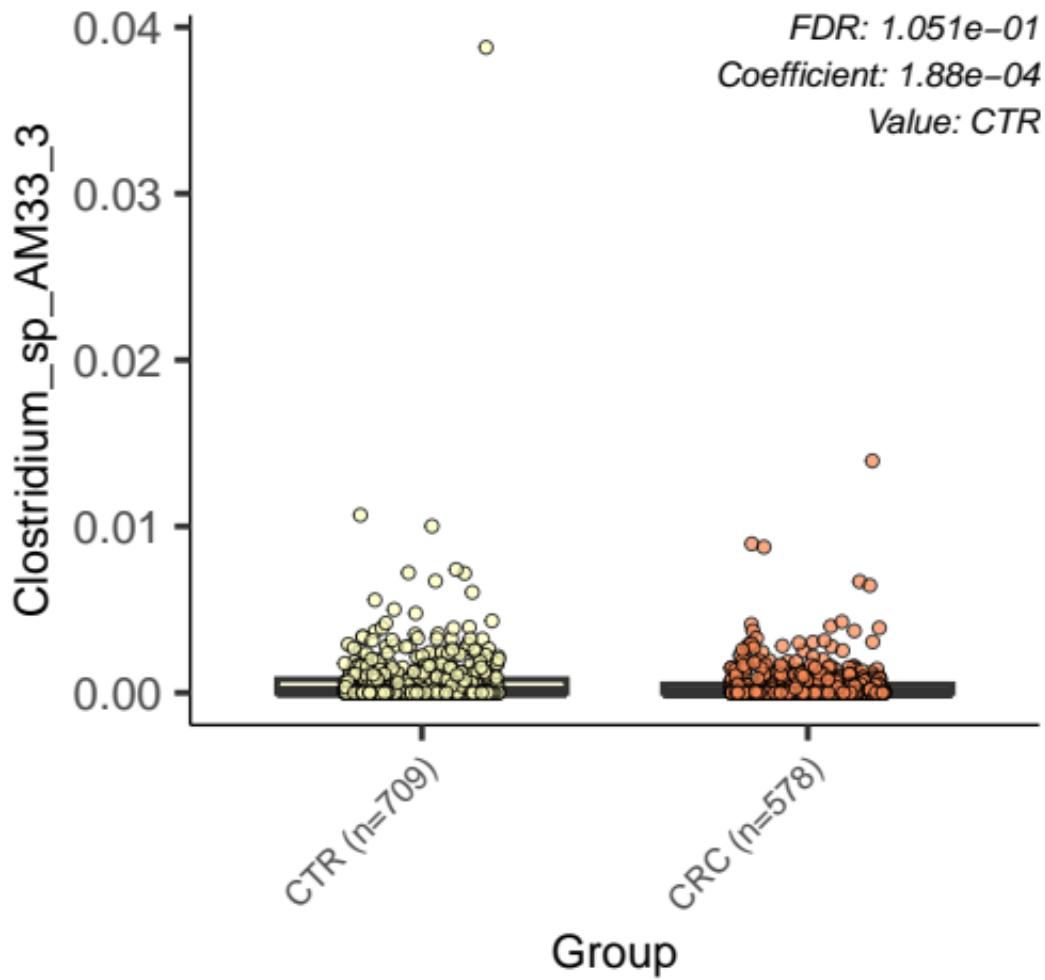
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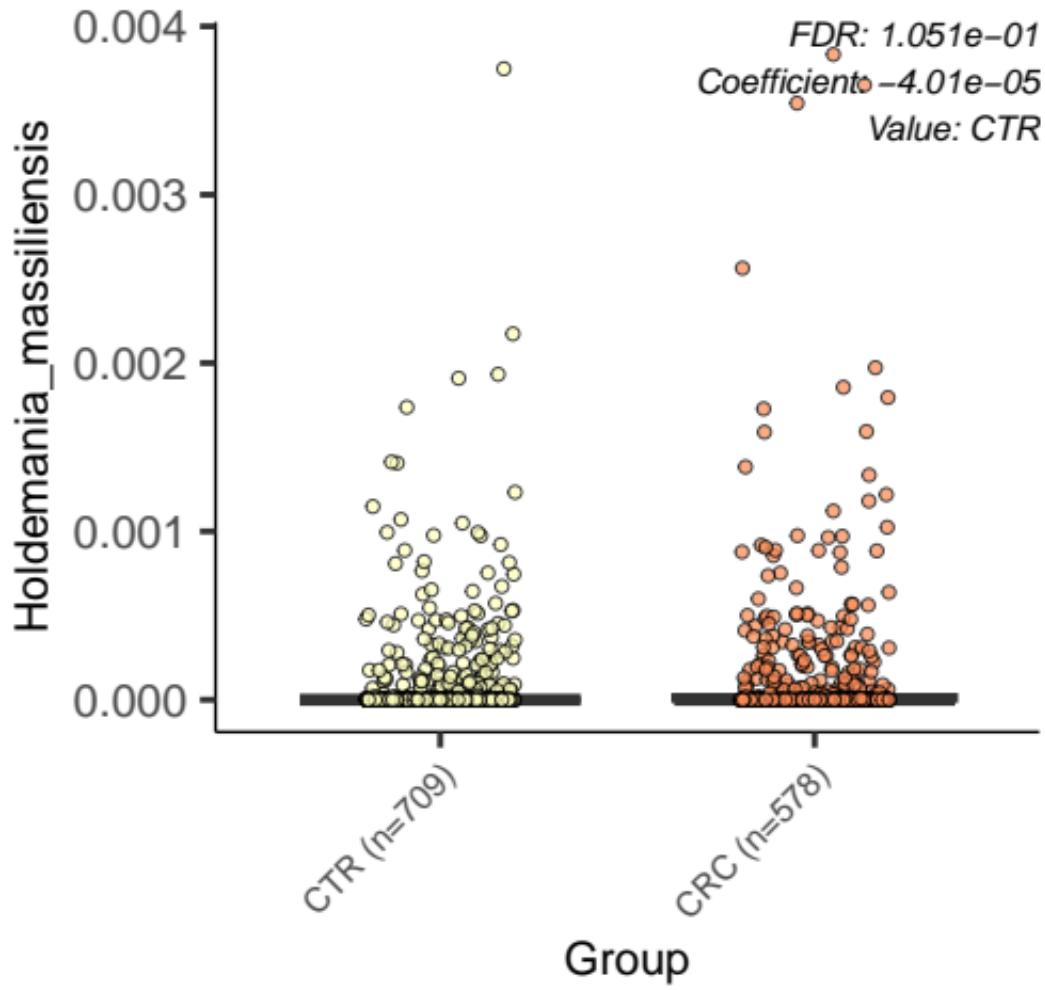


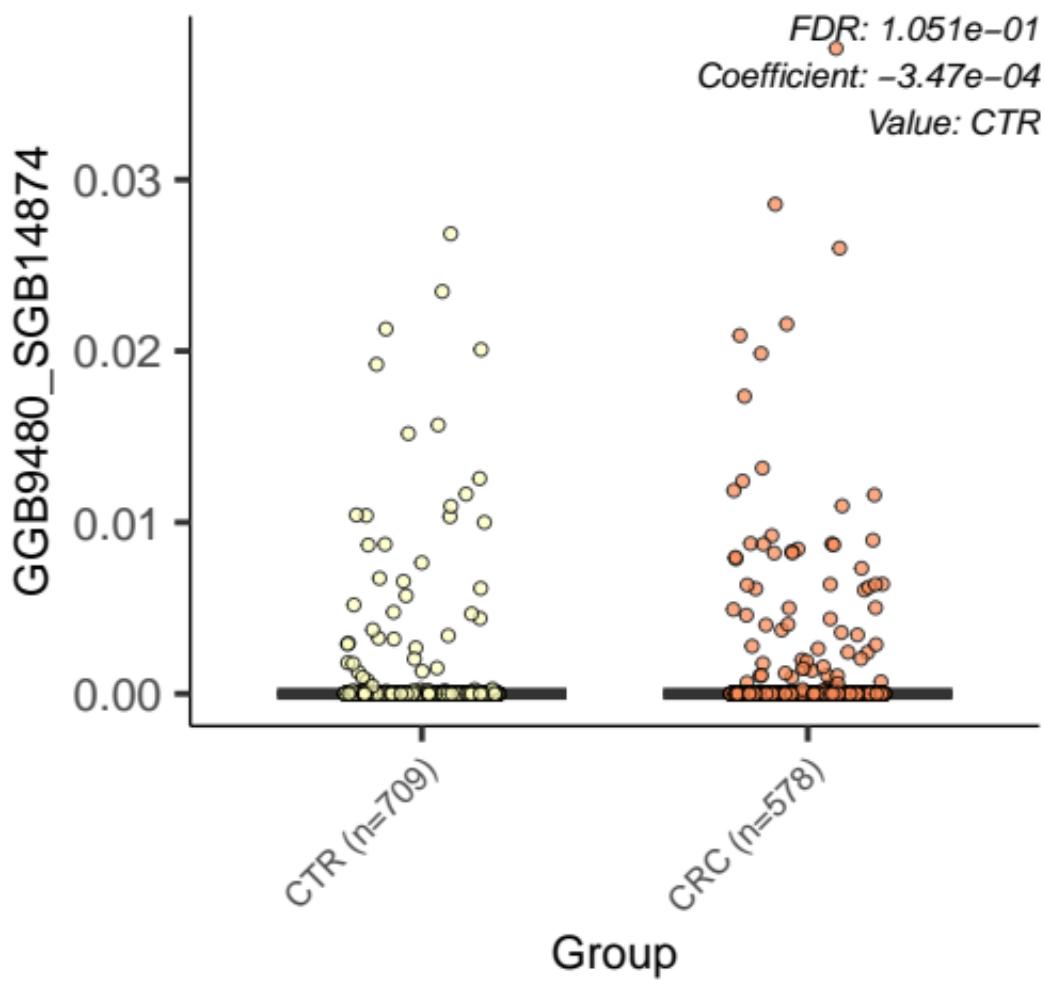


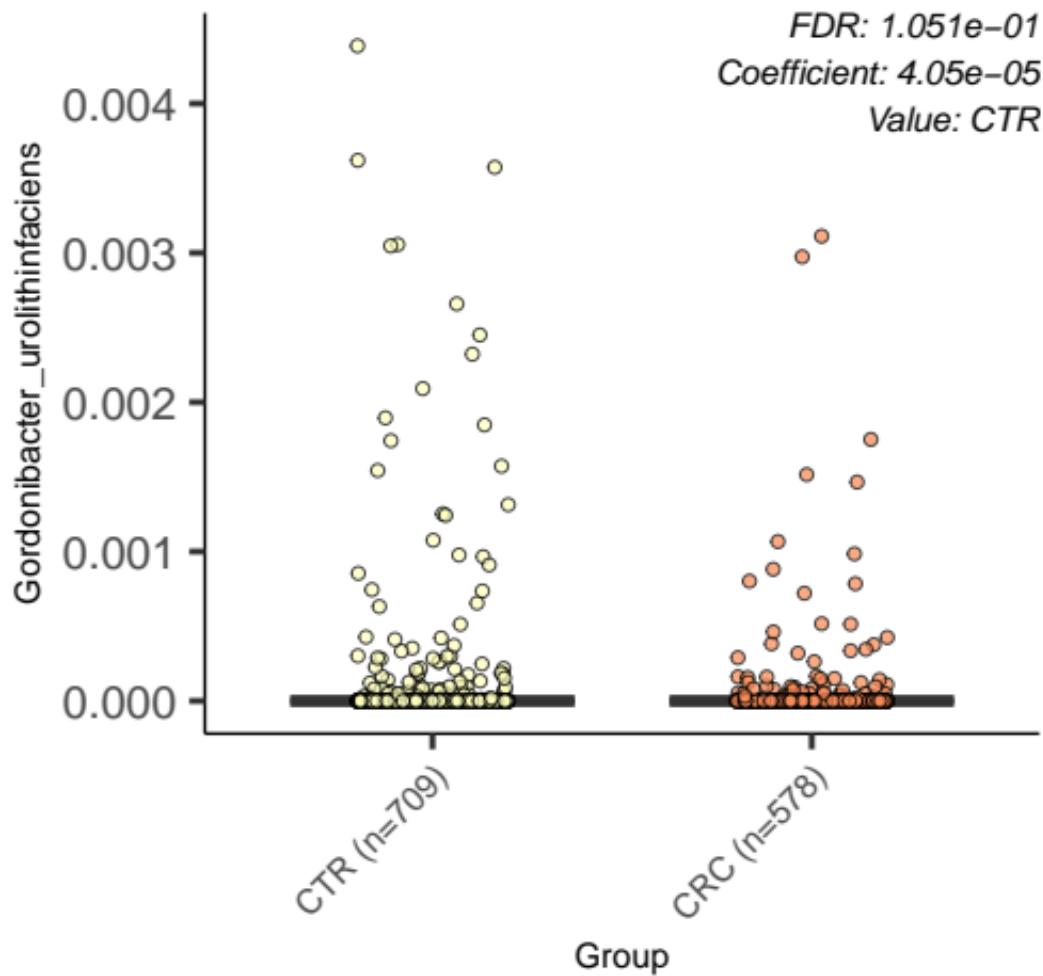


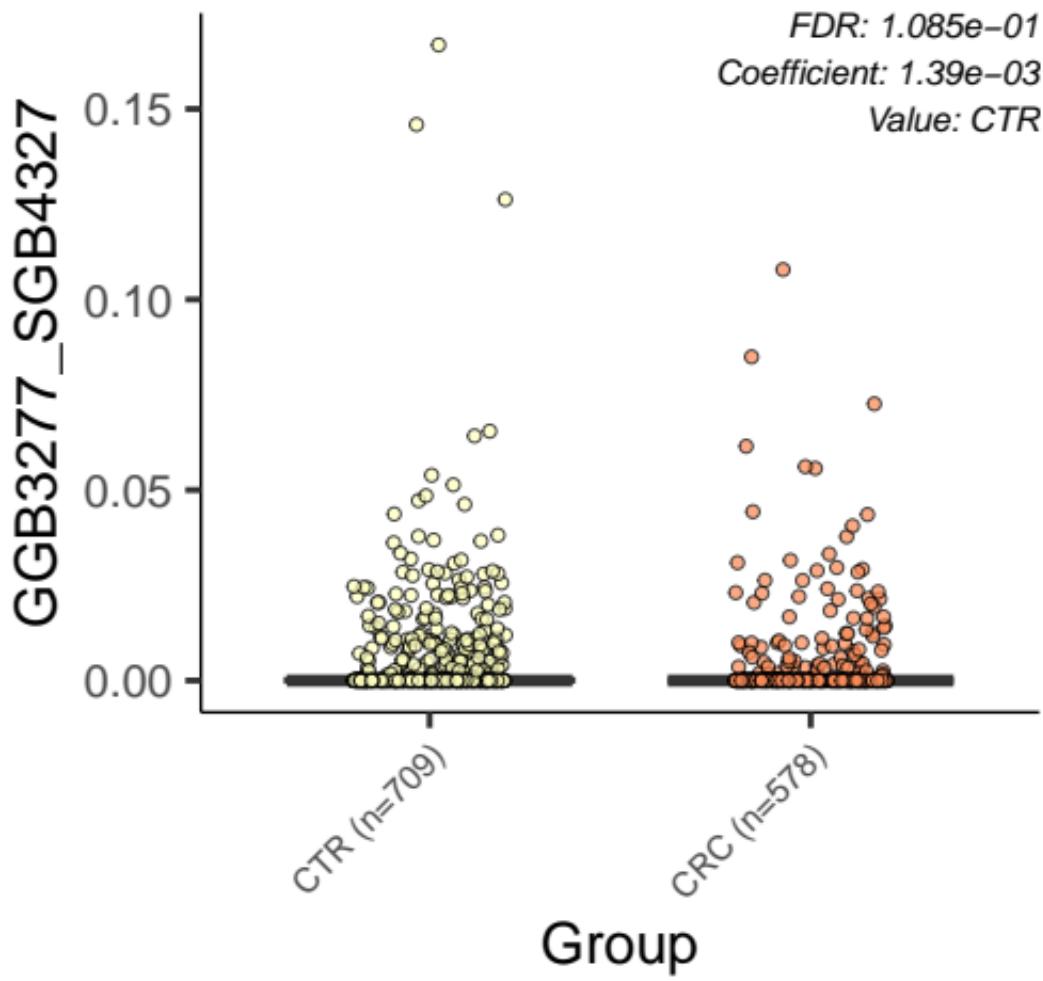


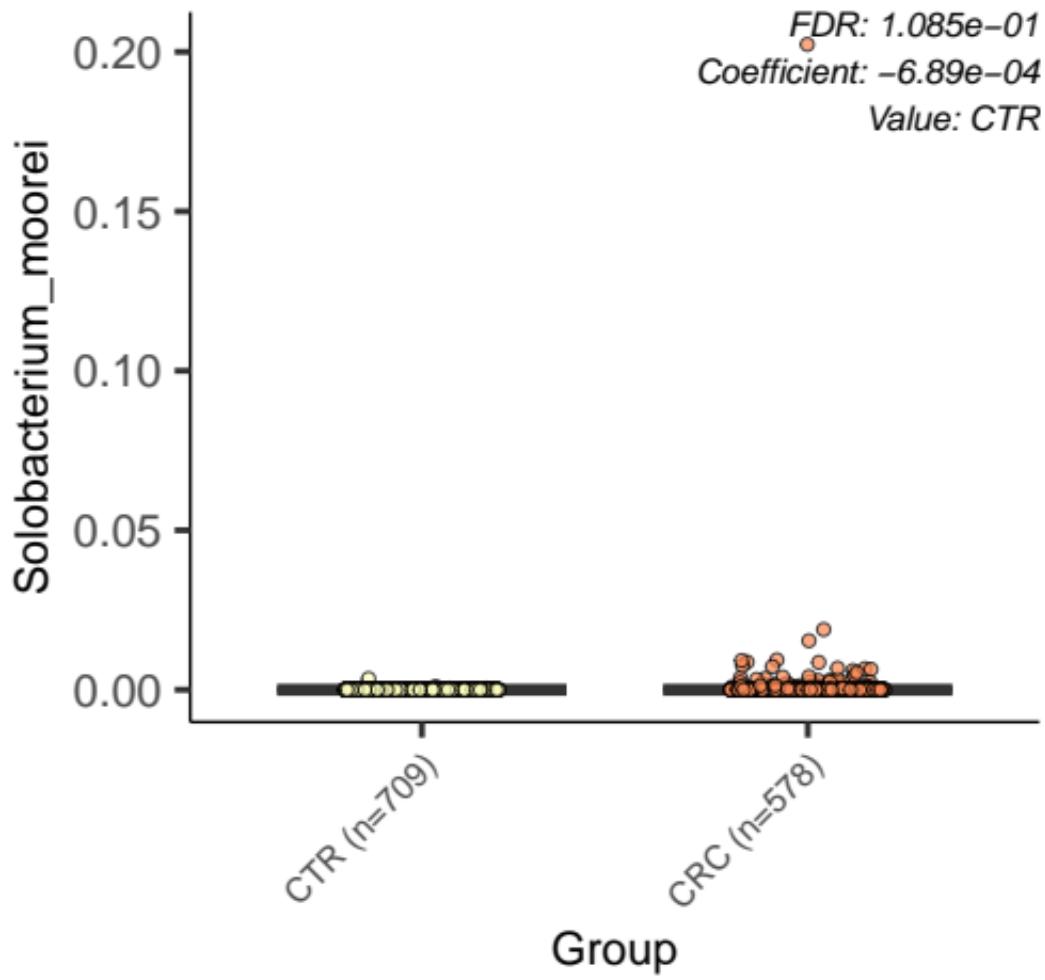


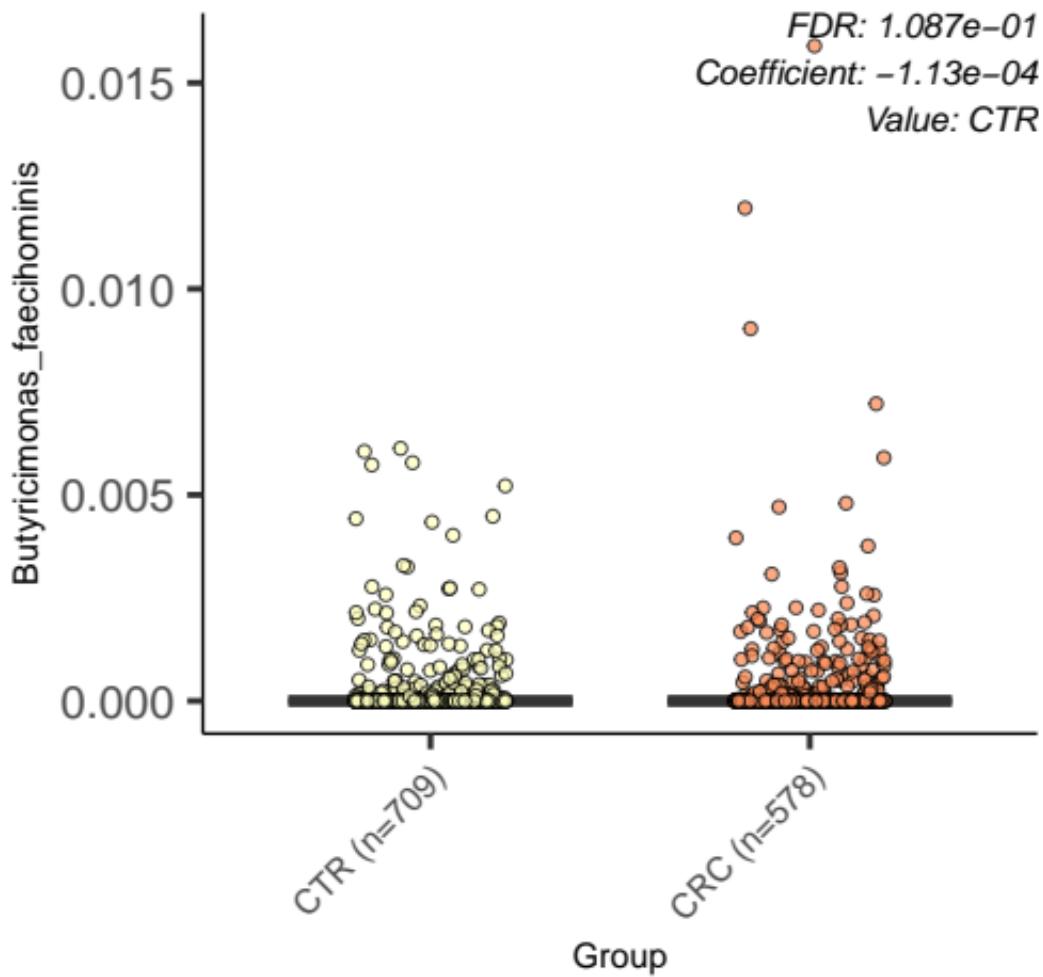


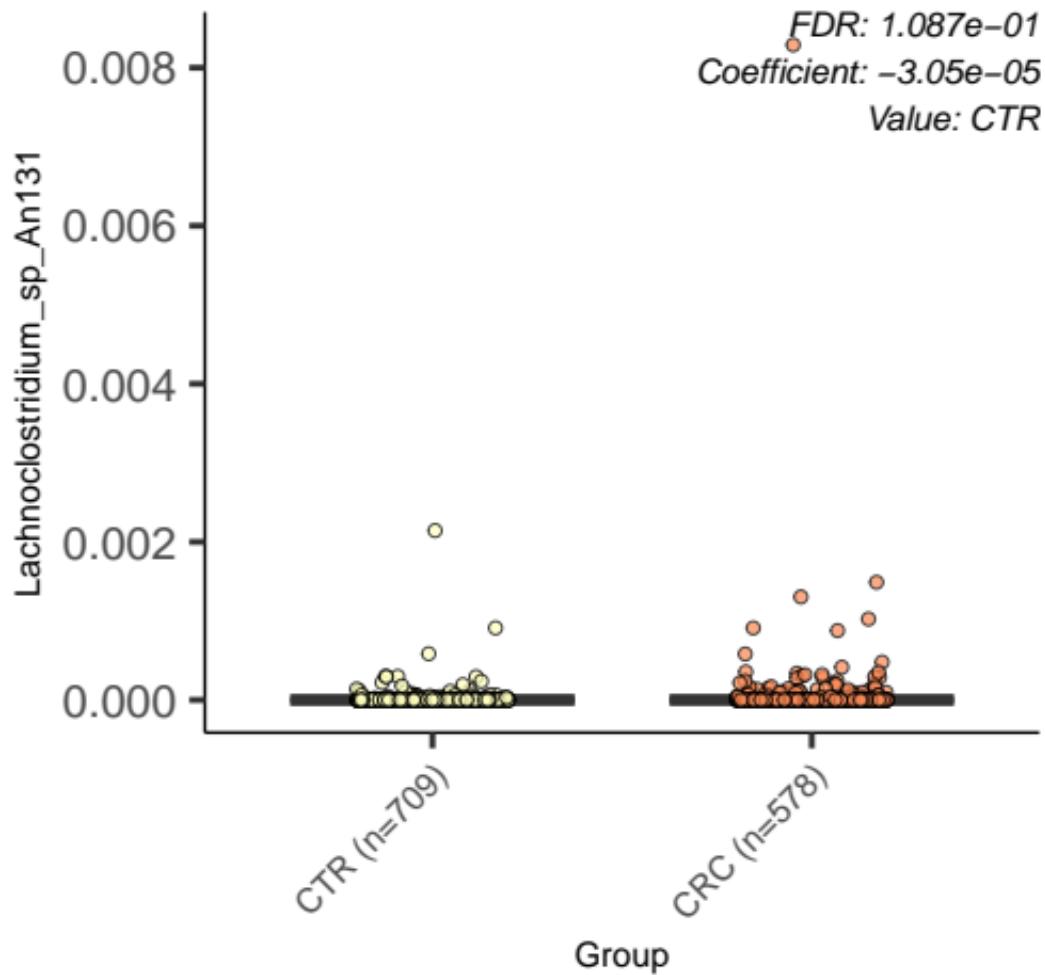


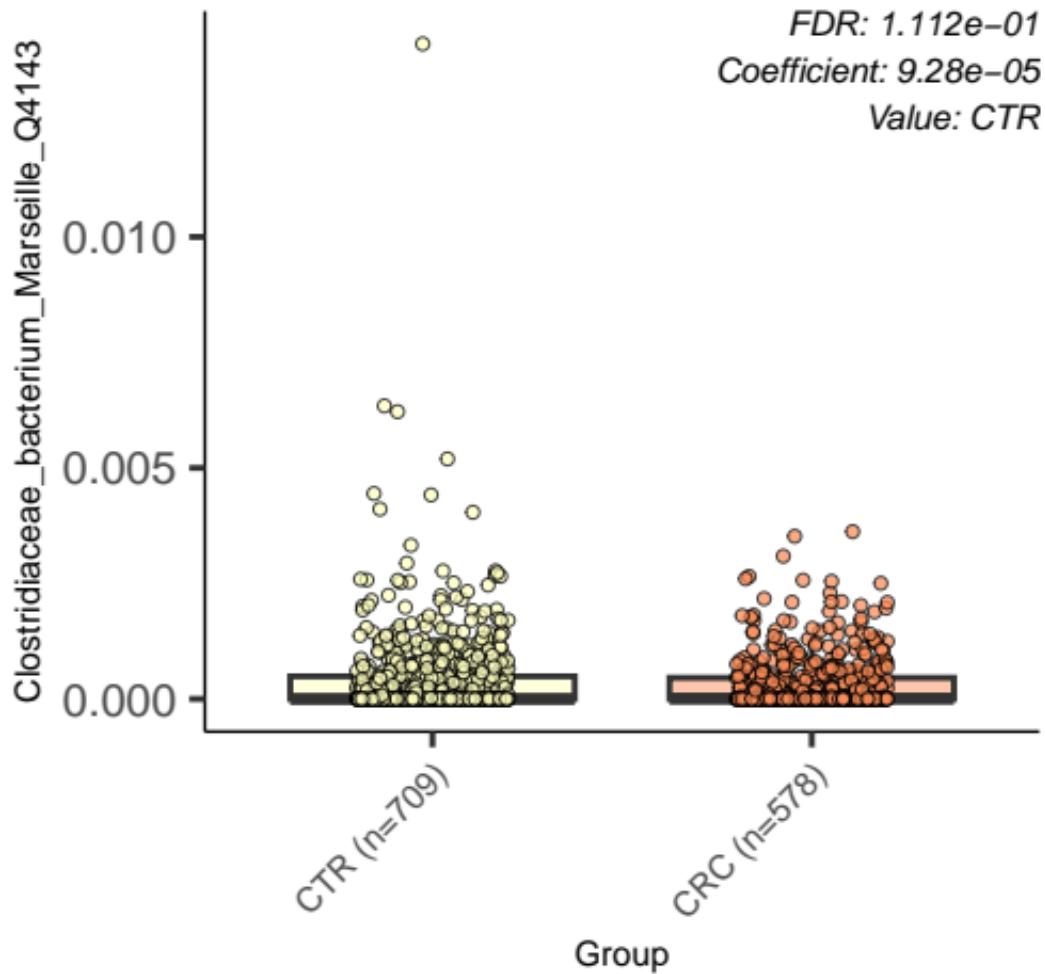


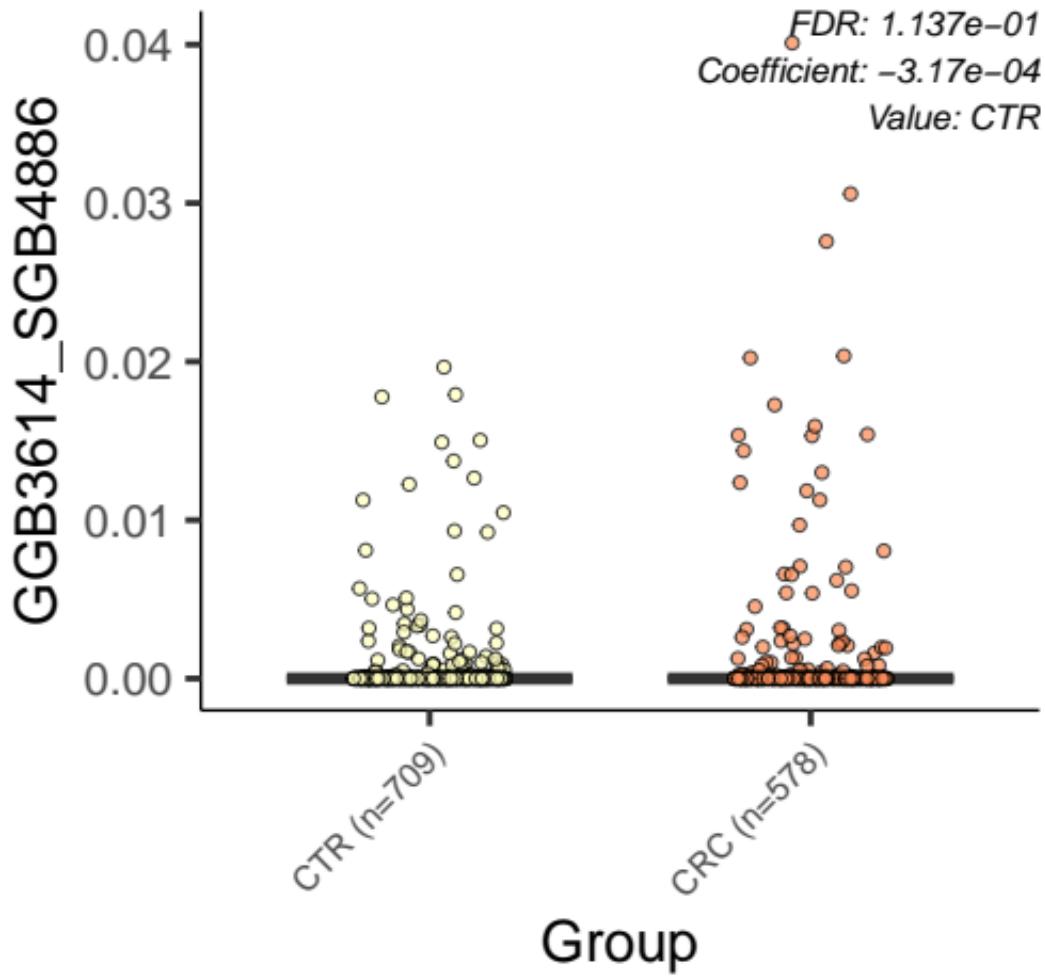


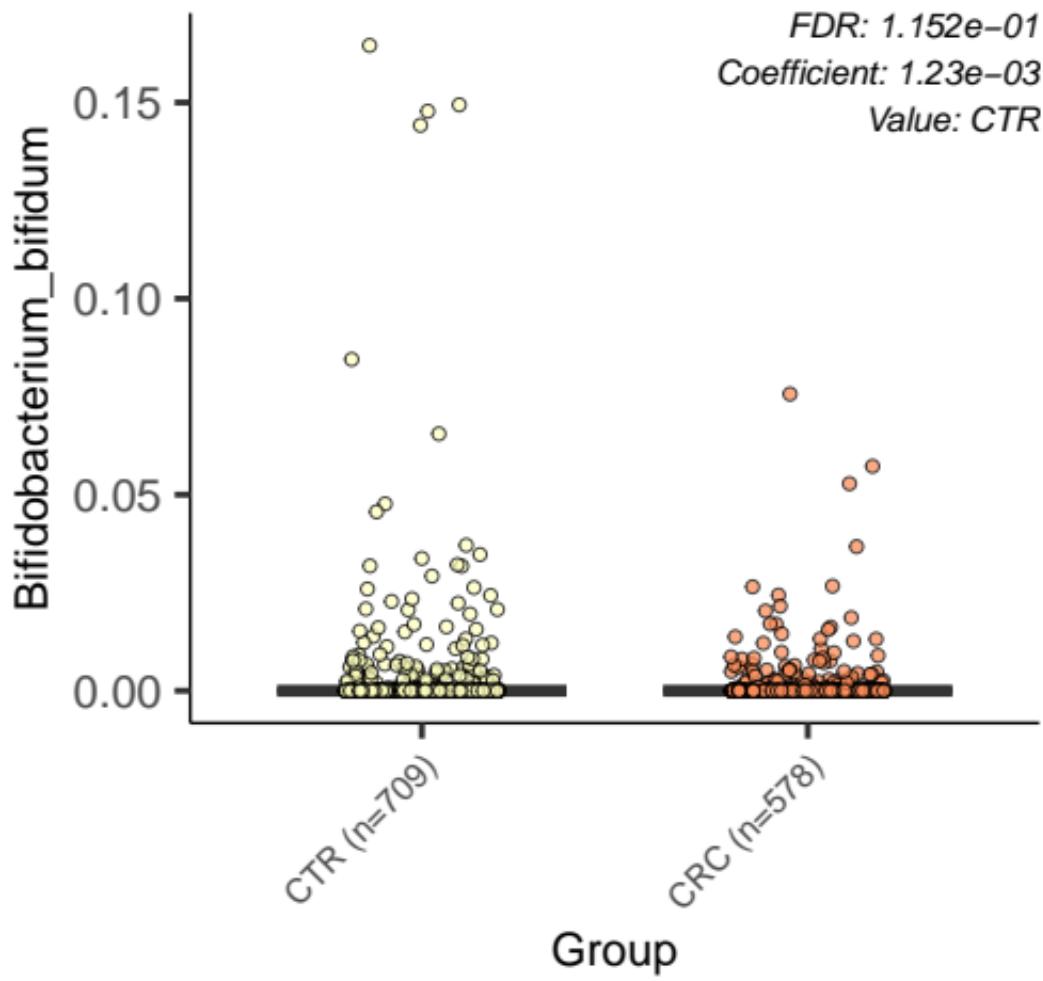


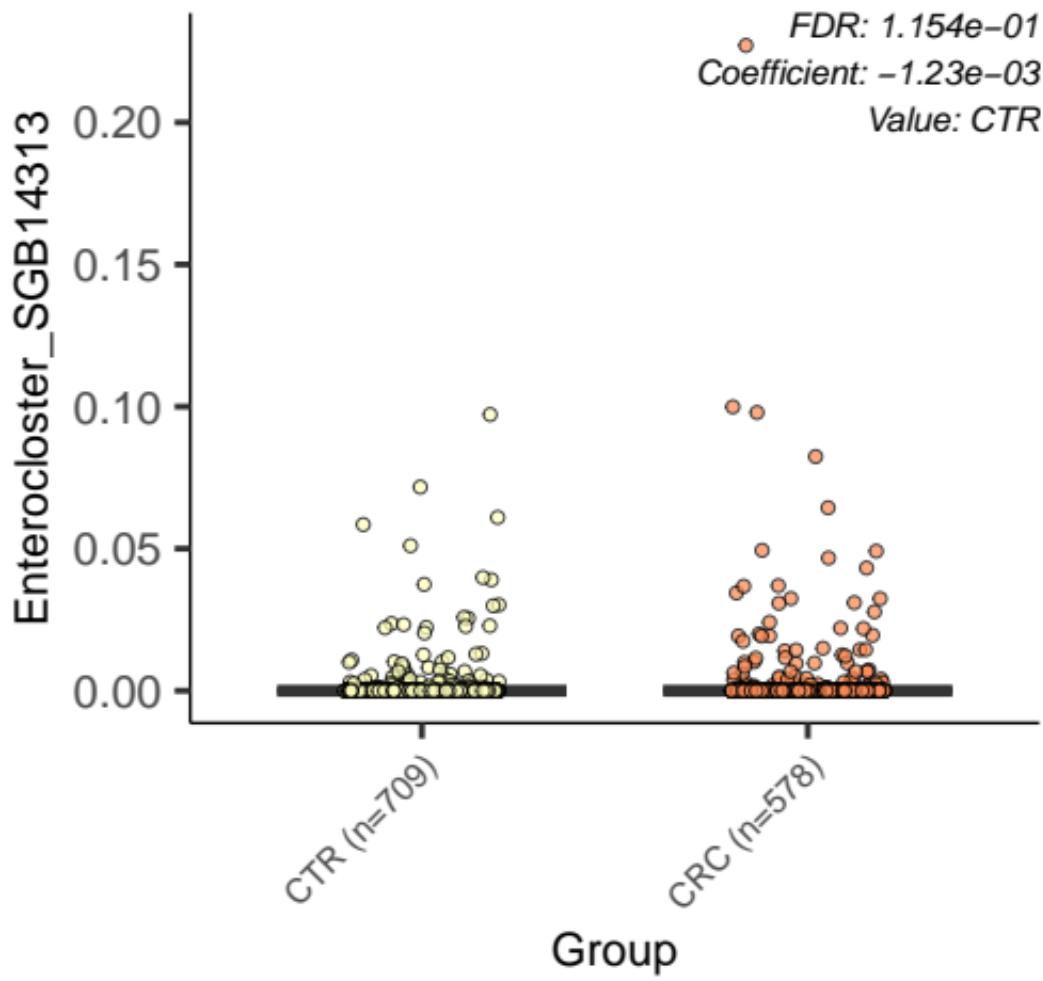


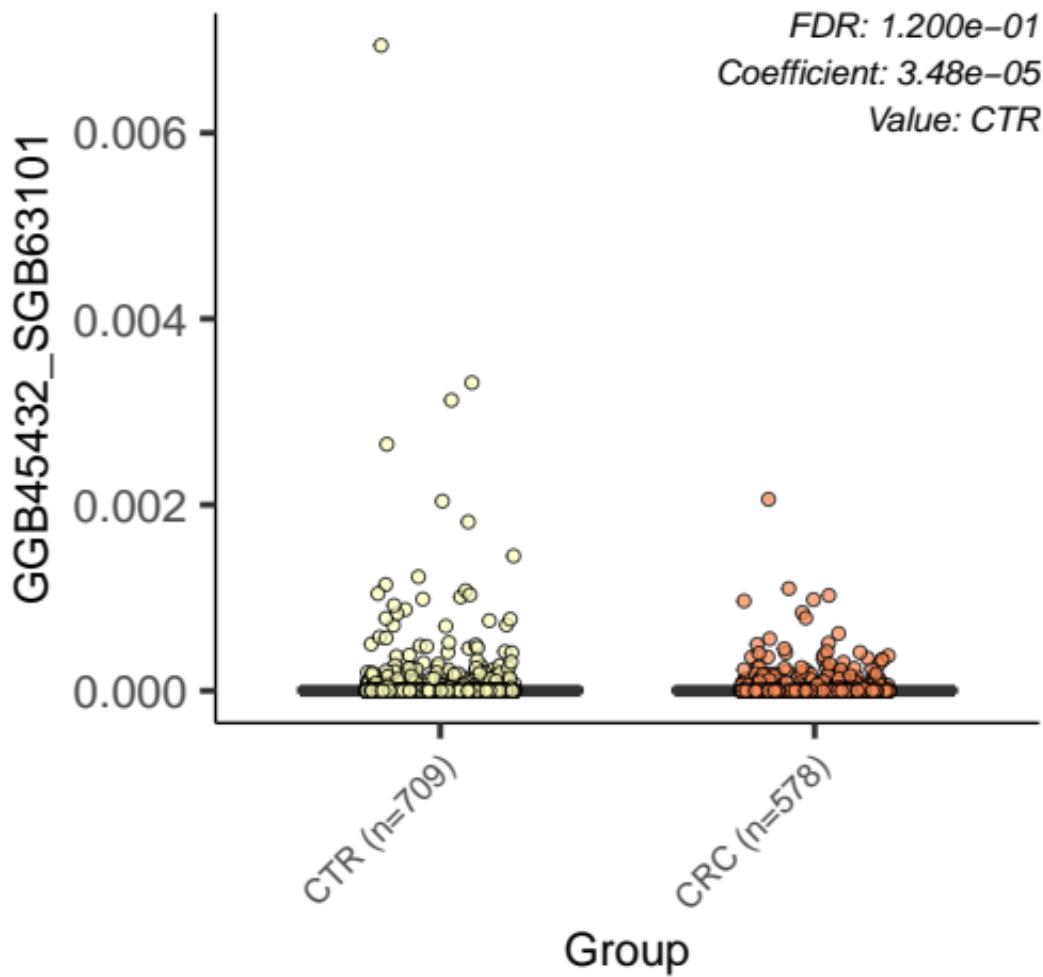


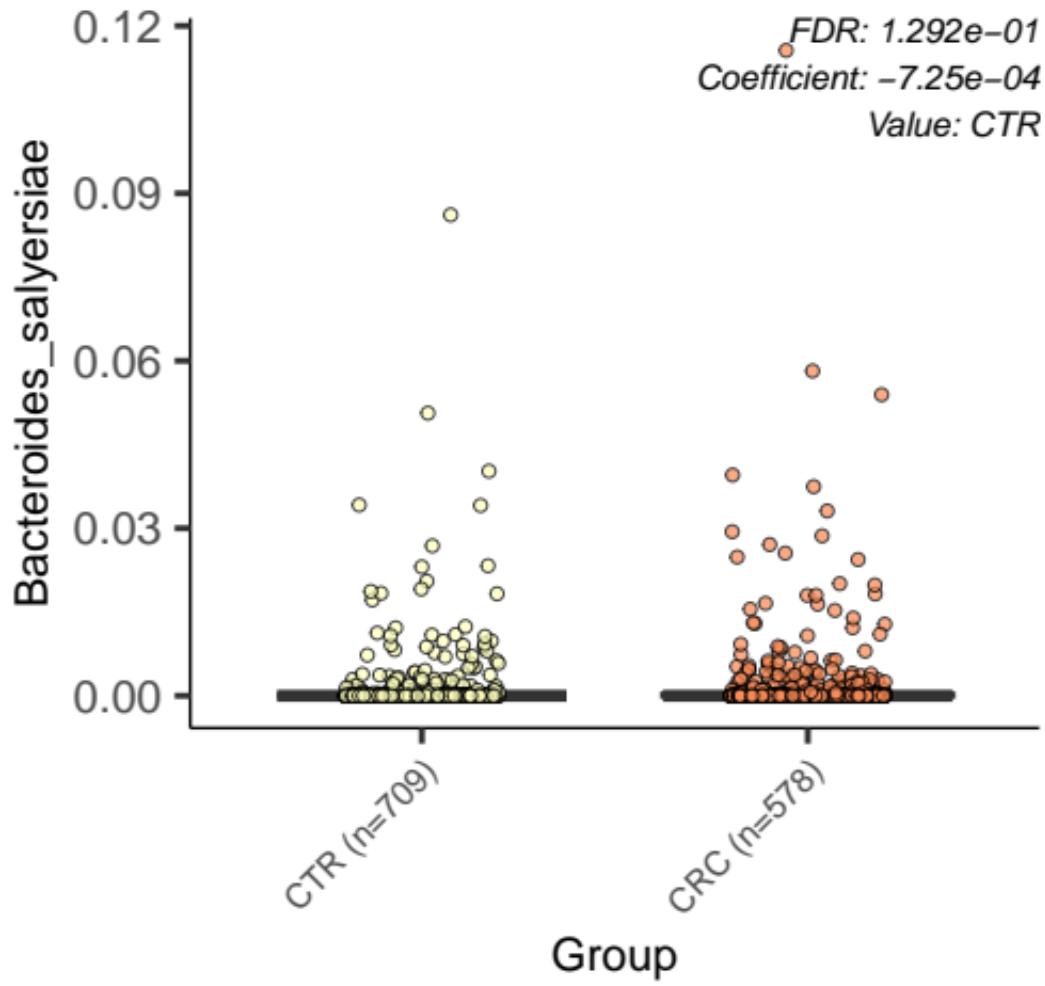


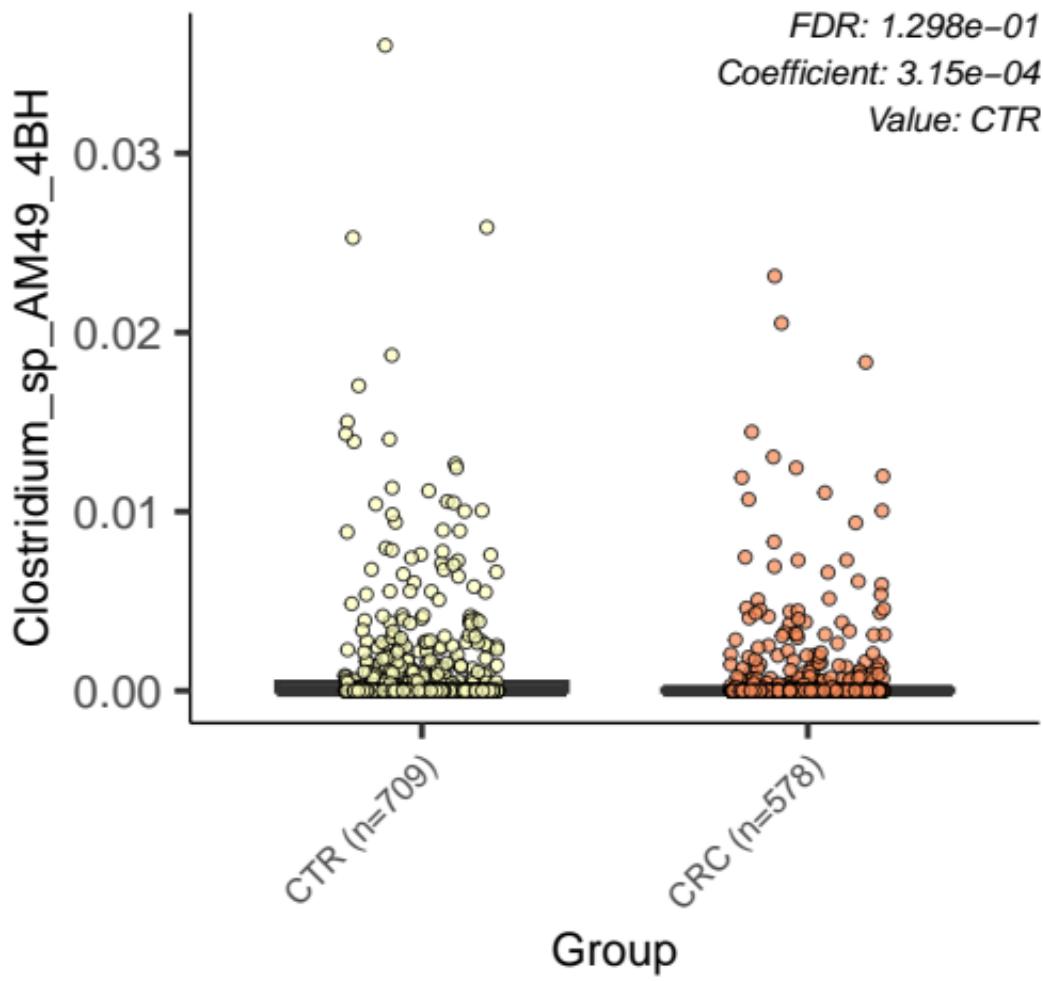


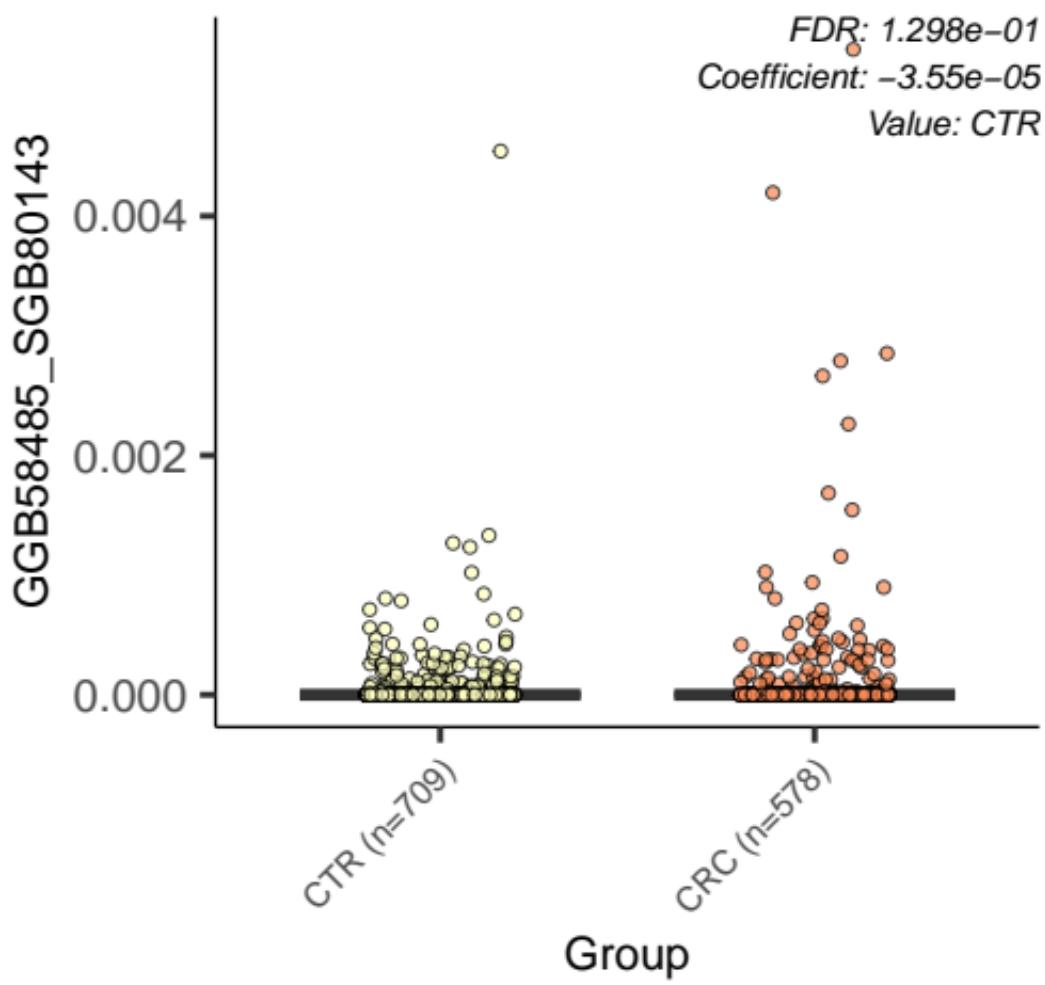


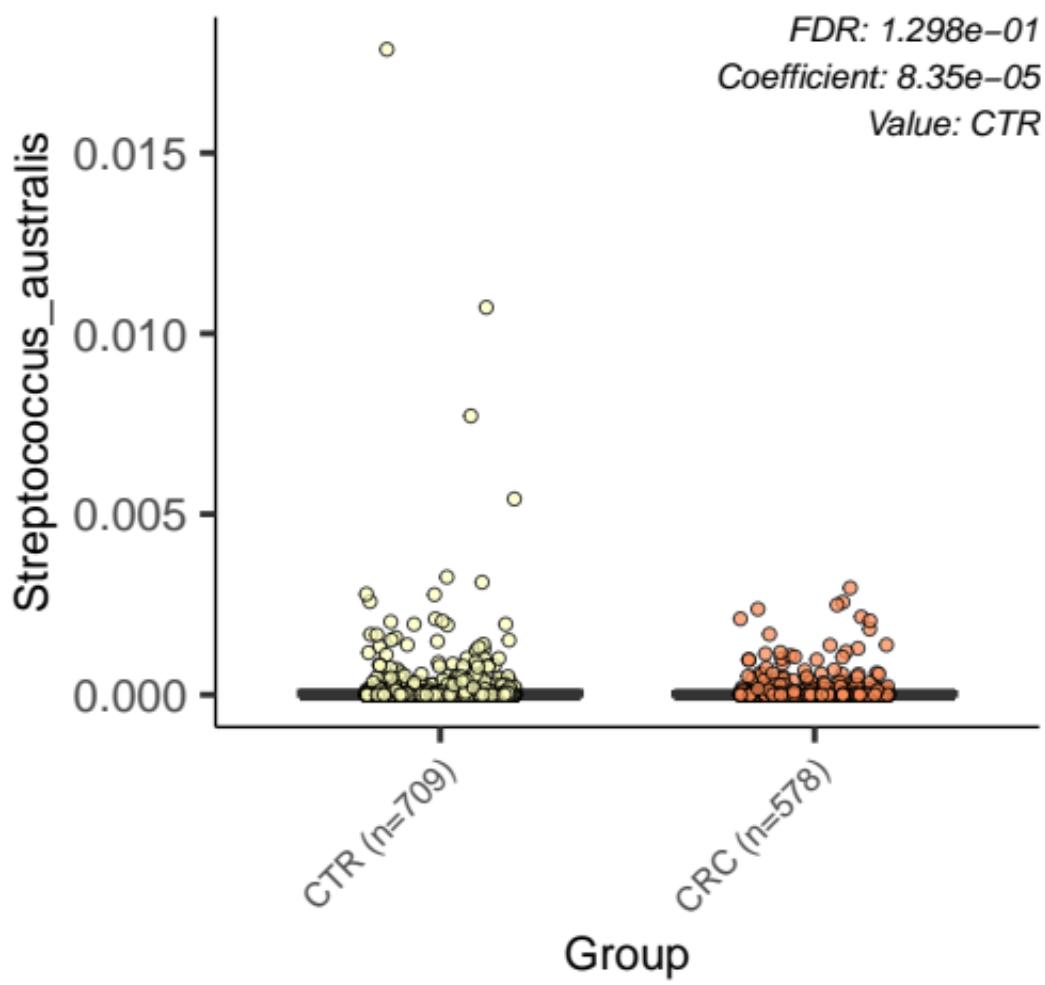


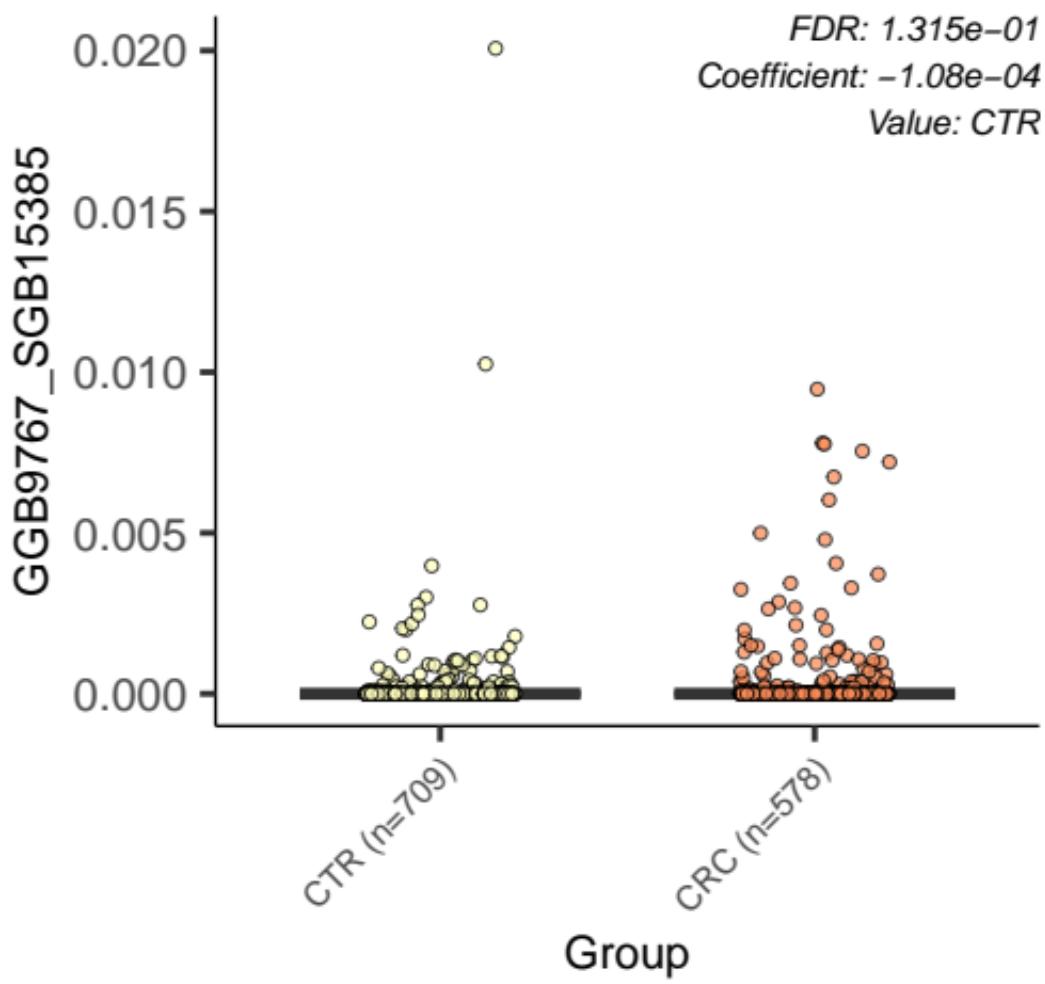


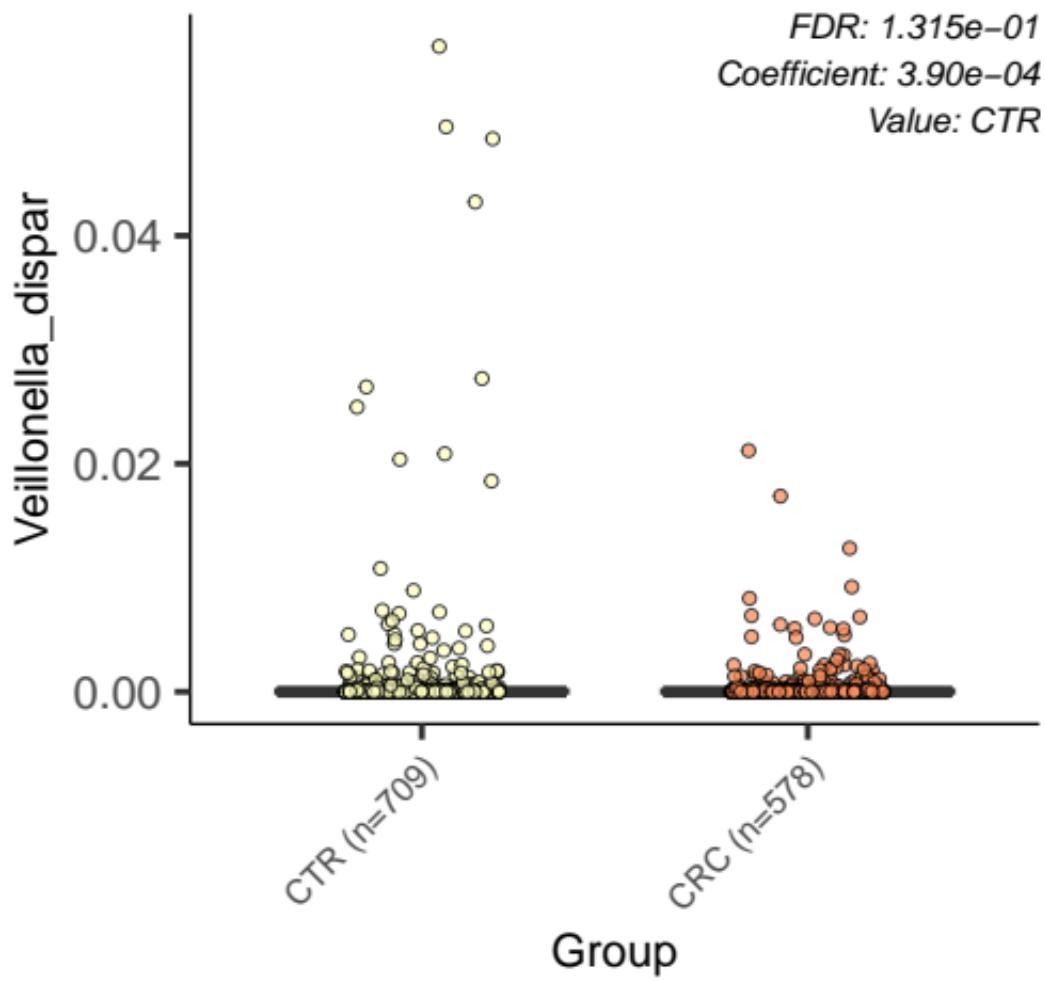


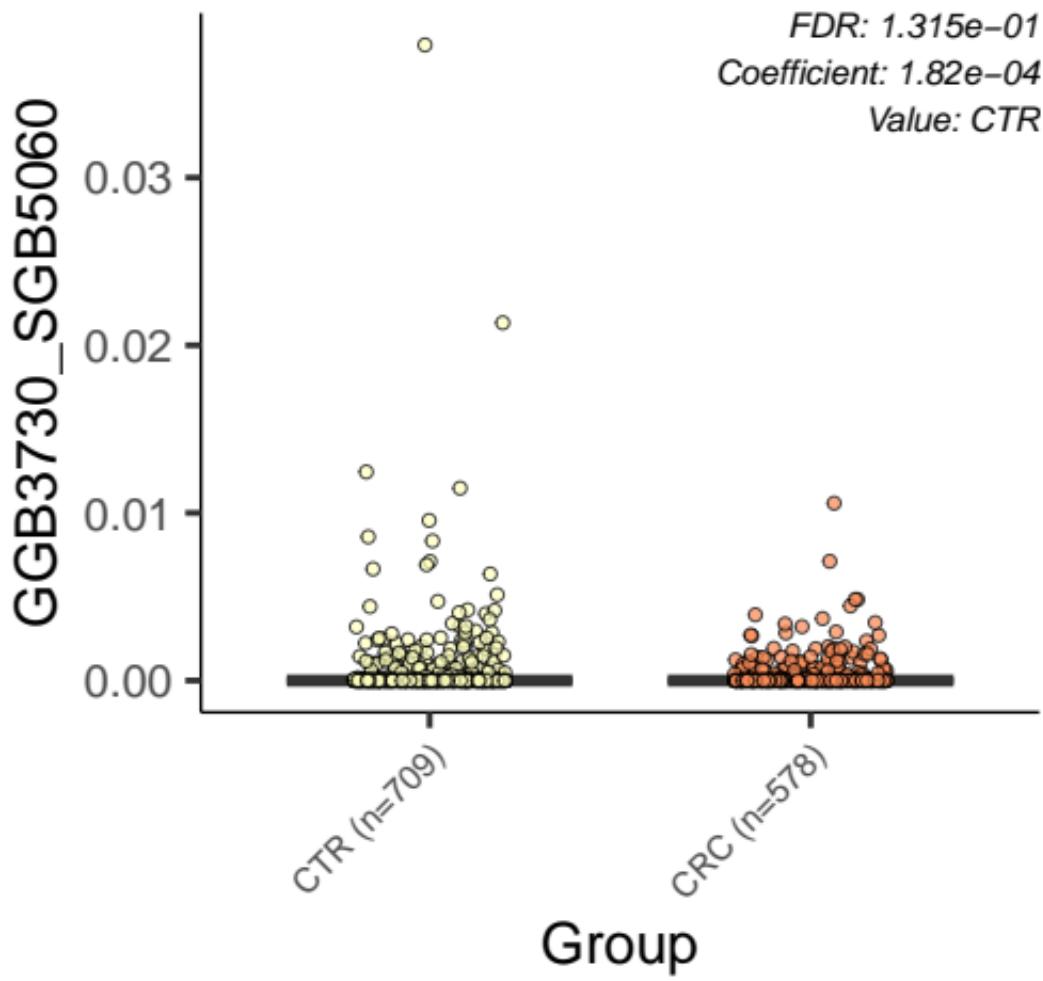


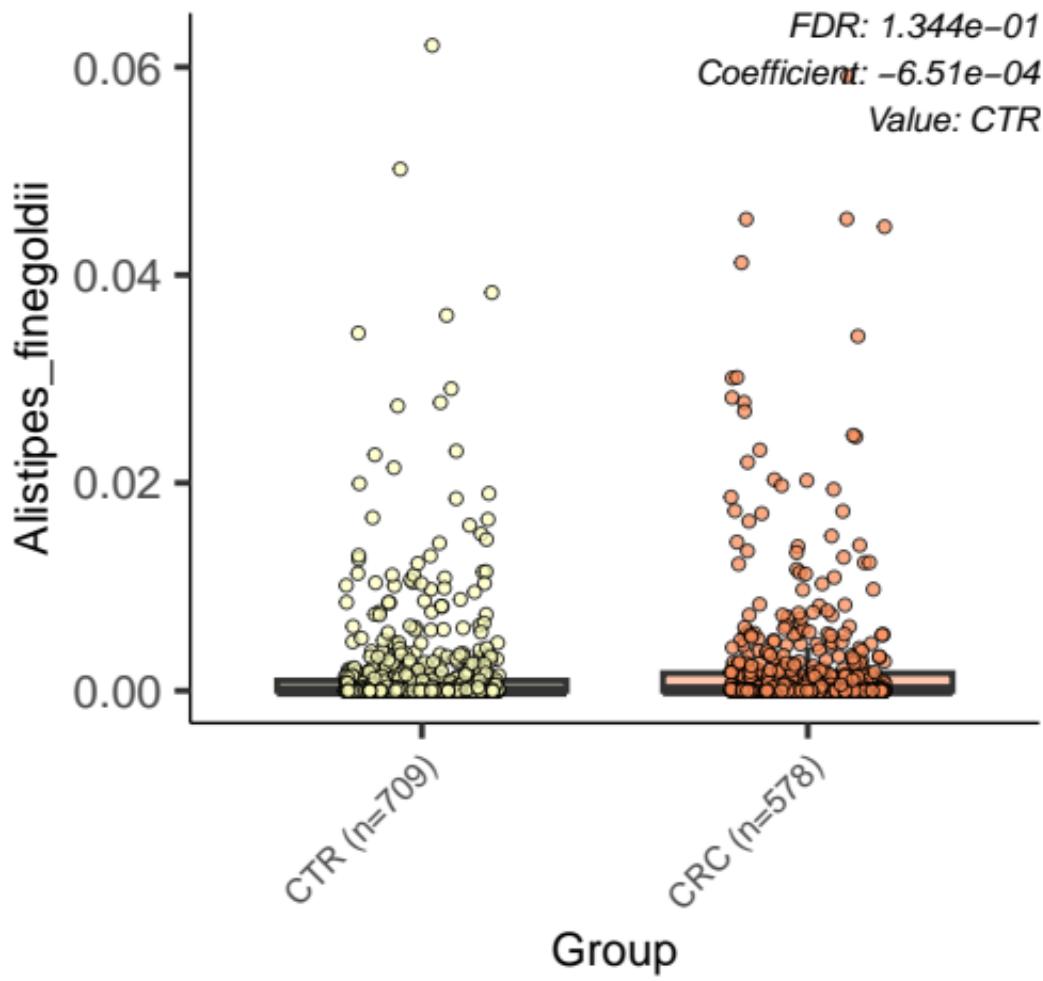


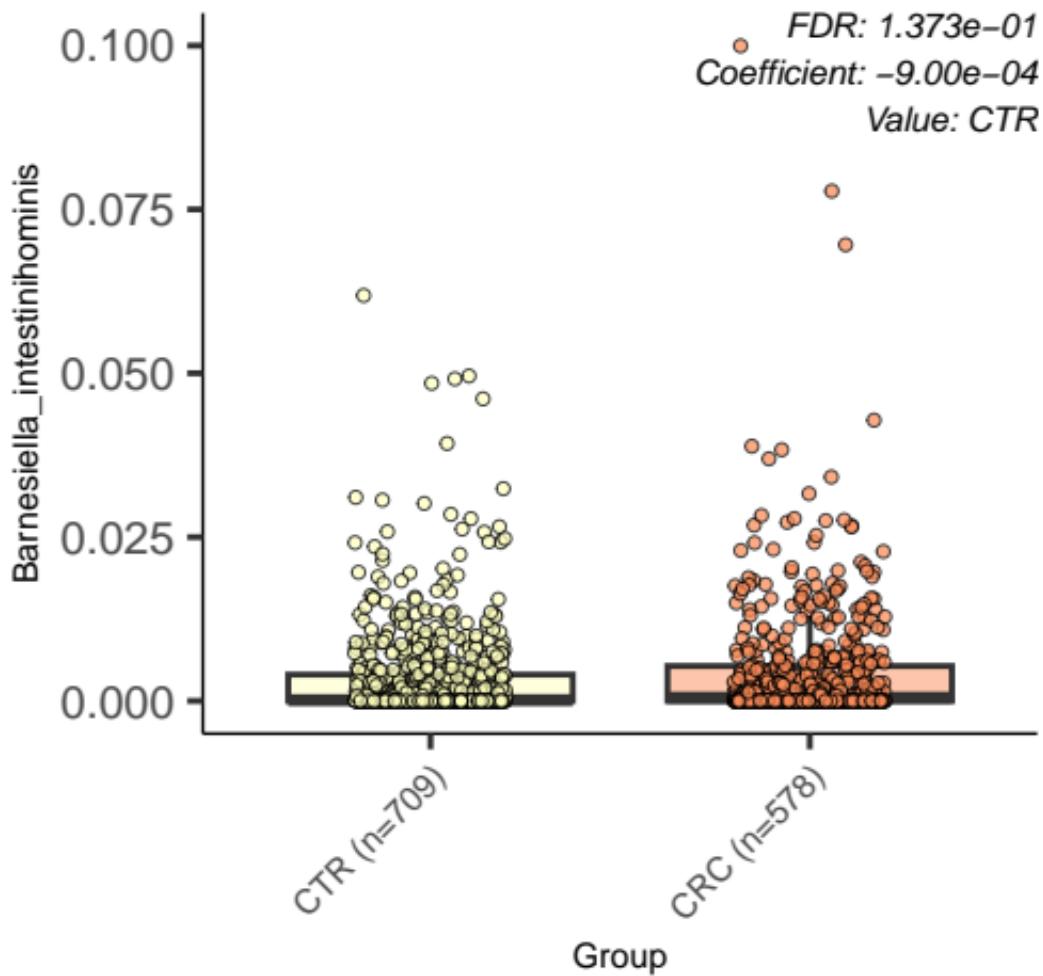


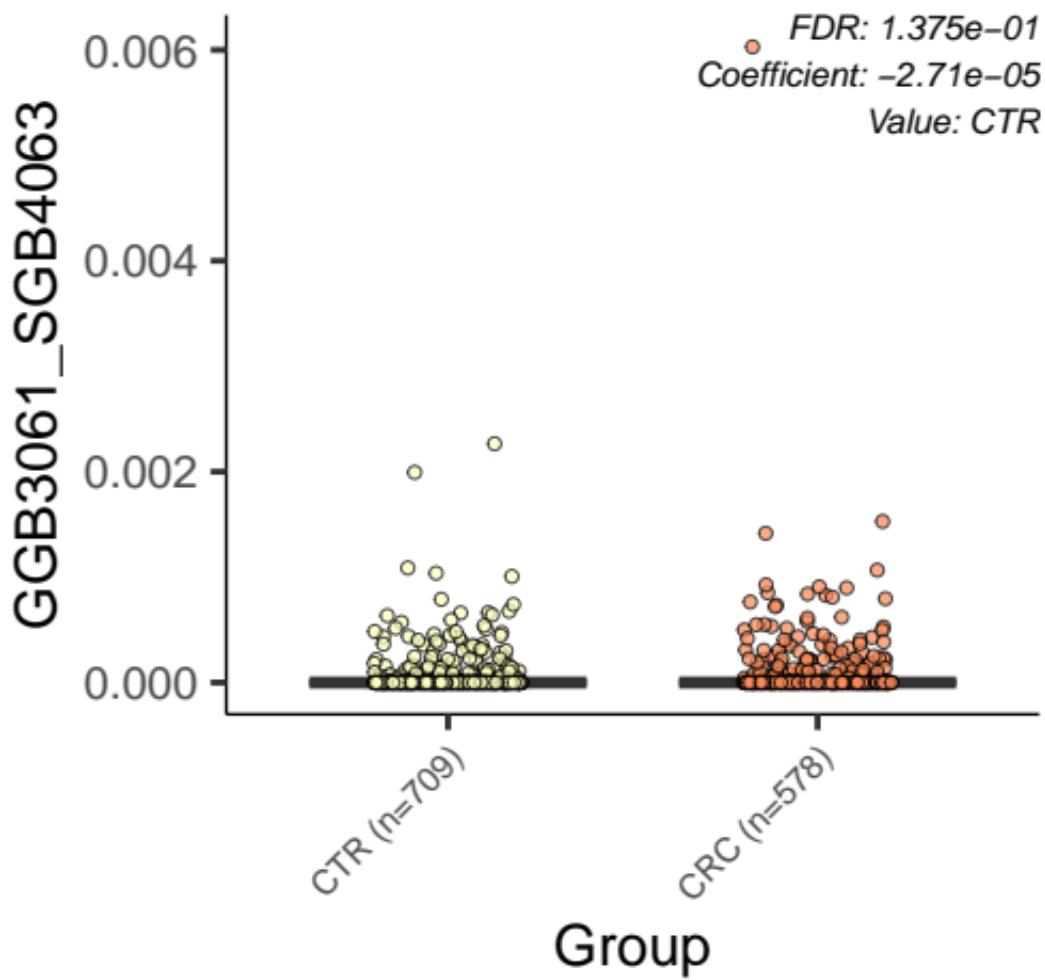


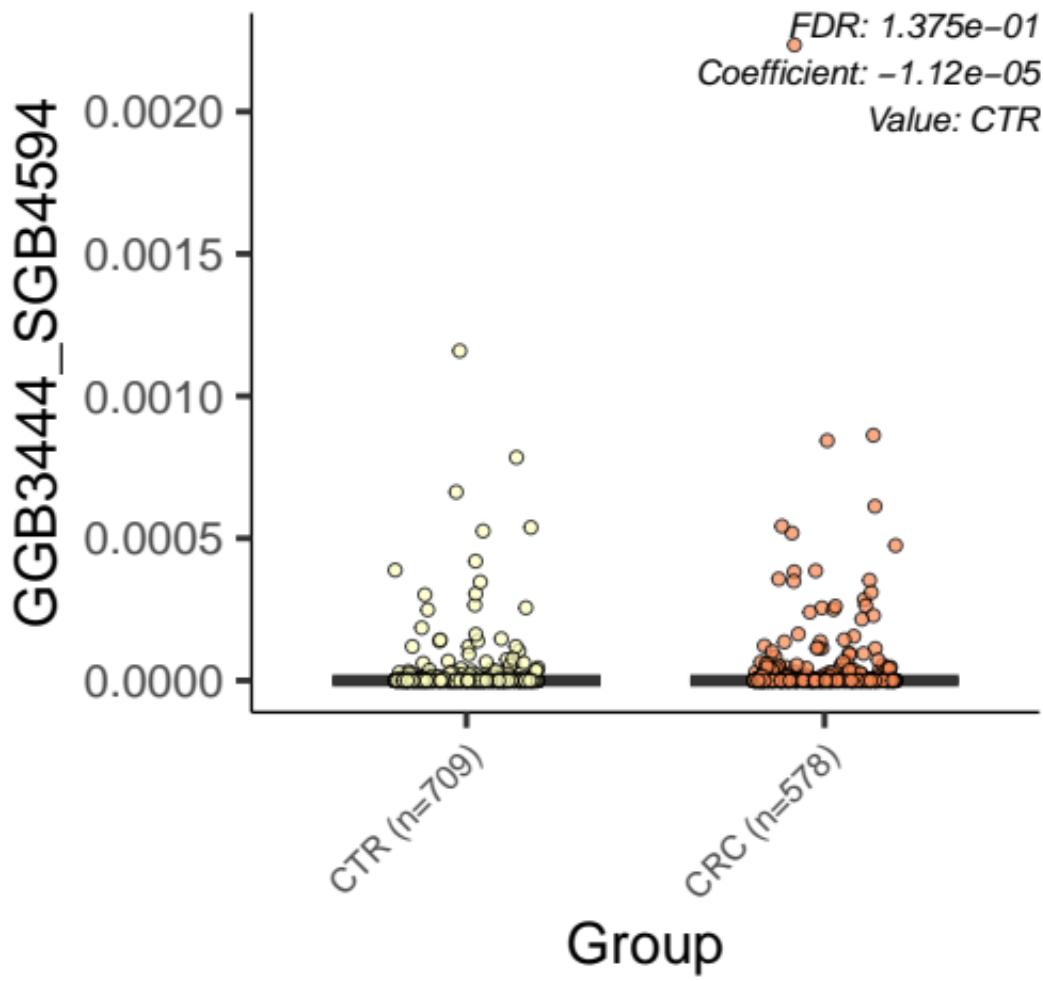


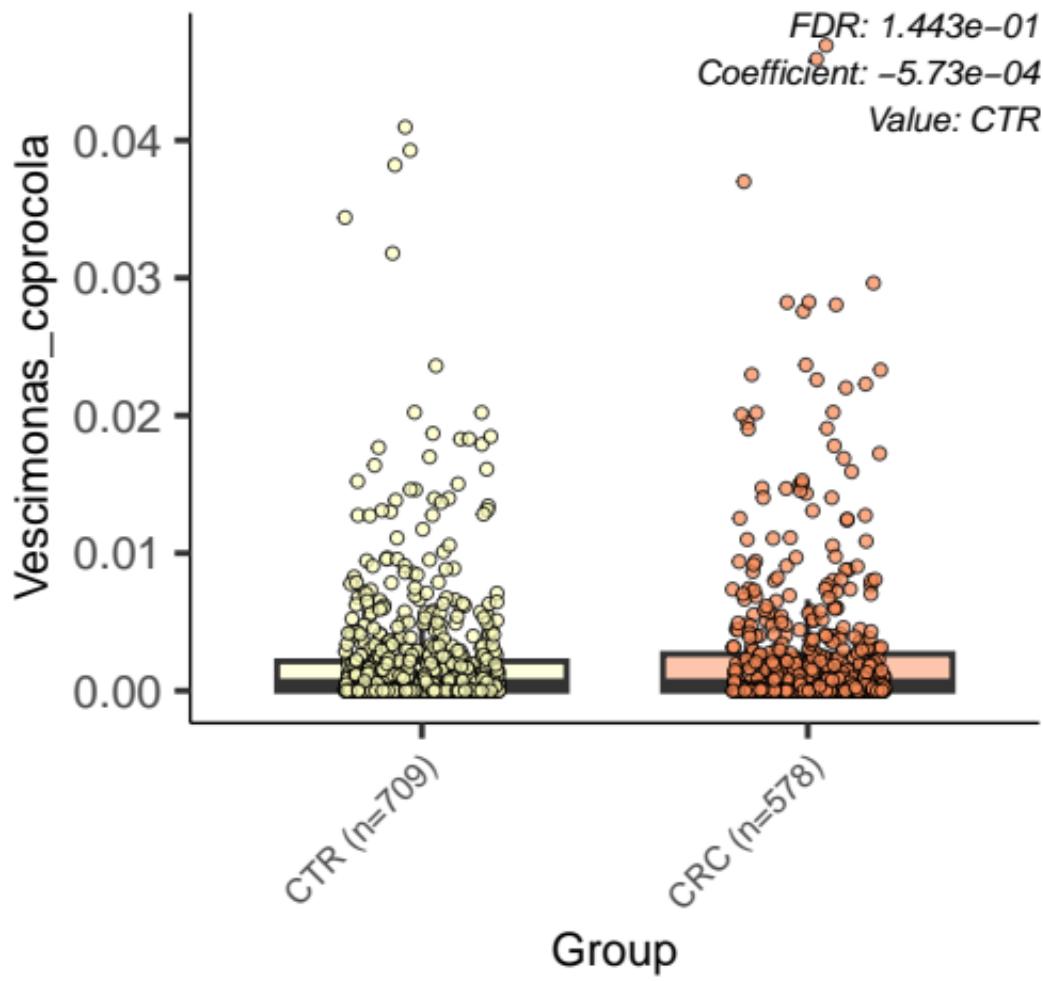


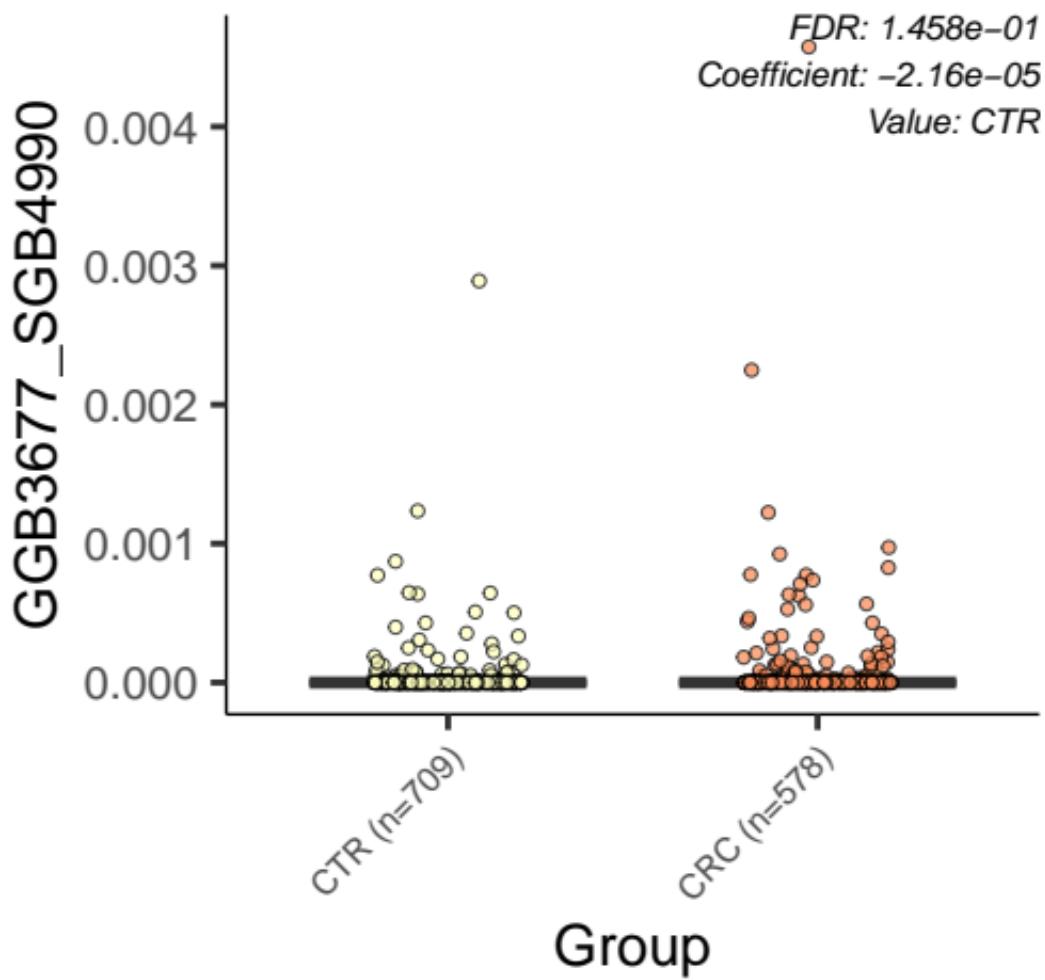


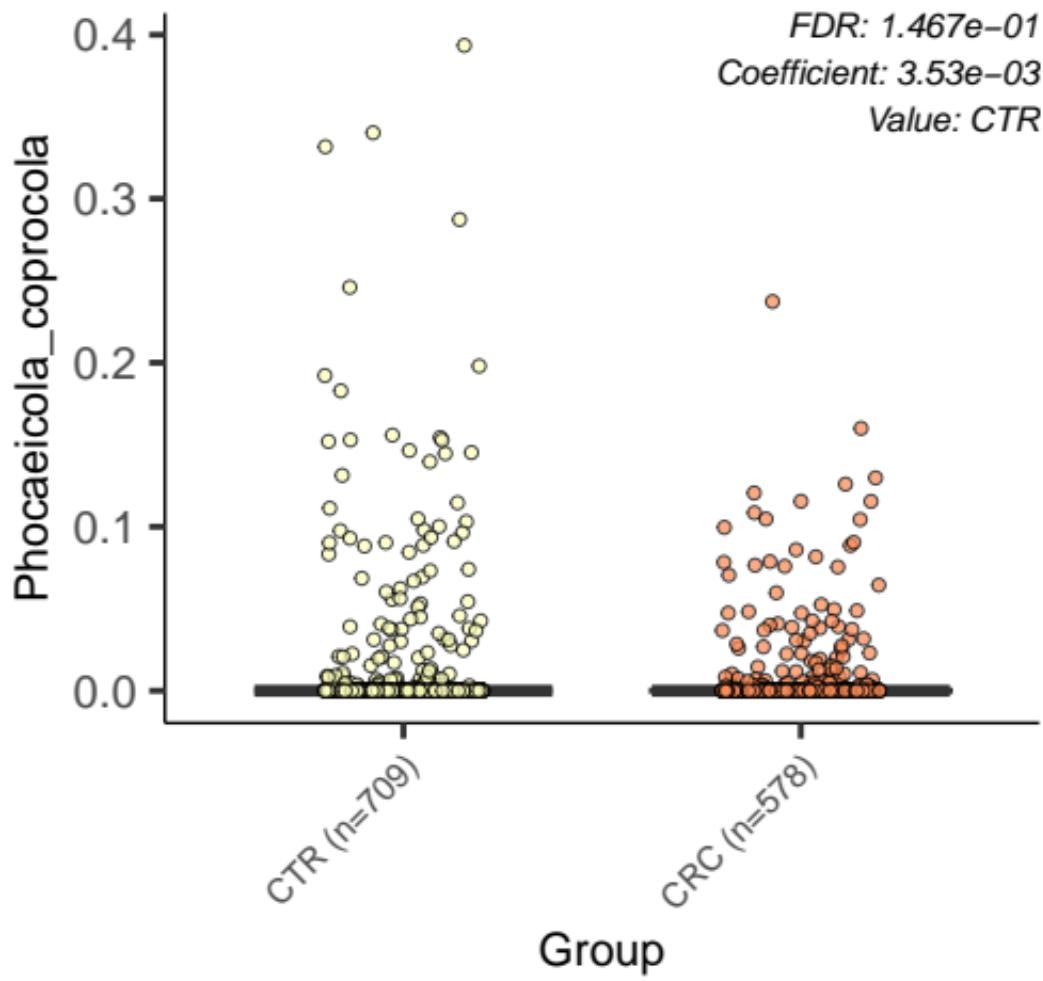


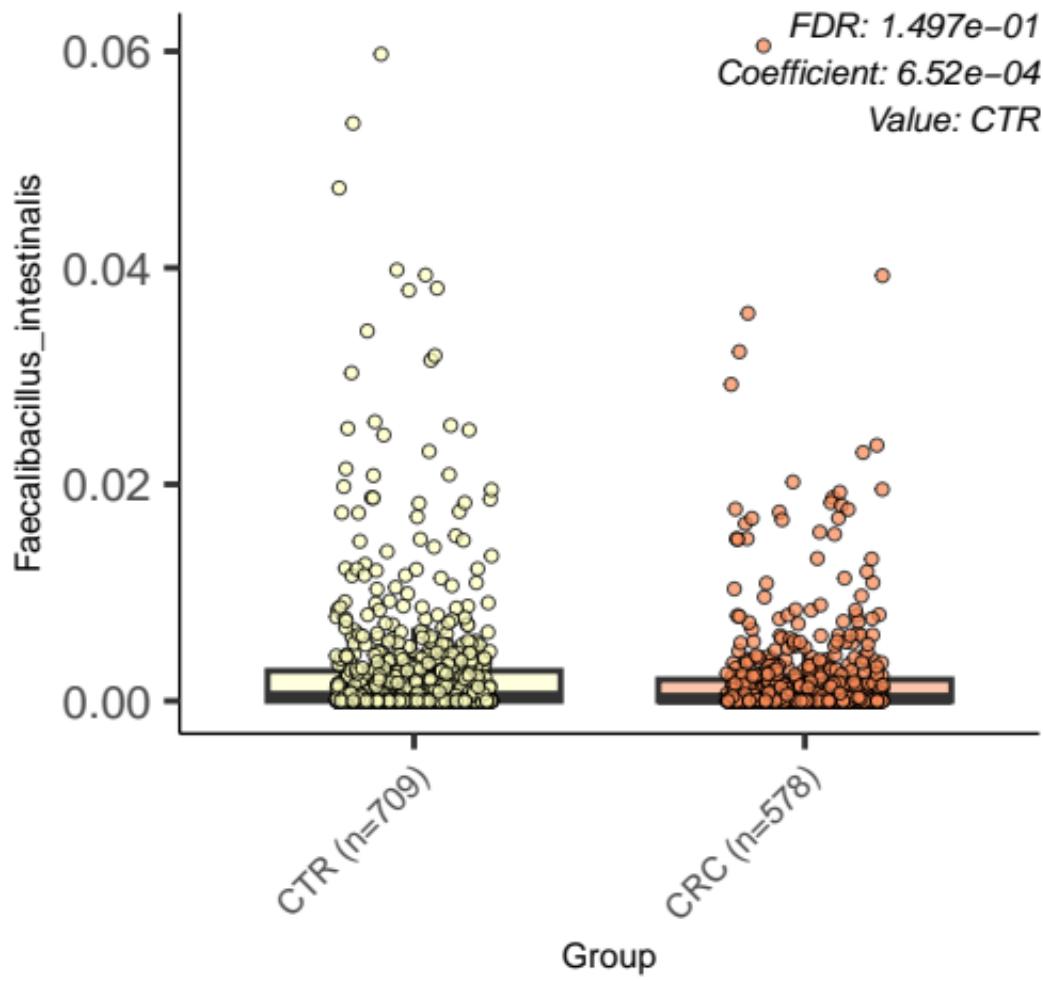


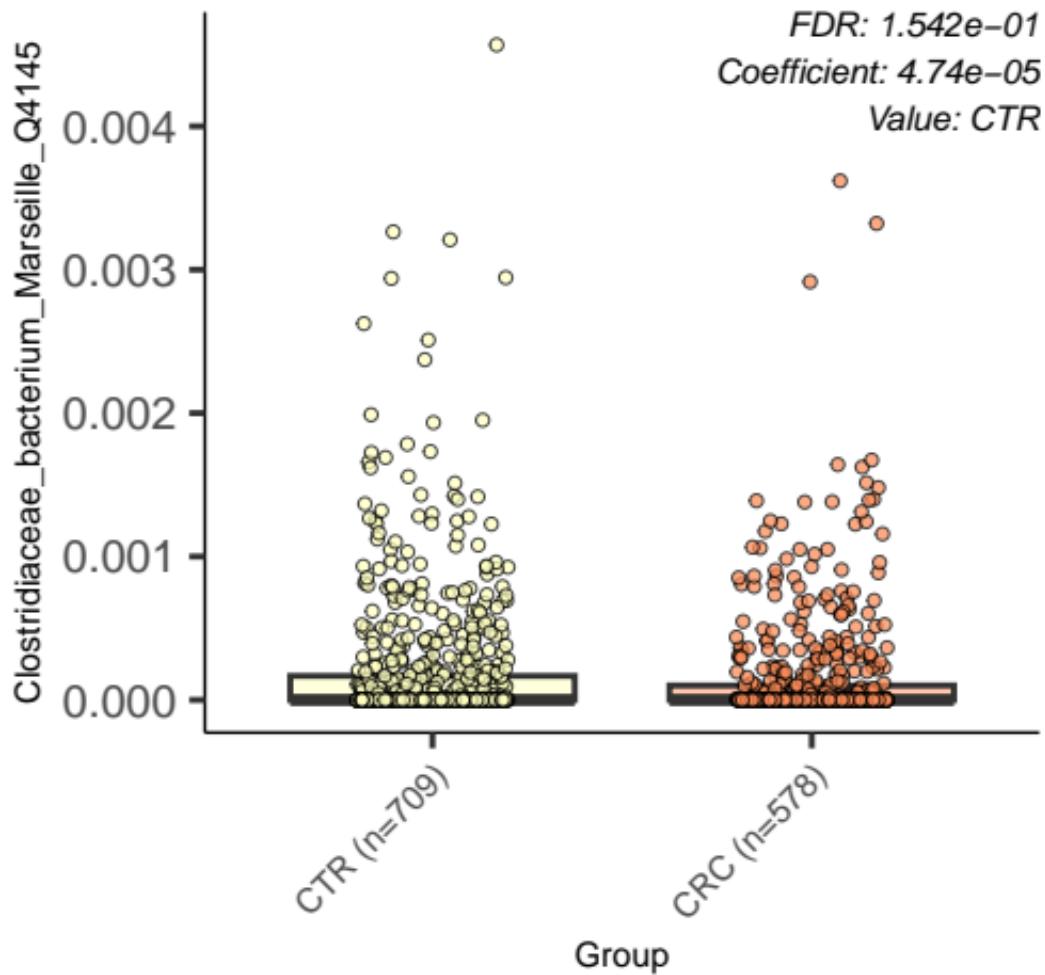


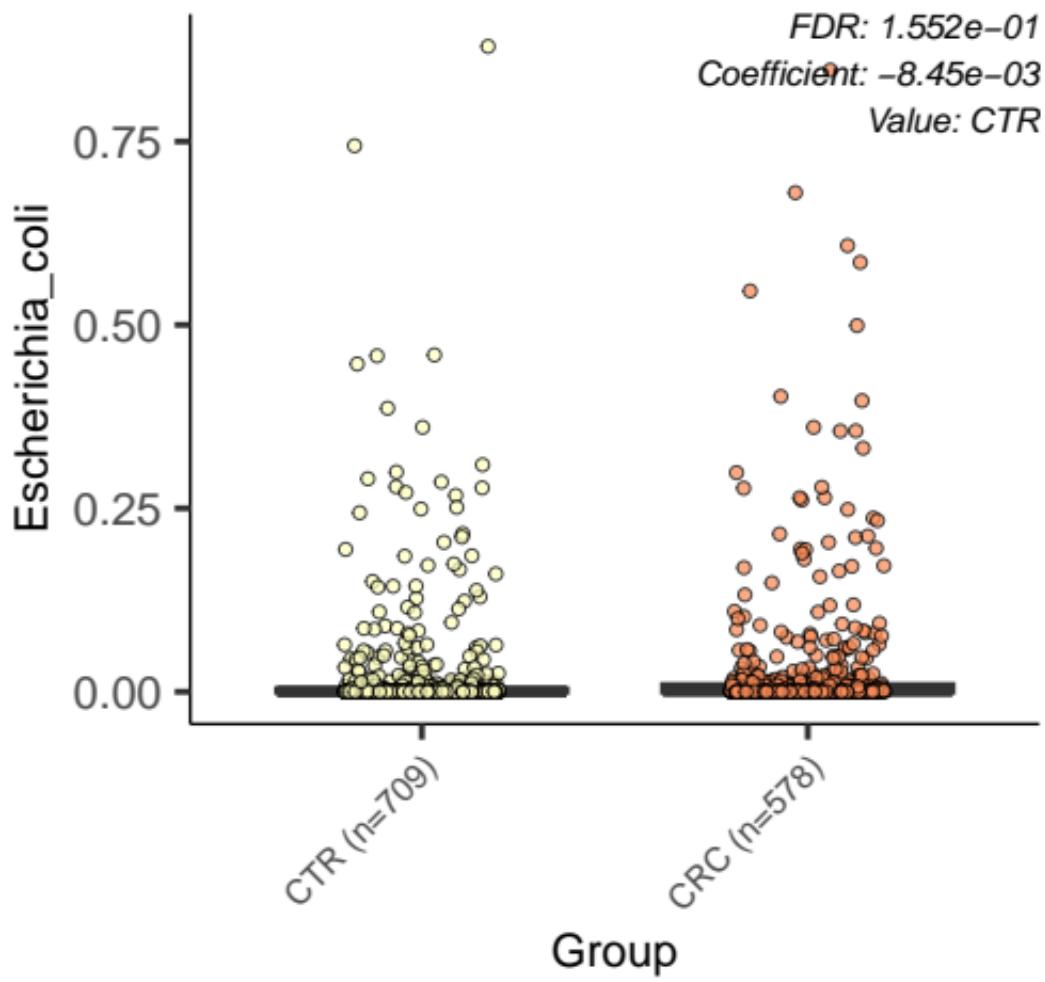


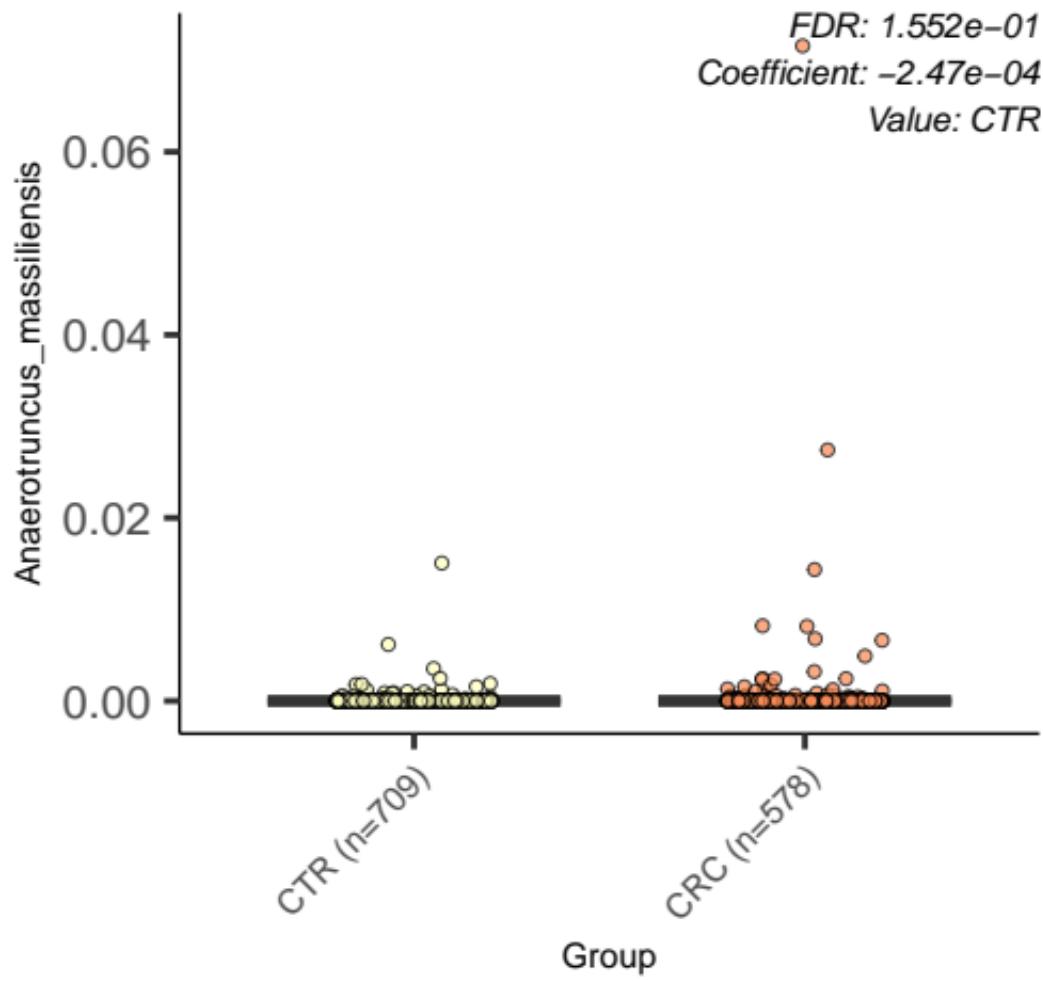


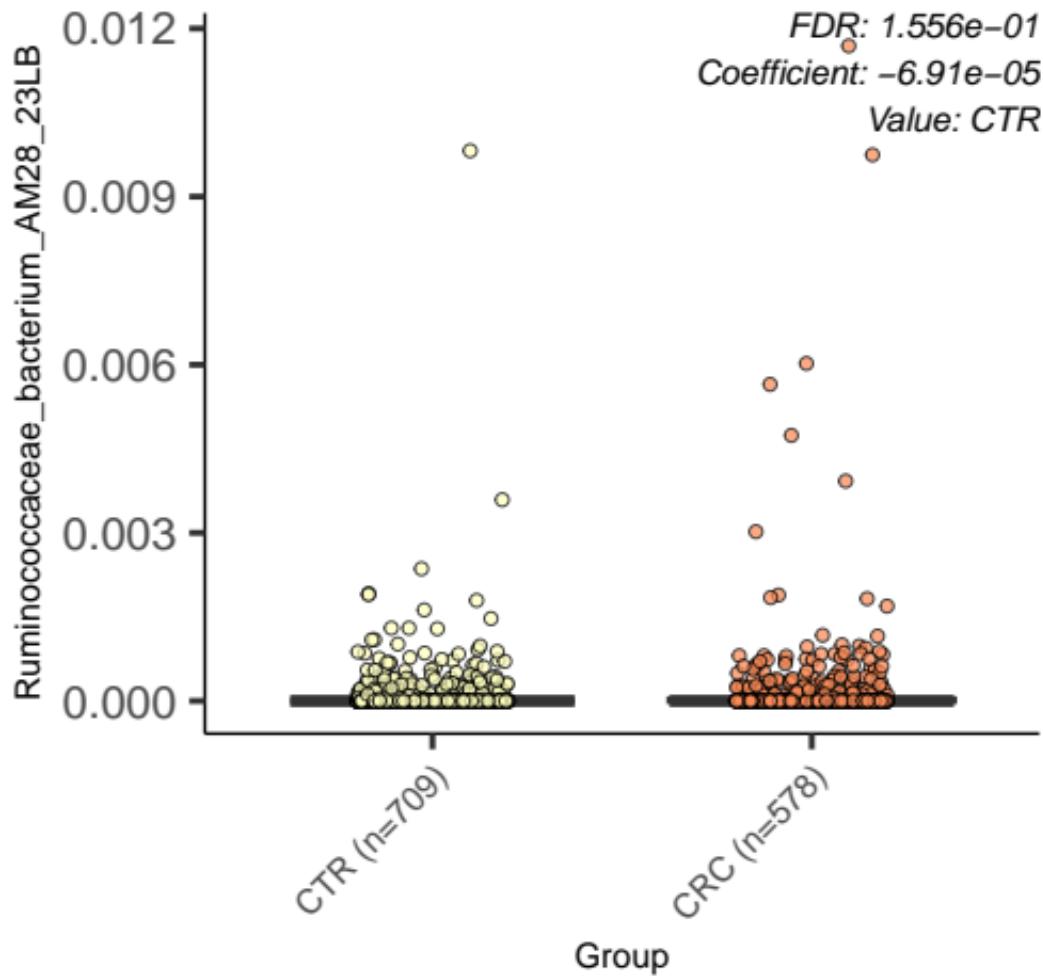


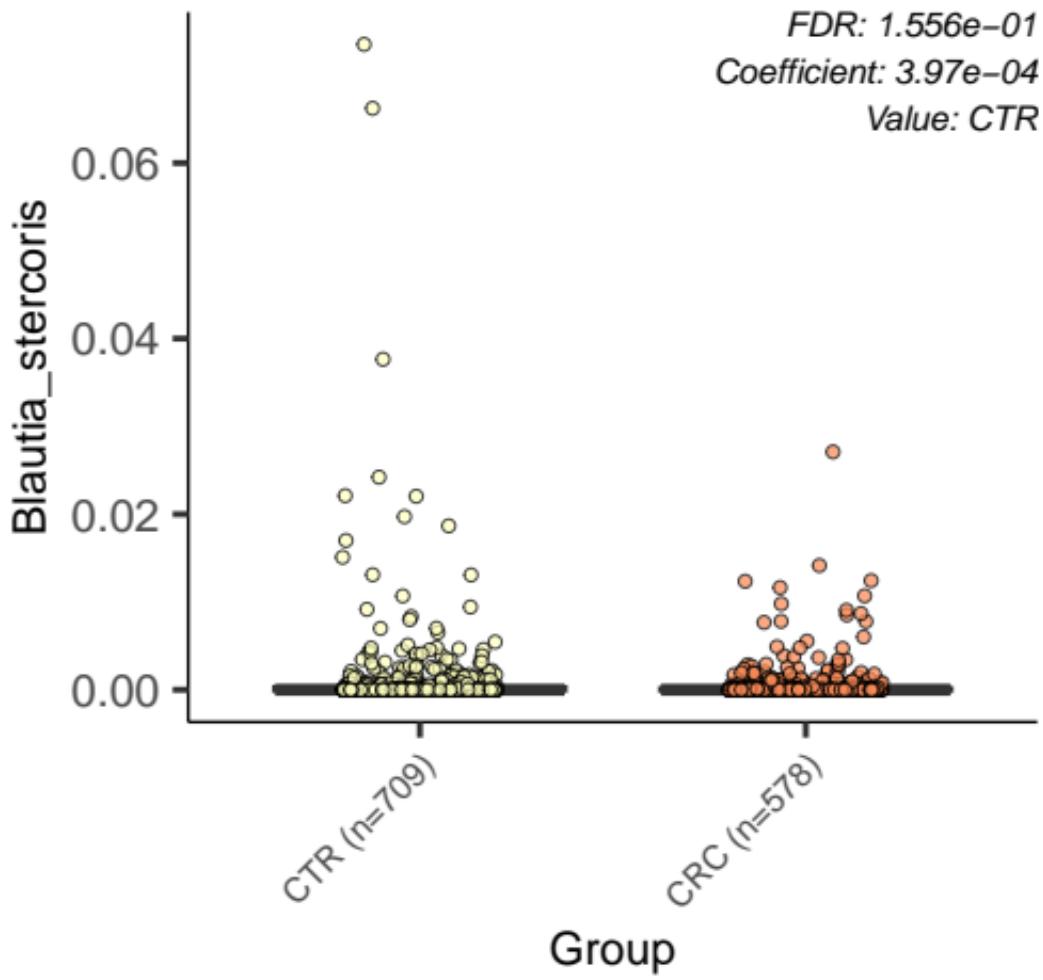


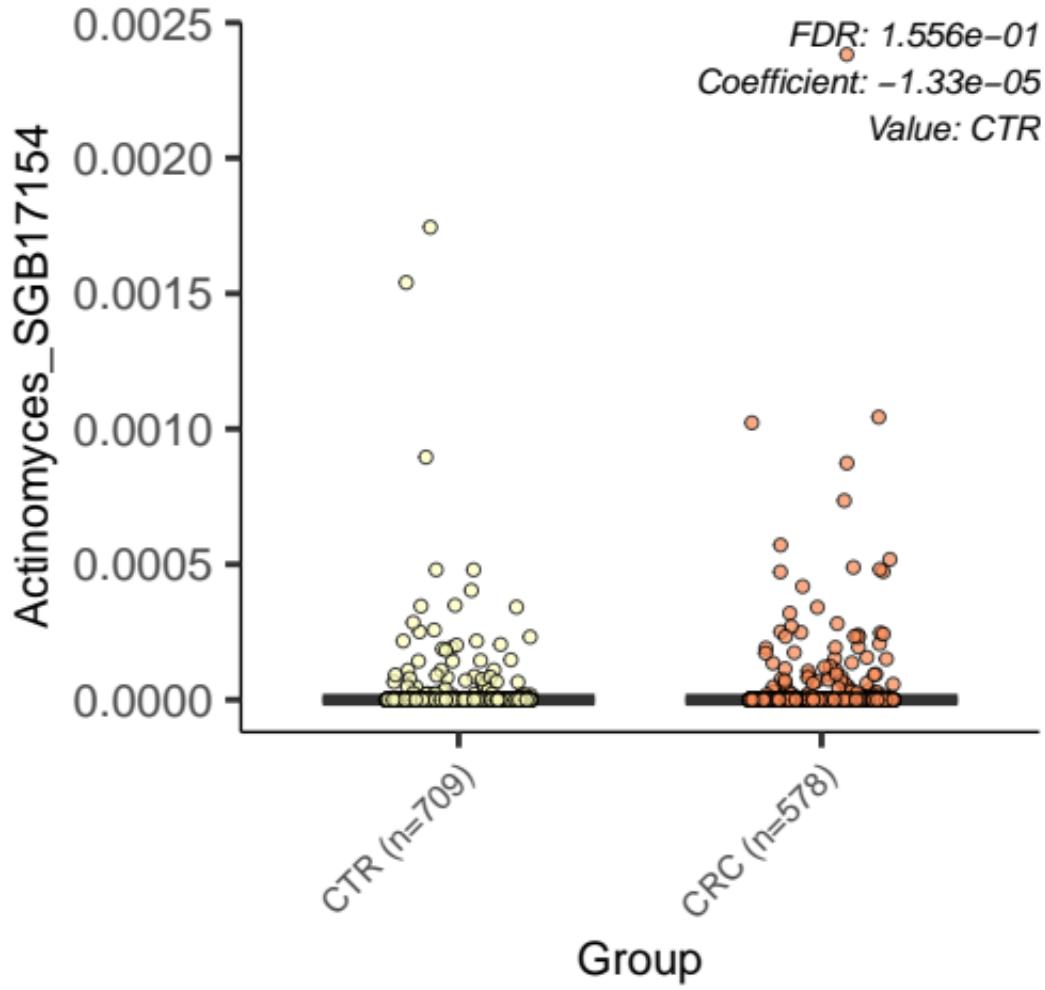


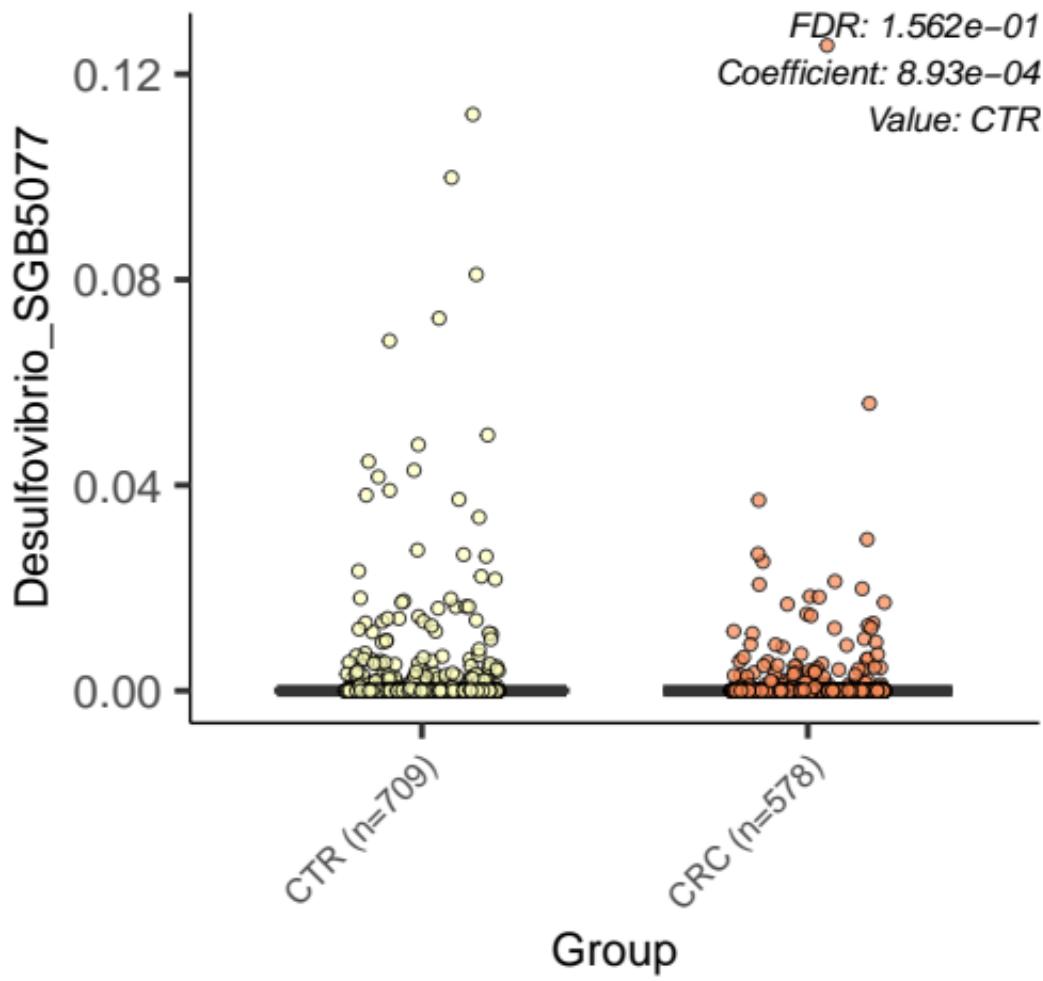


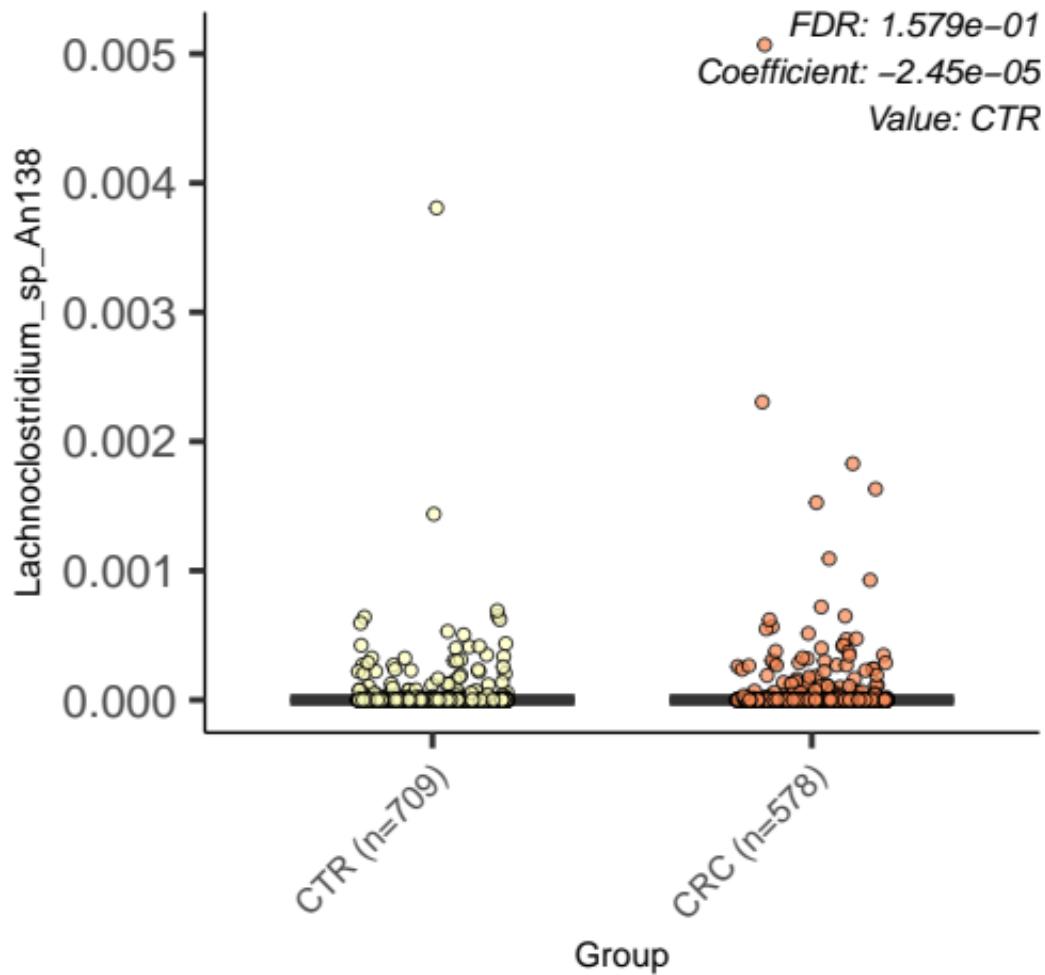


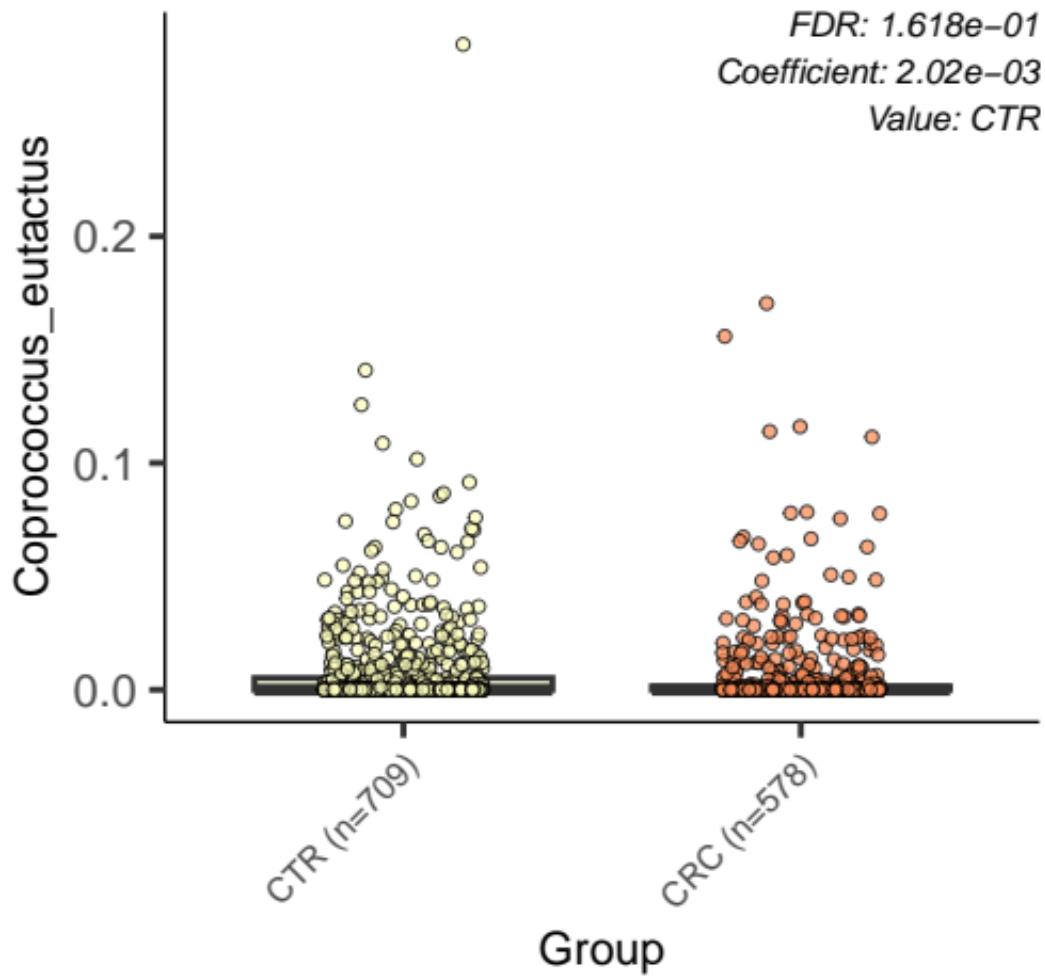


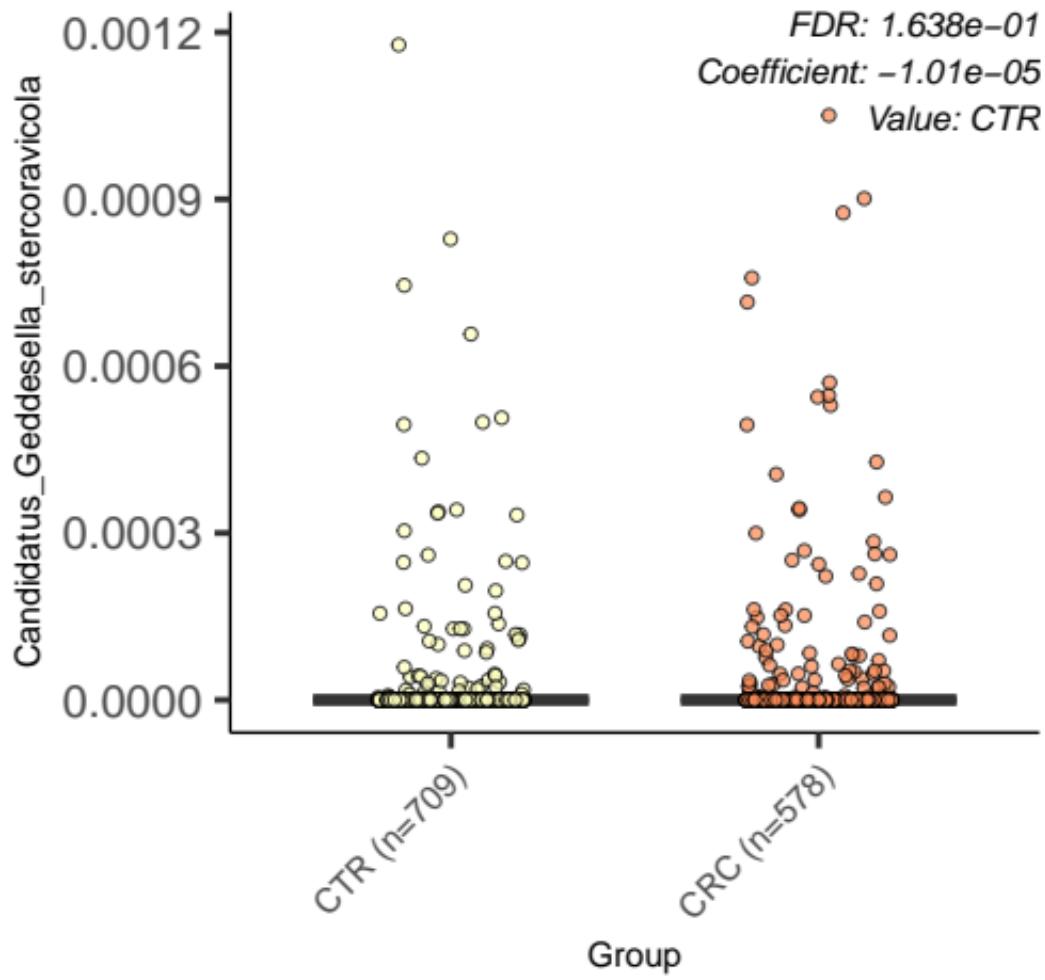


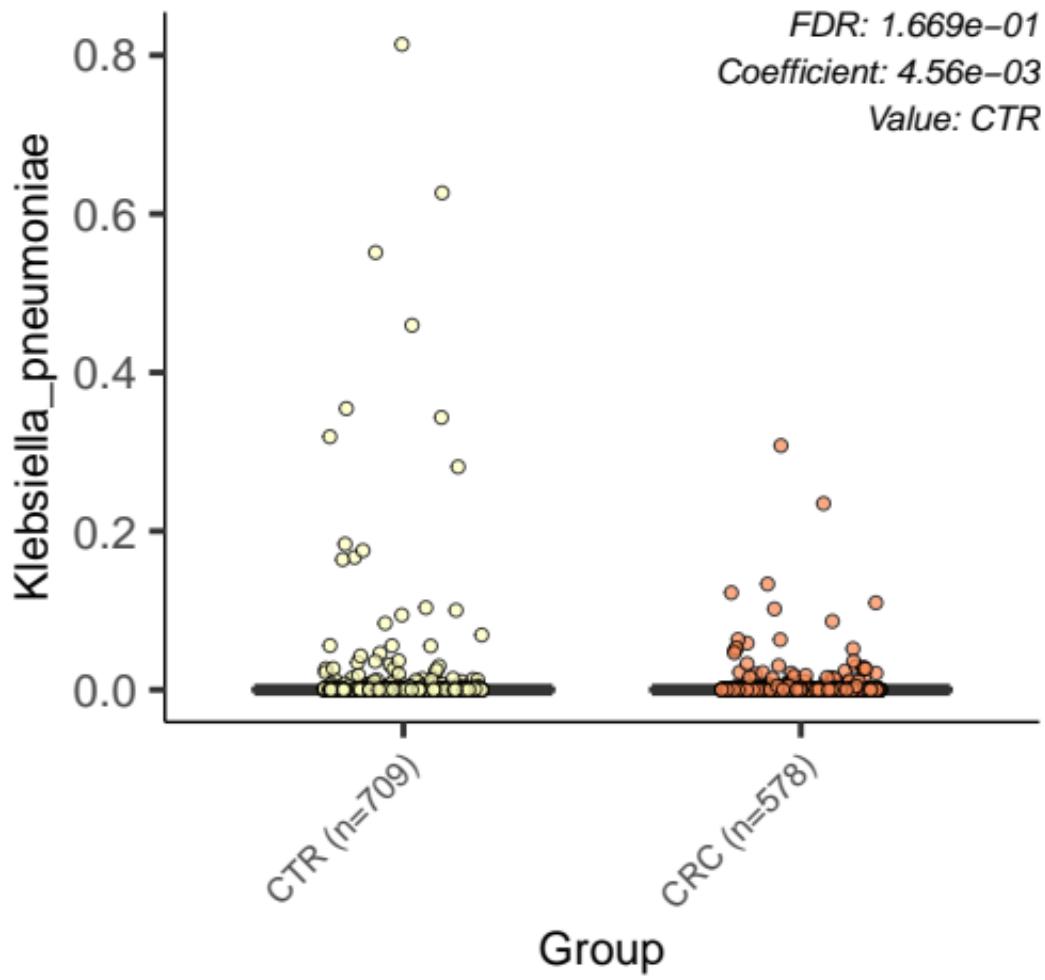


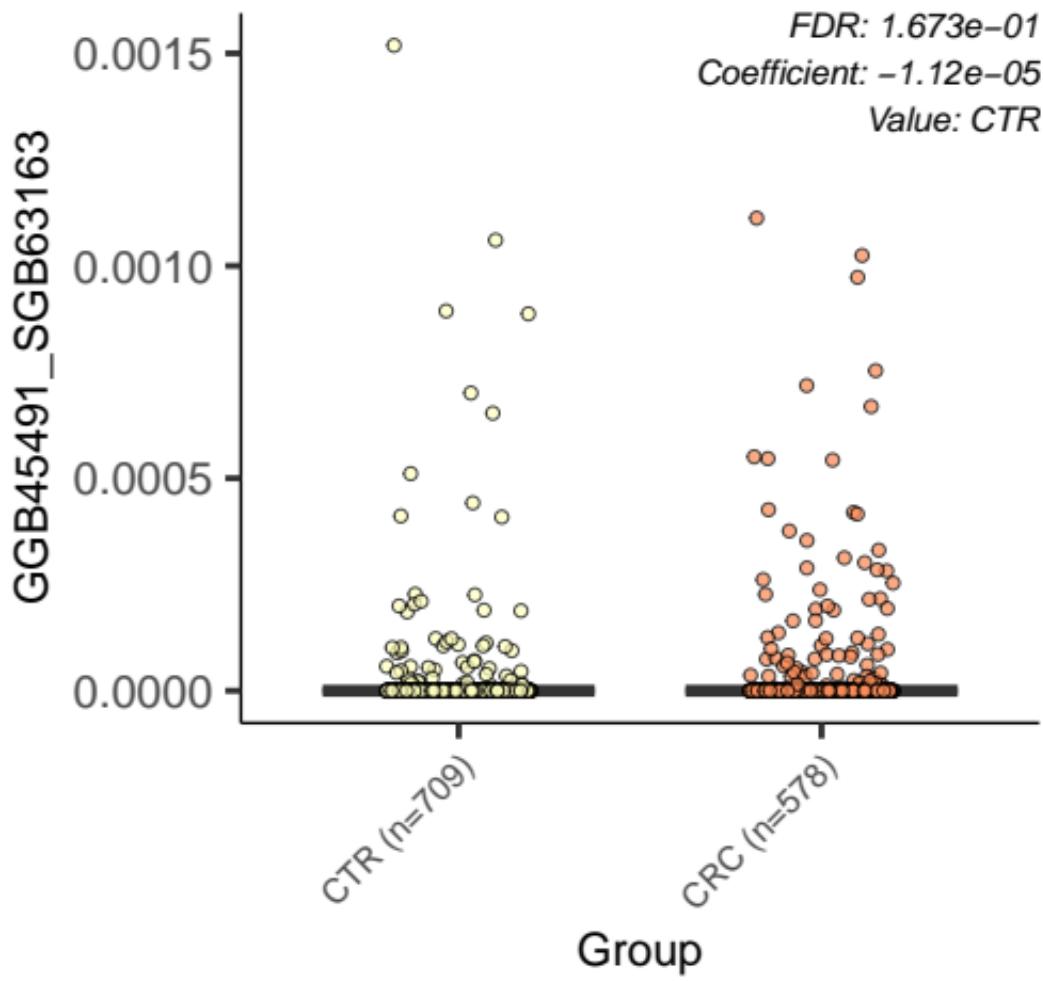


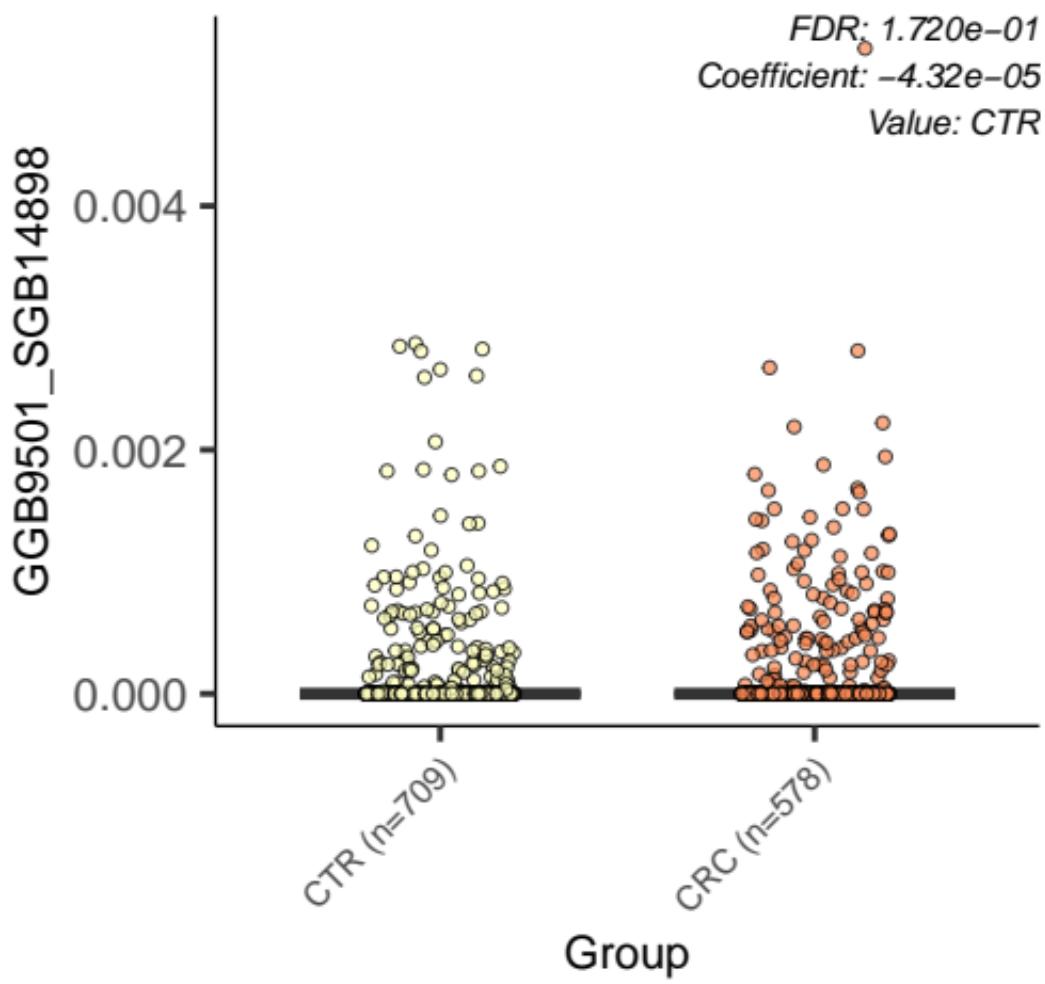


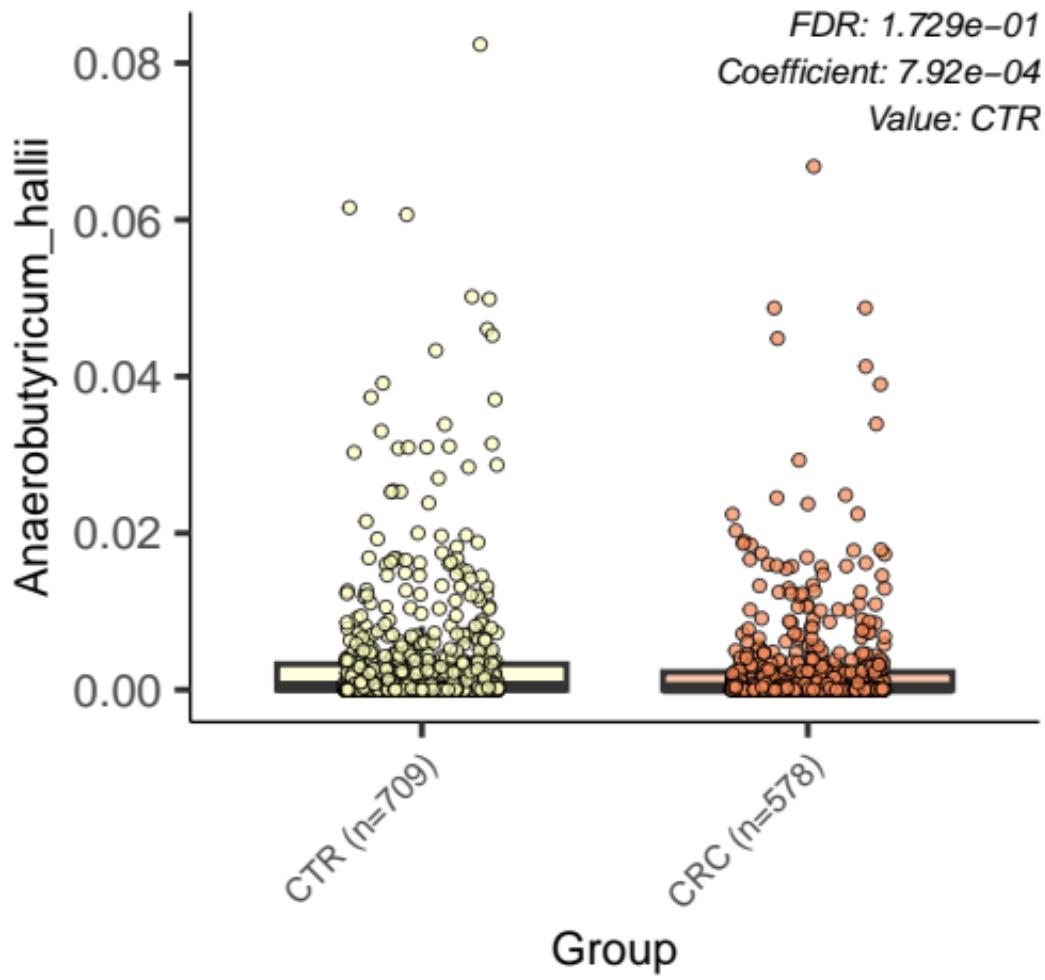


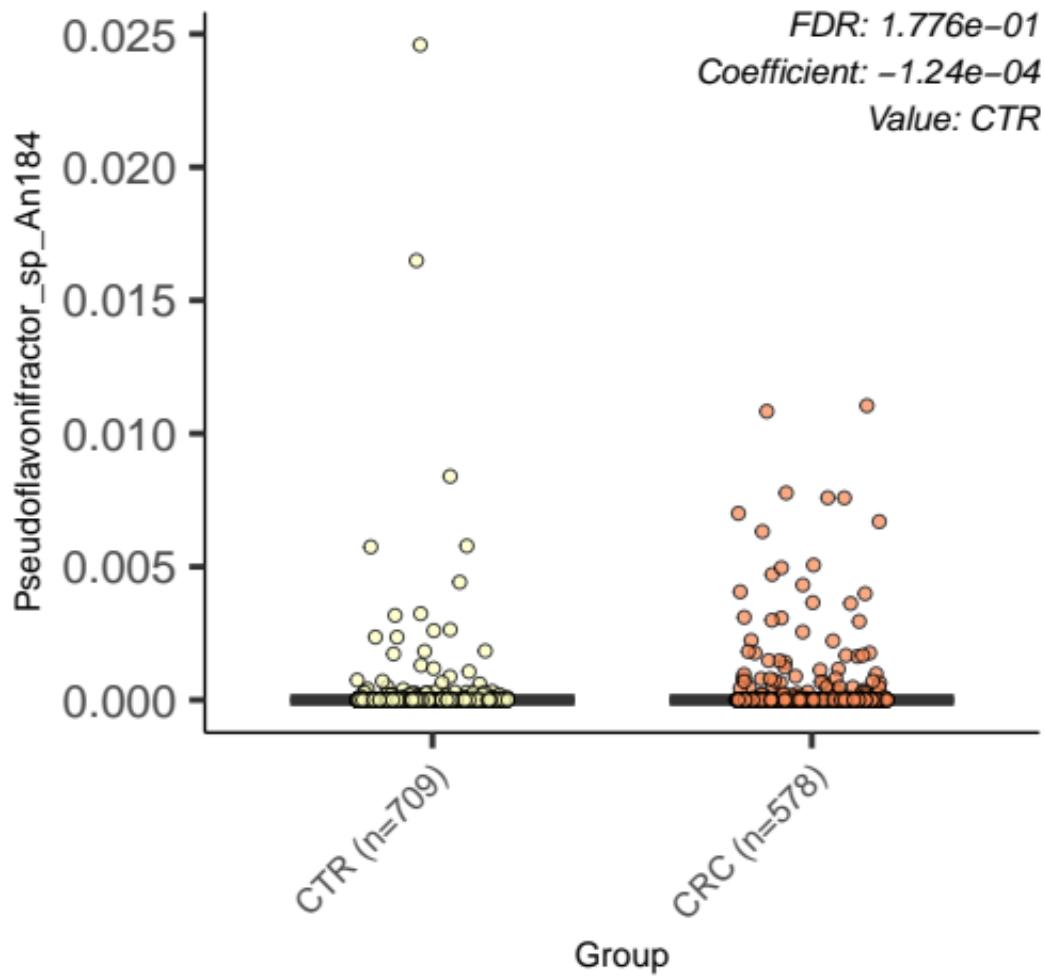


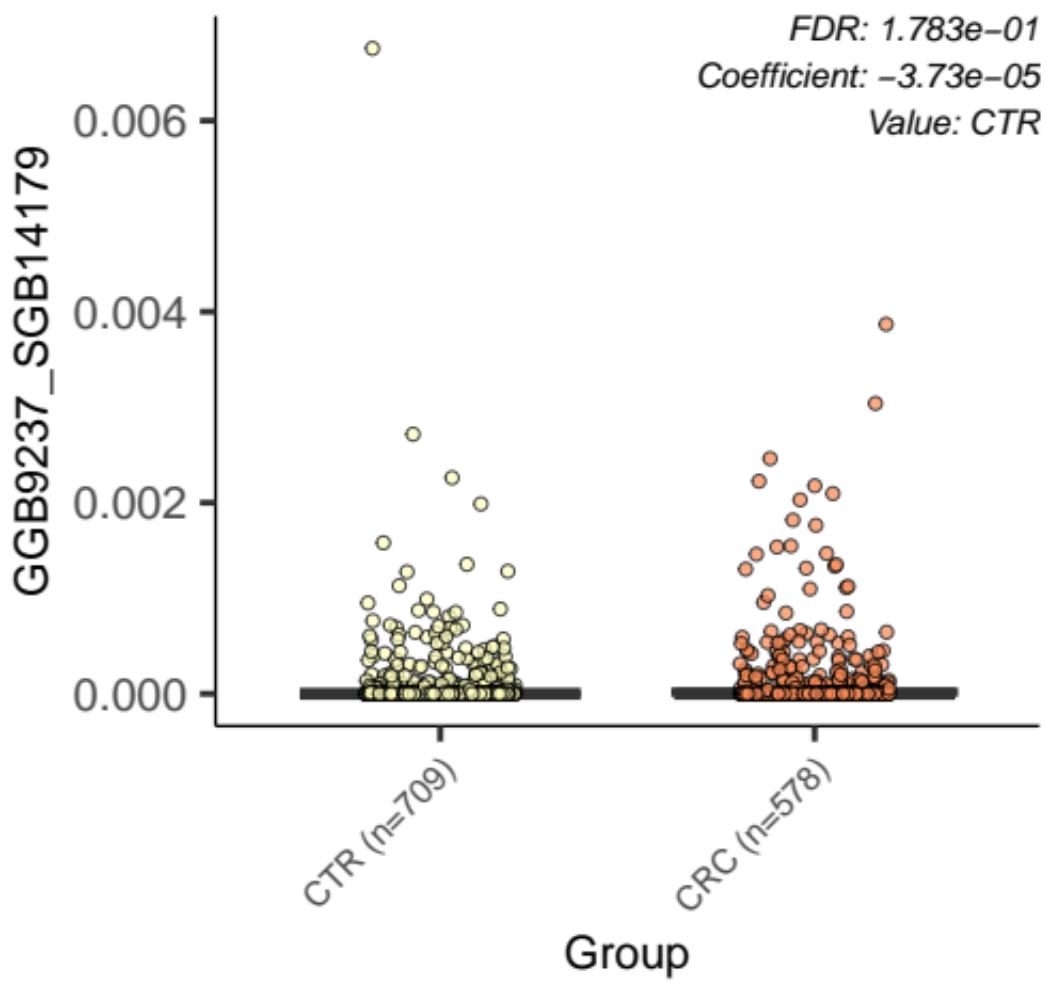


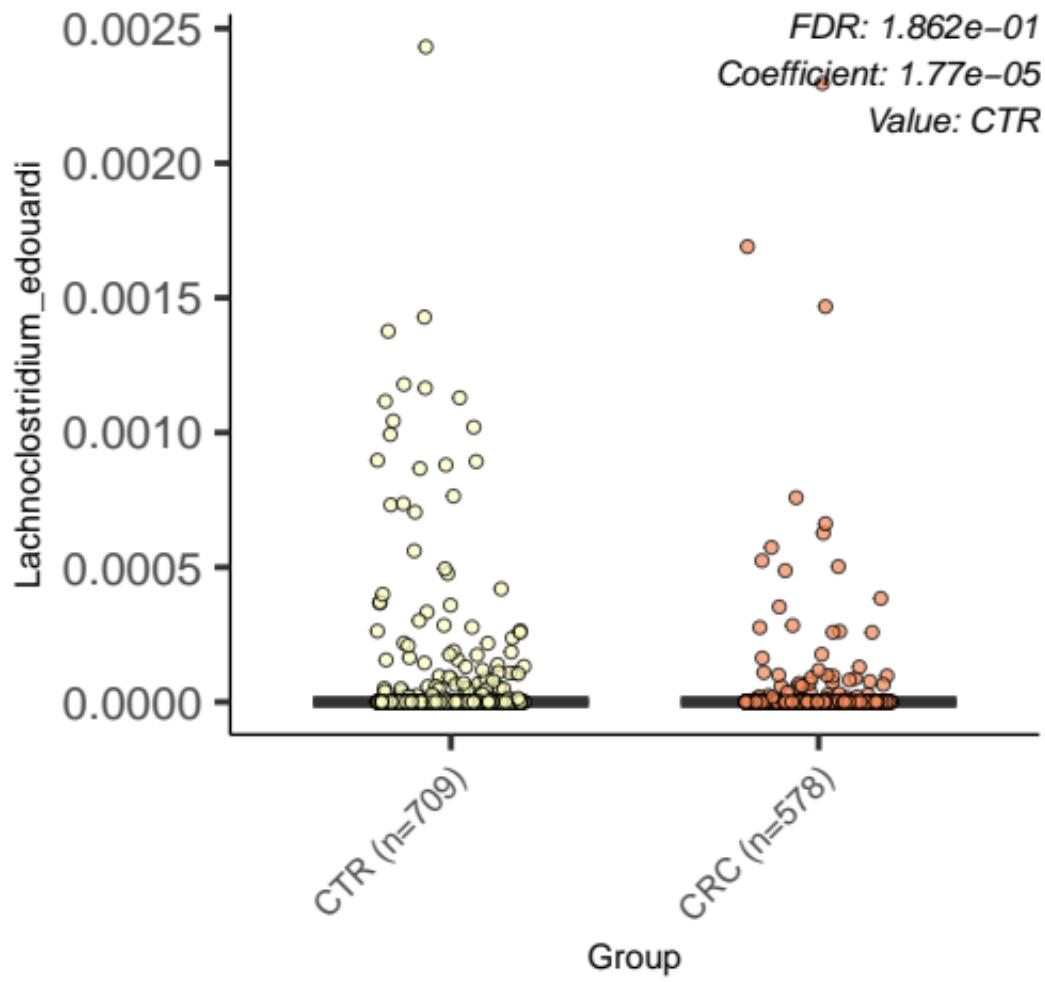


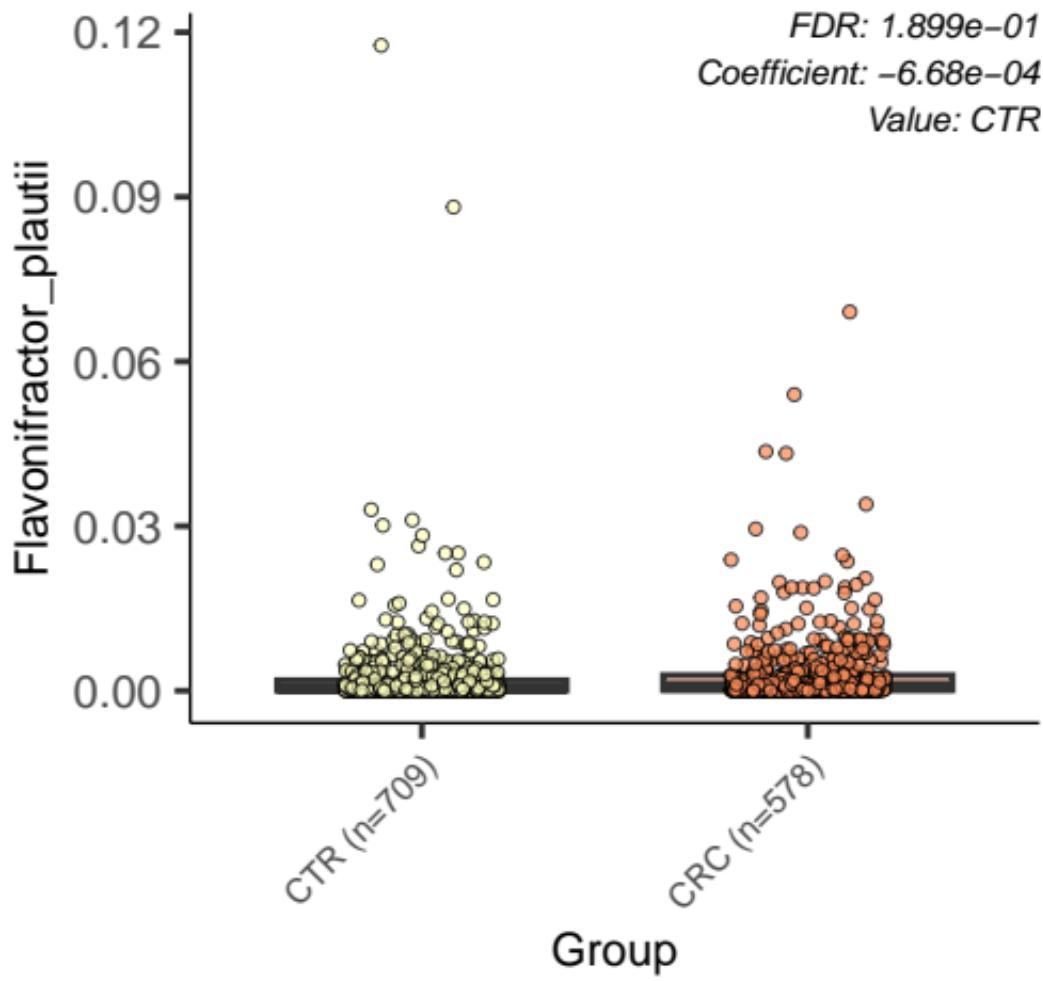


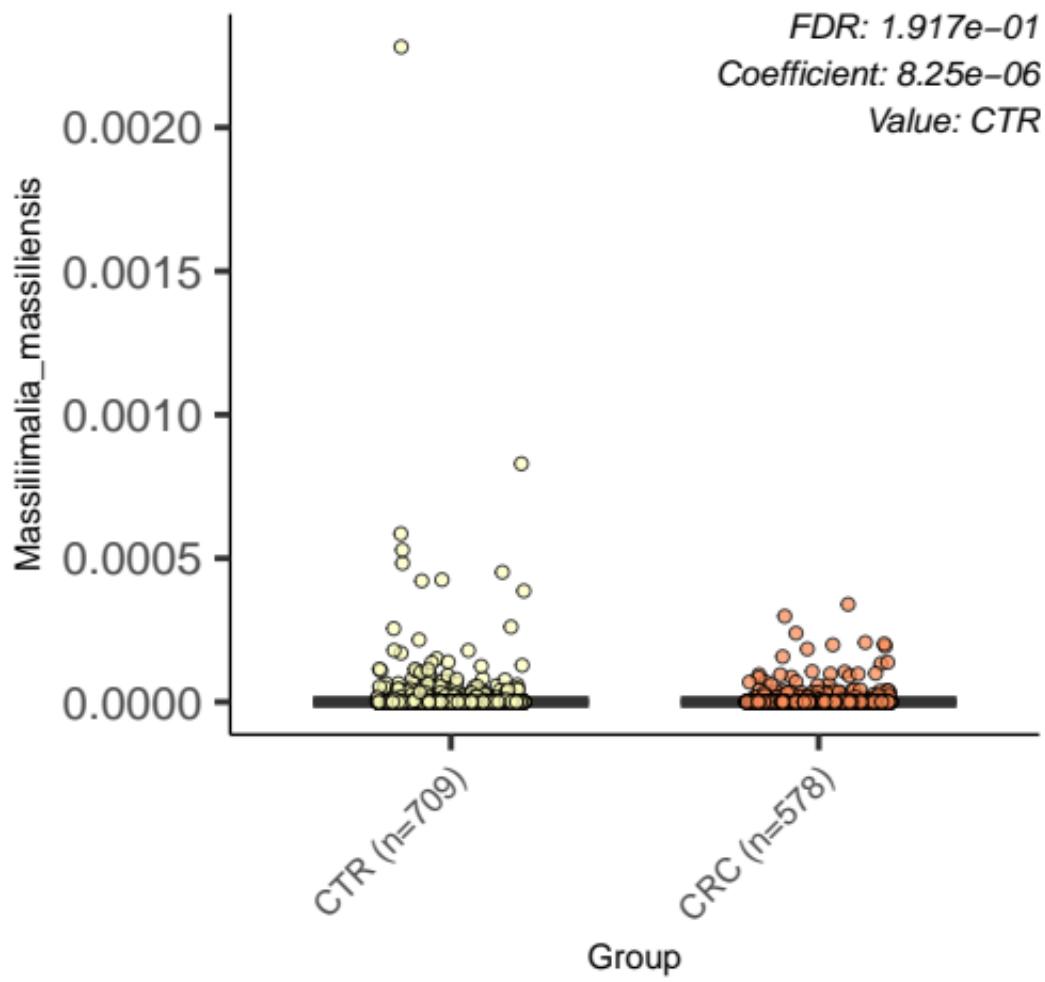


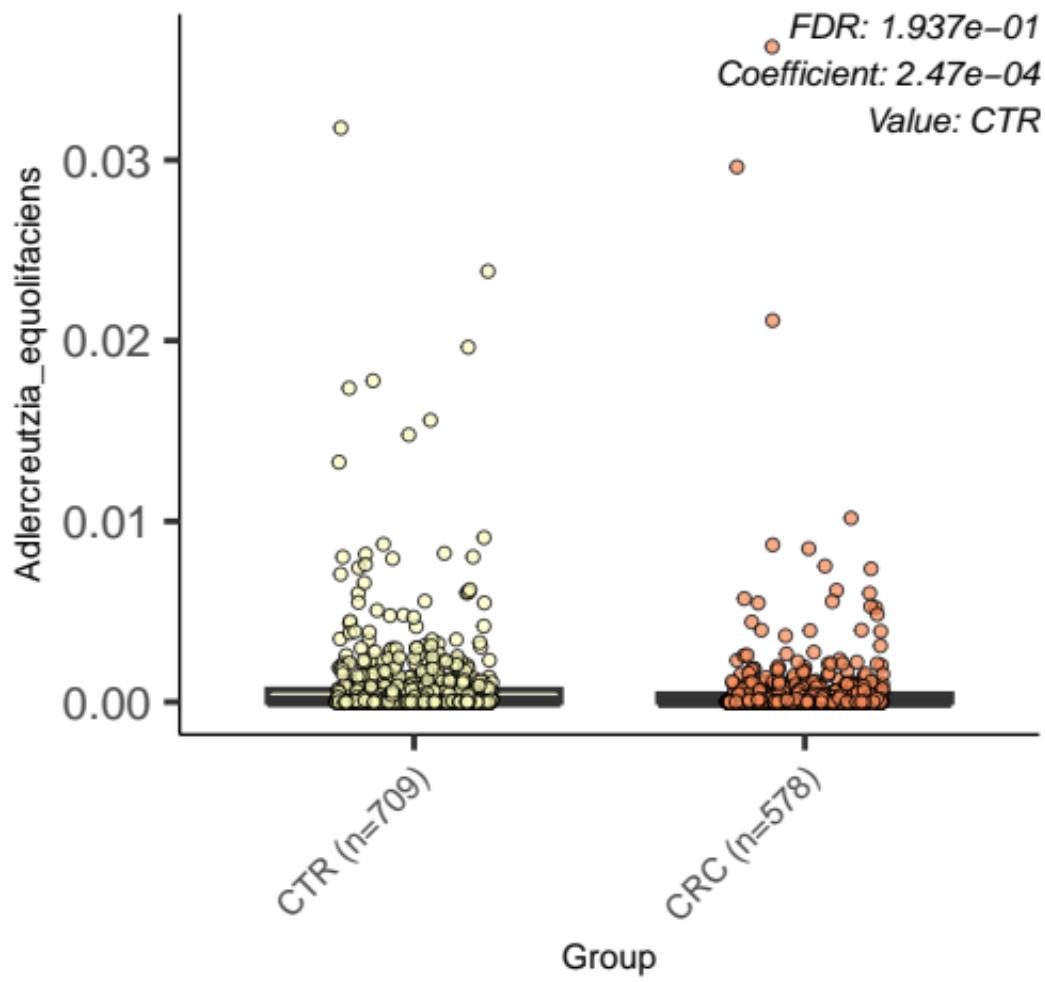


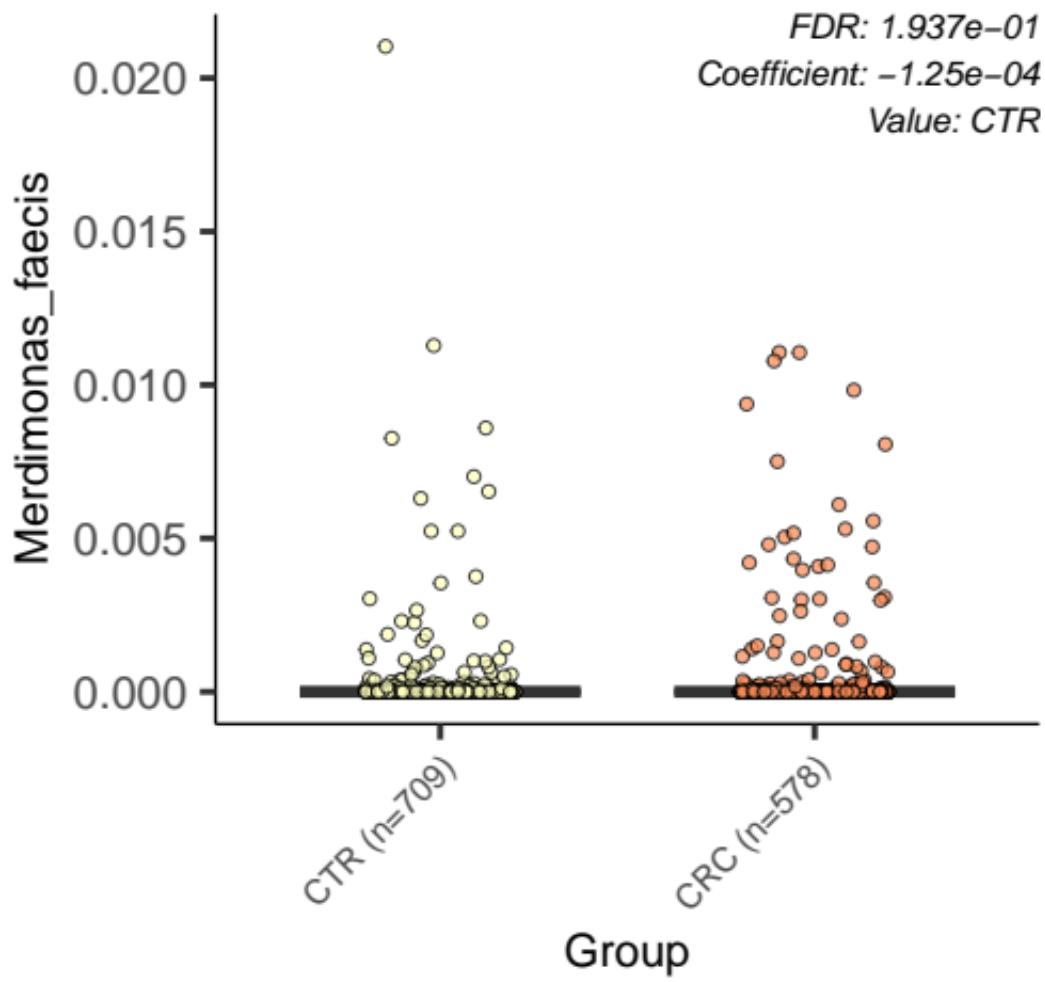












FDR: 1.973e-01
Coefficient: 9.00e-04
Value: CTR

Bifidobacterium_catenulatum

0.2

0.1

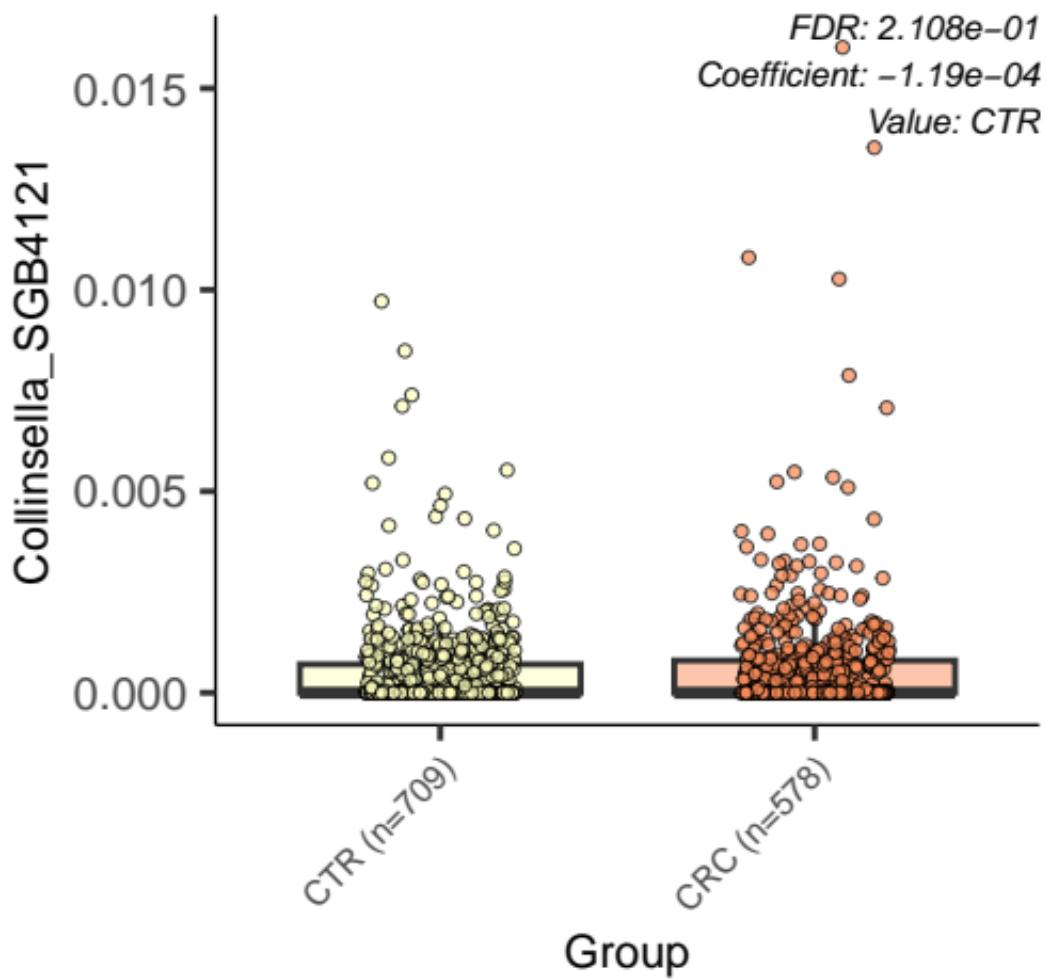
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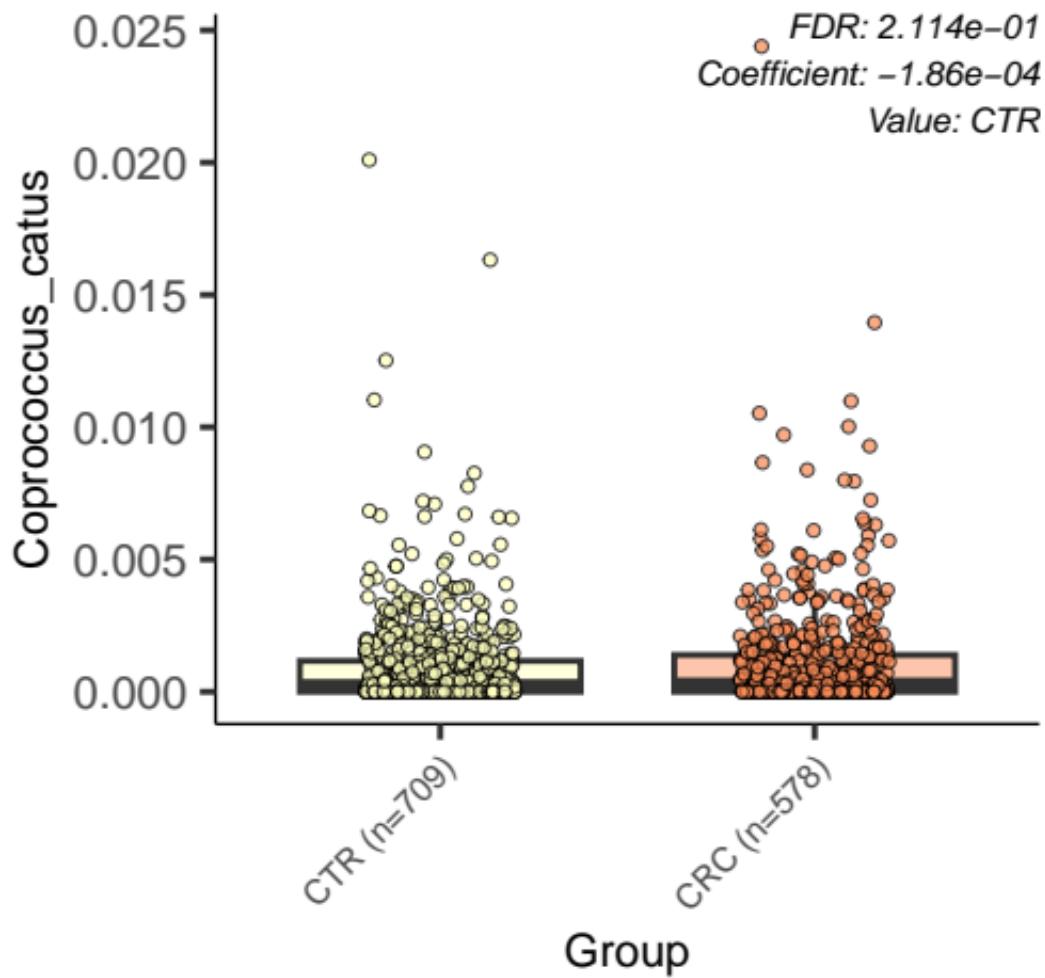
CTR (n=709)

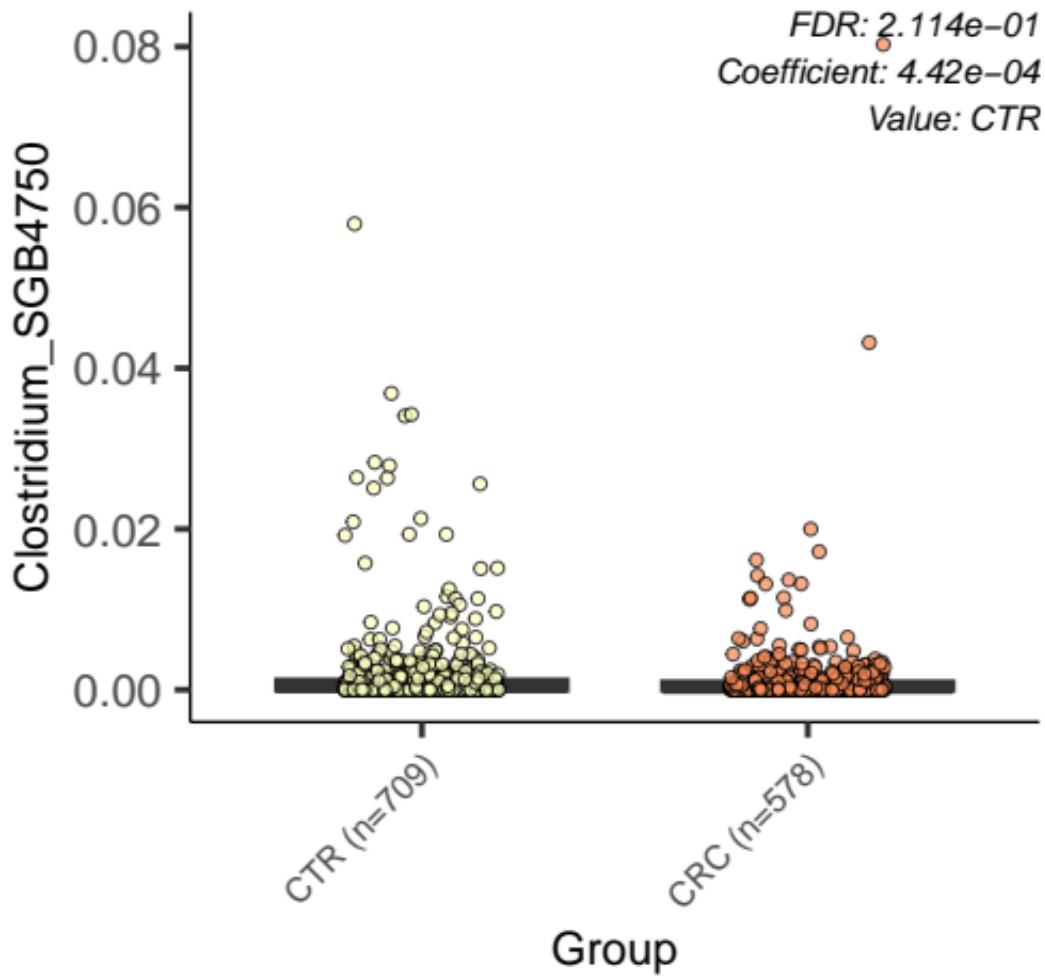
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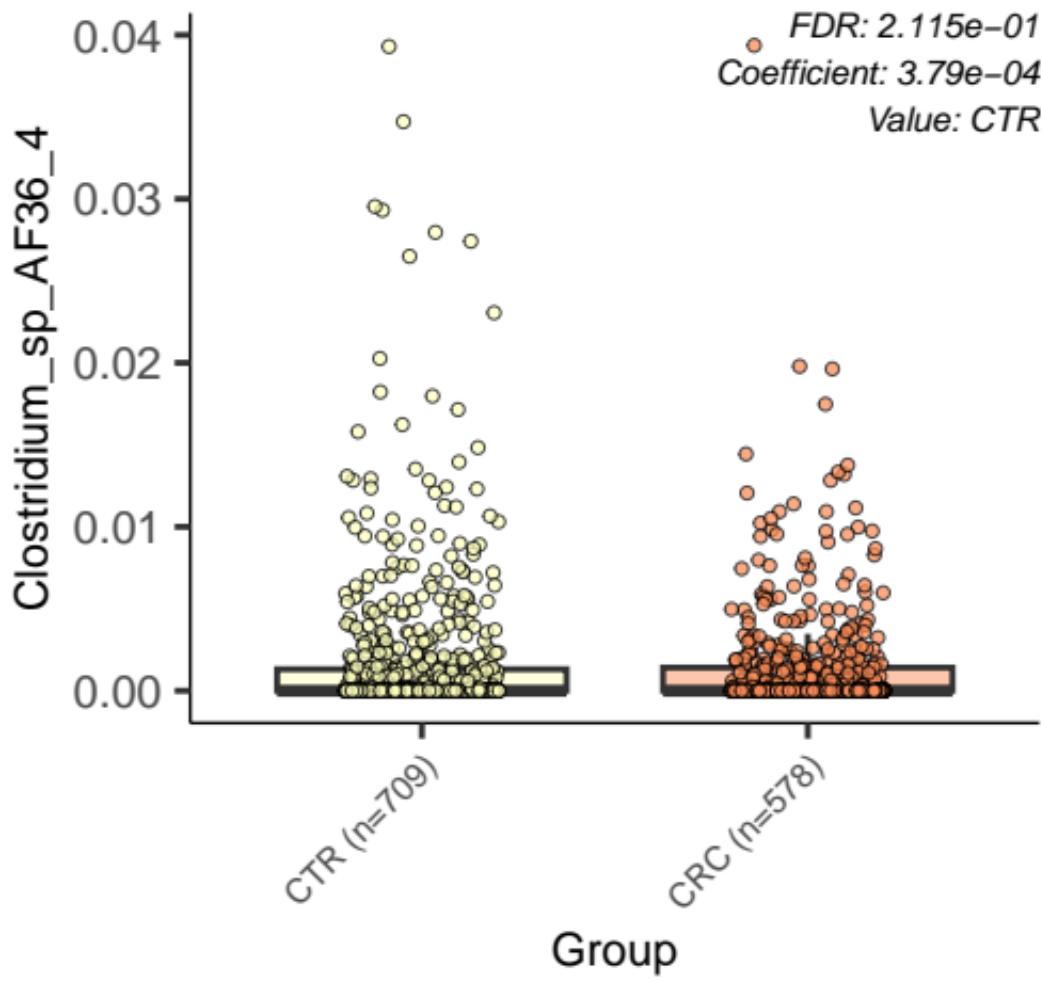
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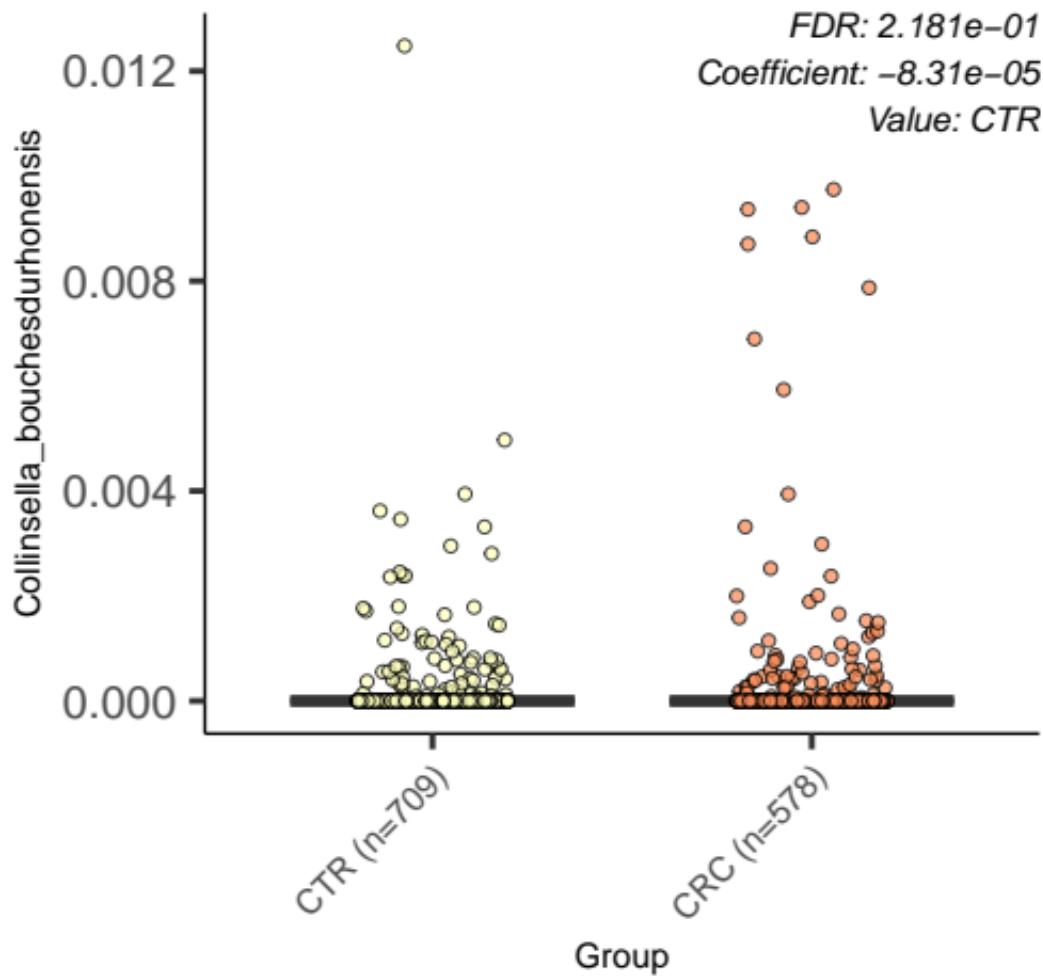


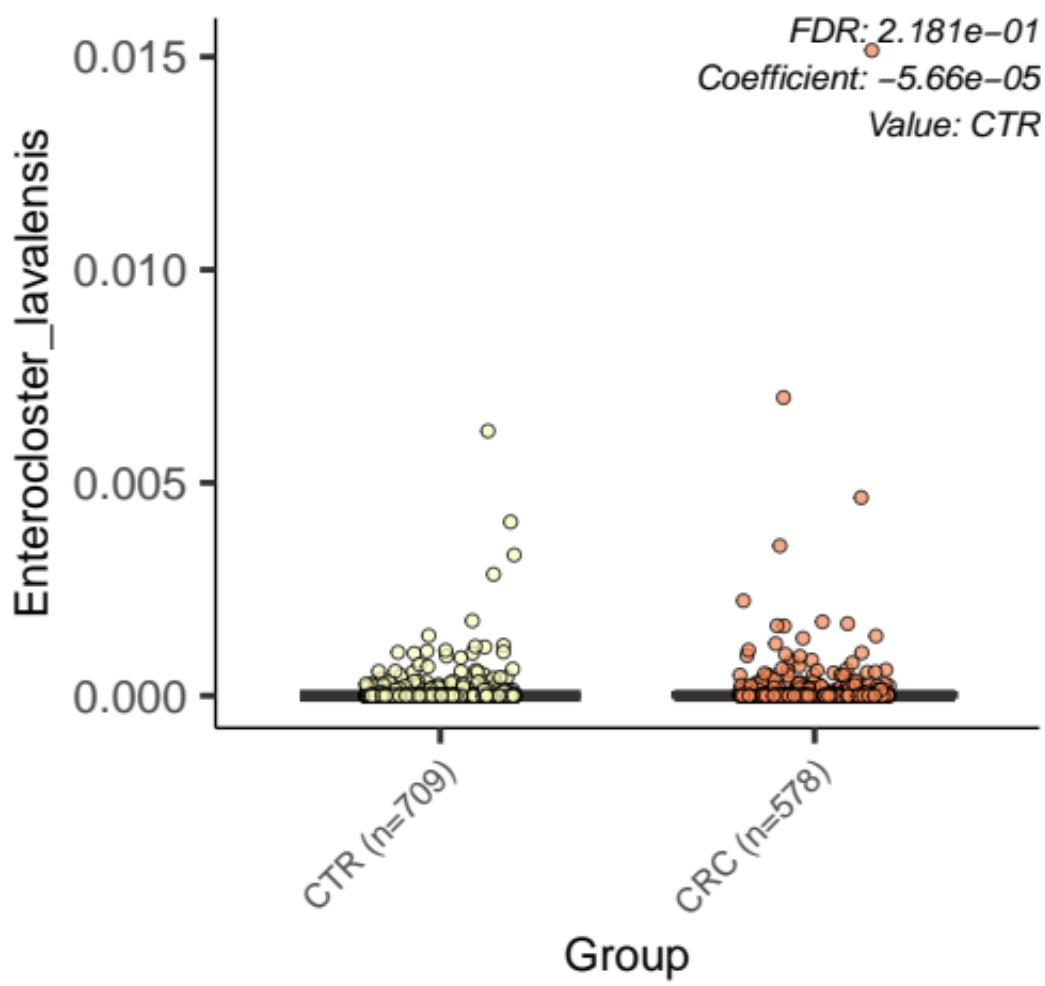


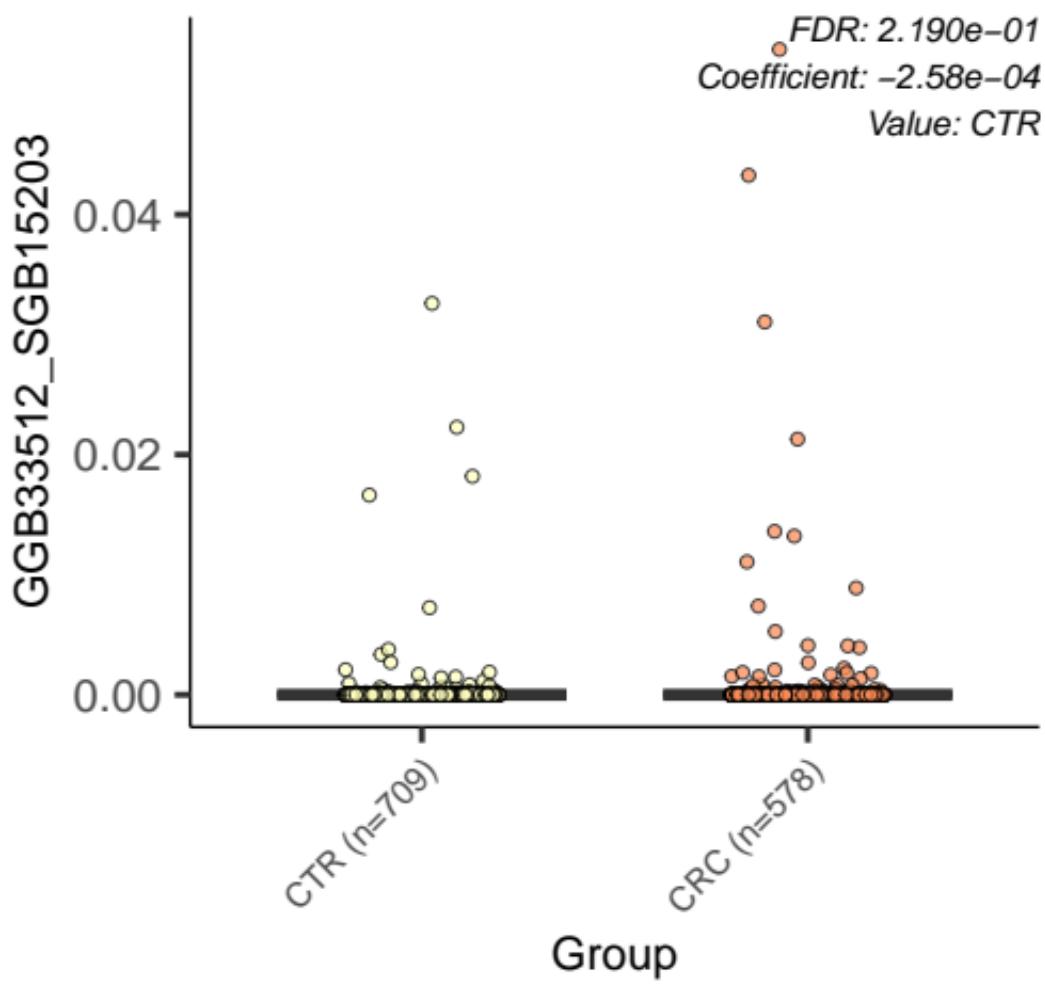


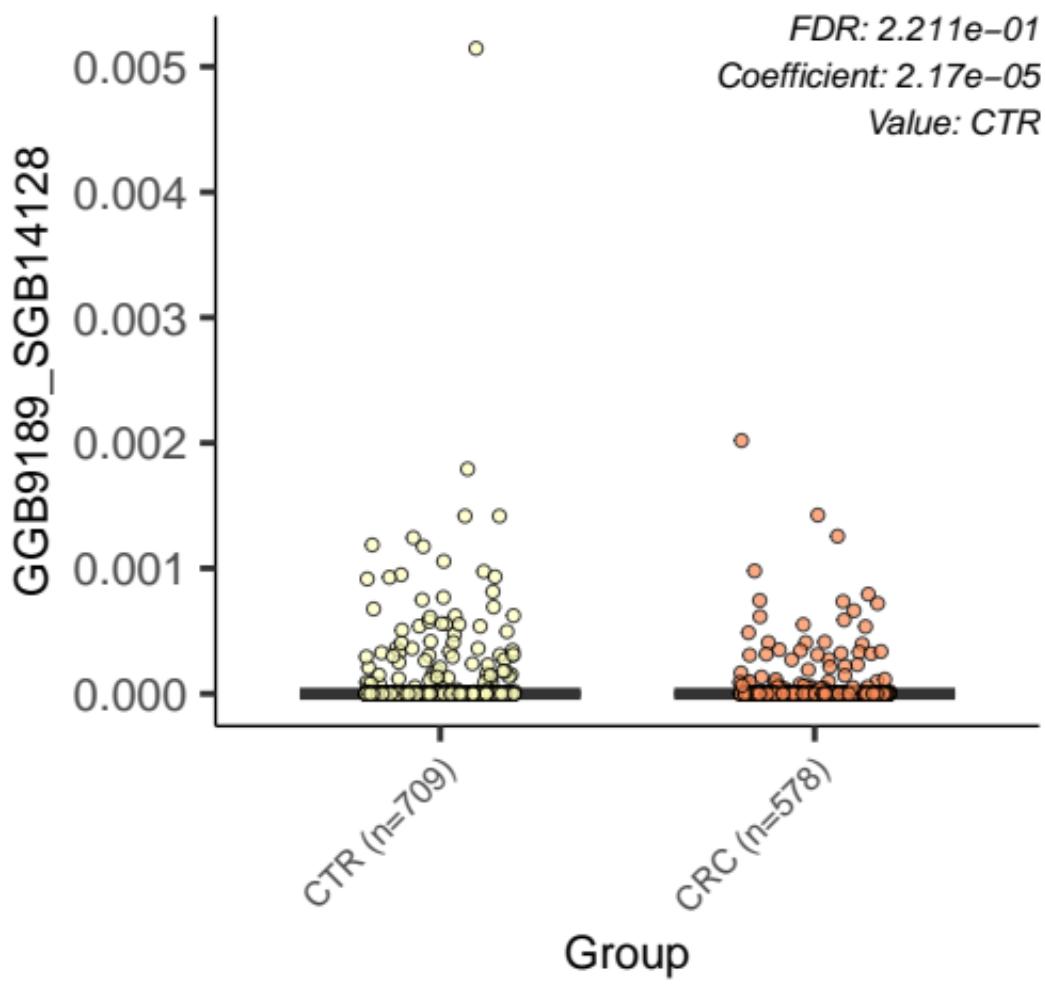


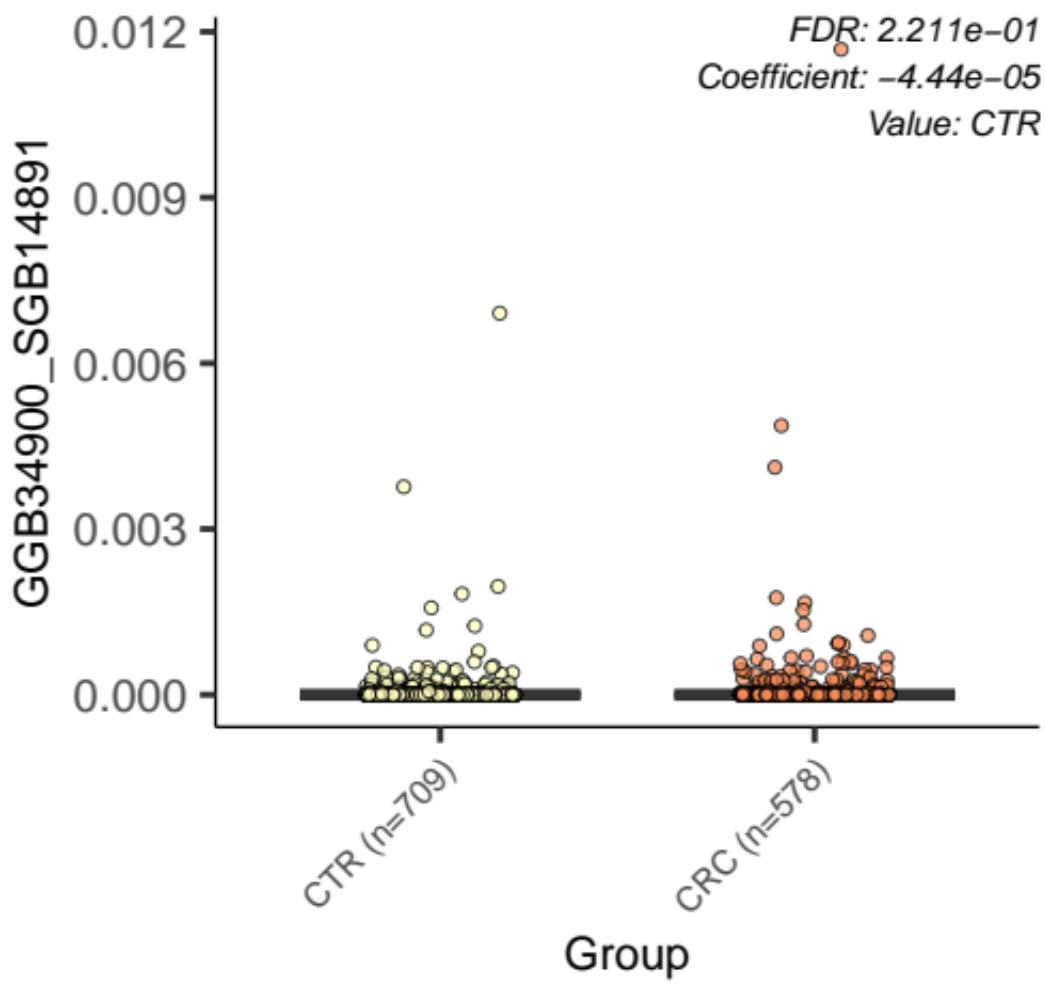


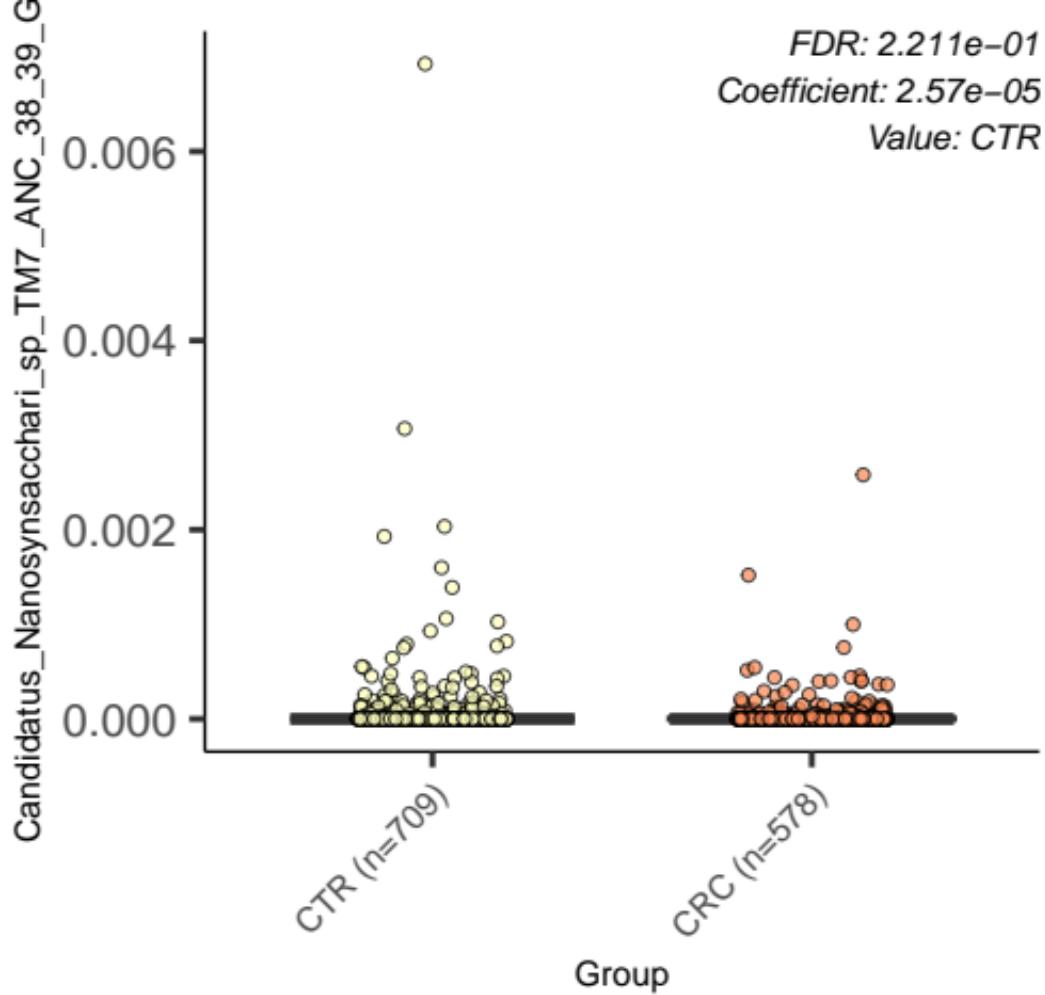












GGB3005_SGB3996

FDR: 2.211e-01
Coefficient: -1.73e-04
Value: CTR

0.02
0.01
0.00

CTR (n=709)

CRC (n=578)

Group

