

Peptostreptococcus

*FDR: 2.182e-04*

*Coefficient: -1.48e-03*

*Value: CTR*

0.03

0.02

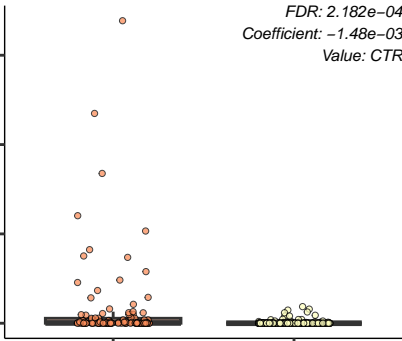
0.01

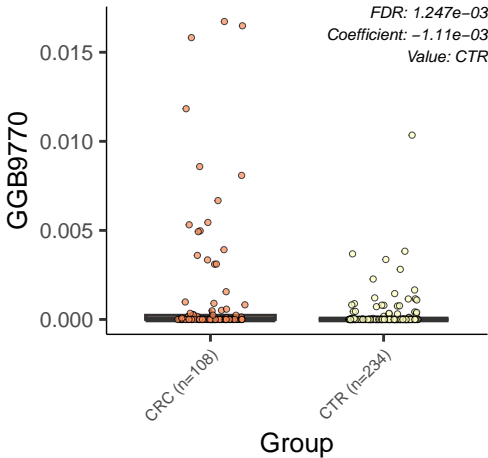
0.00

CRC (n=108)

CTR (n=234)

Group





Solobacterium

*FDR: 4.279e-03*

*Coefficient: -3.72e-04*

*Value: CTR*

0.0075

0.0050

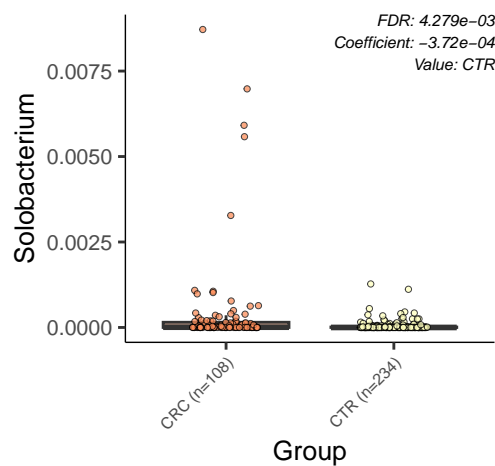
0.0025

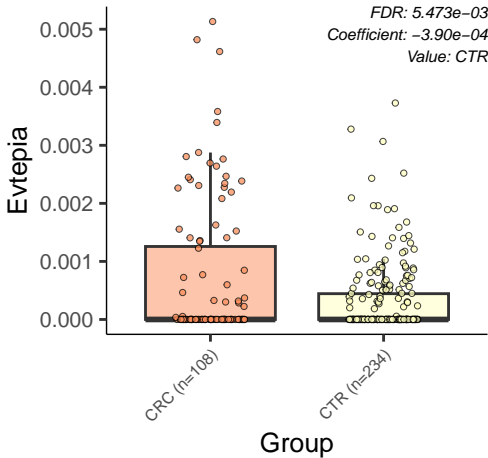
0.0000

CRC (n=108)

CTR (n=234)

Group





Anaerotignum

FDR:  $9.426e-03$

Coefficient:  $-2.36e-03$

Value: CTR

0.03

0.02

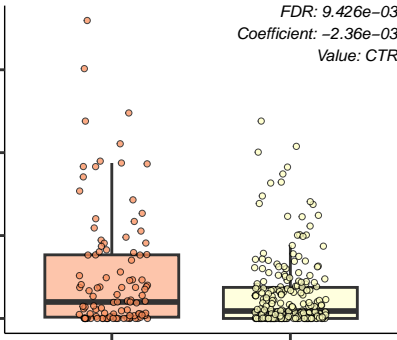
0.01

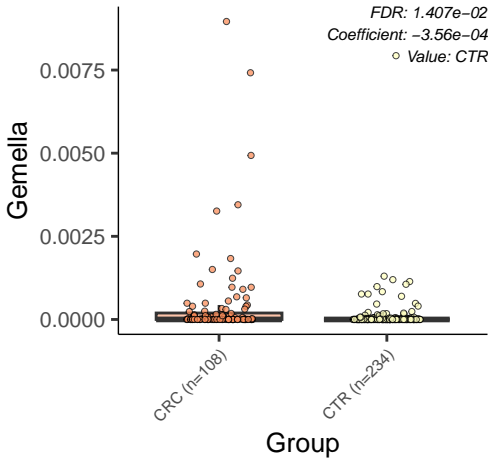
0.00

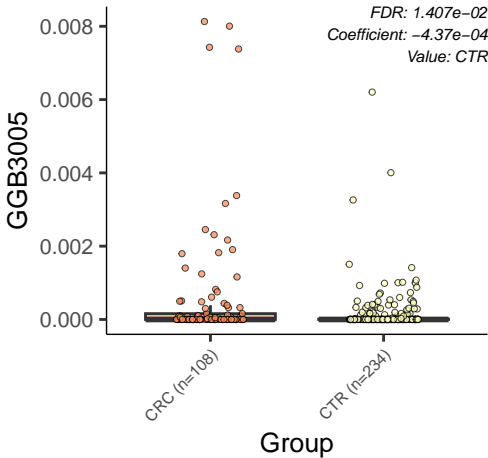
CRC (n=108)

CTR (n=234)

Group









Odoribacter

*FDR: 1.526e-02*

*Coefficient: -2.91e-03*

*Value: CTR*

0.075

0.050

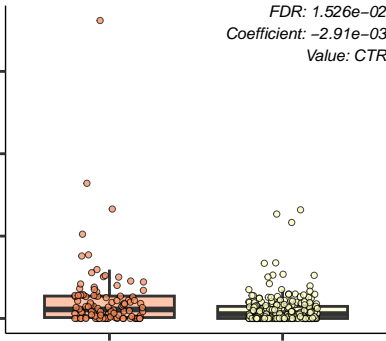
0.025

0.000

CRC (n=108)

CTR (n=234)

Group



Lachnospira

FDR: 1.770e-02  
Coefficient: 1.54e-02  
Value: CTR

0.2

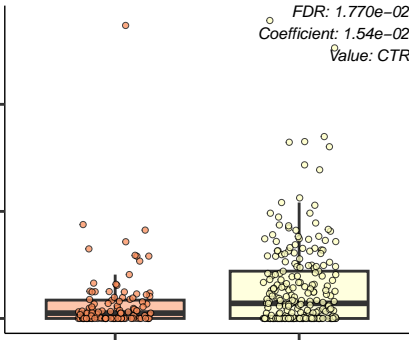
0.1

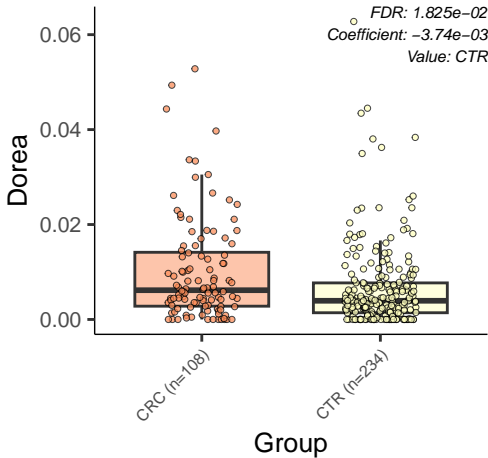
0.0

CRC (n=108)

CTR (n=234)

Group





Sanguibacter

*FDR: 2.109e-02*

*Coefficient: -3.33e-04*

*Value: CTR*

0.006

0.004

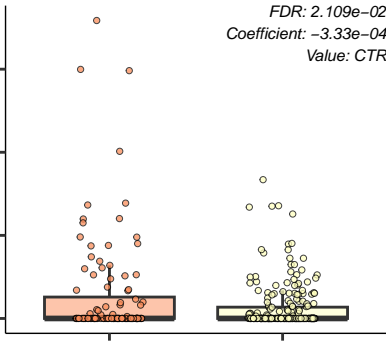
0.002

0.000

CRC (n=108)

CTR (n=234)

Group



GGB2998

*FDR: 4.499e-02*

*Coefficient: -1.70e-04*

*Value: CTR*

CRC (n=108)

CTR (n=234)

Group

0.004

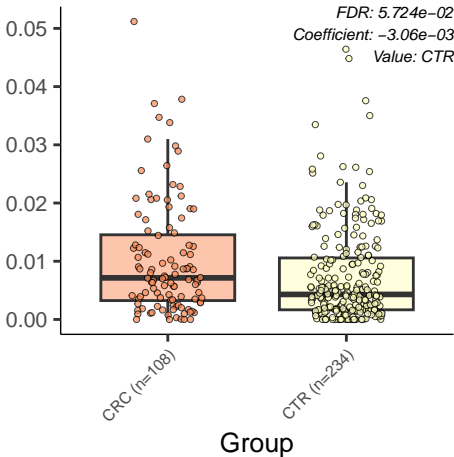
0.003

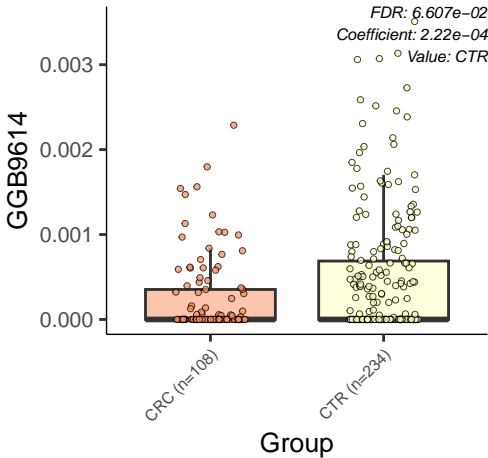
0.002

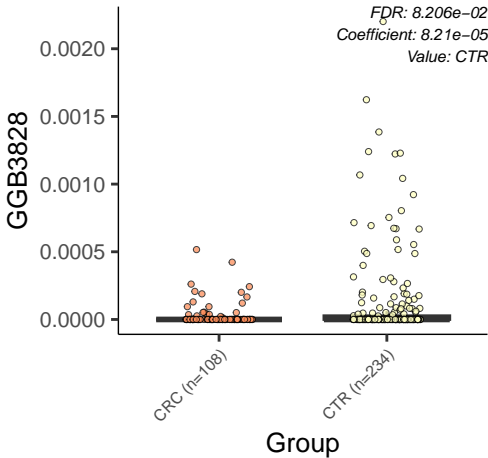
0.001

0.000

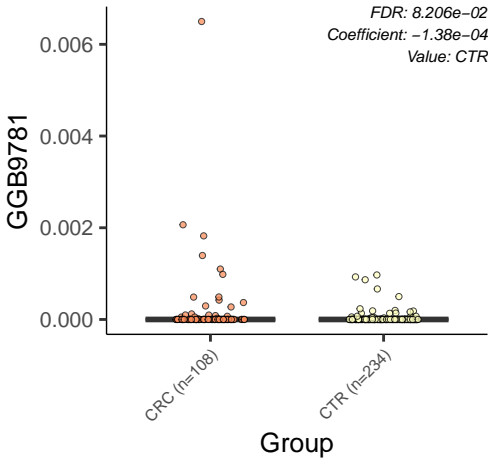
Collinsella











Cloacibacillus

*FDR: 8.206e-02*

*Coefficient: -4.08e-04*

*Value: CTR*

0.015

0.010

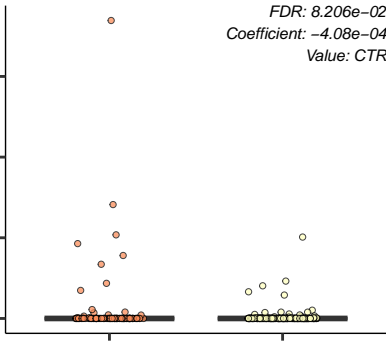
0.005

0.000

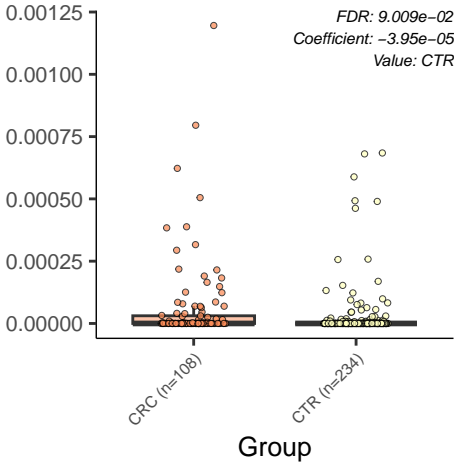
CRC (n=108)

CTR (n=234)

Group



Christensenella

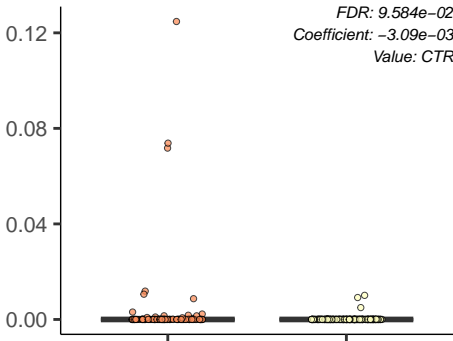




*FDR: 9.584e-02*

Coefficient:  $-3.09e-03$

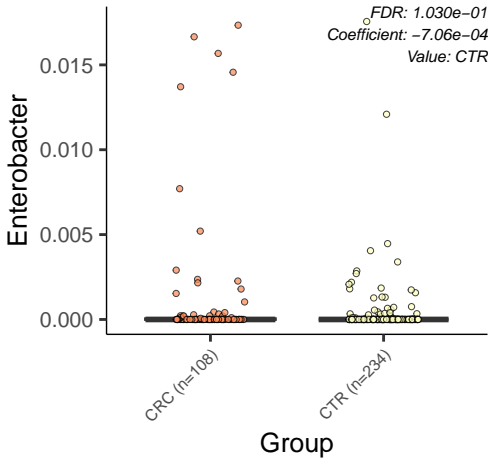
Value: CTR



CRC (n=108)

CTR (n=234)

Group



Candidatus\_Pararuminococcus

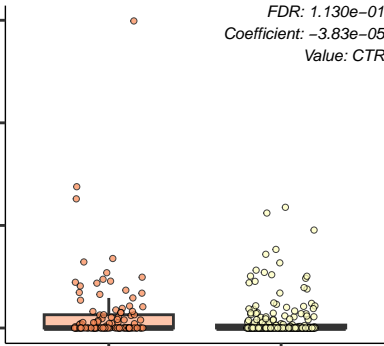
0.0015  
0.0010  
0.0005  
0.0000

*FDR: 1.130e-01*  
*Coefficient: -3.83e-05*  
*Value: CTR*

CRC (n=108)

CTR (n=234)

Group



Harryflintia

4e-04

2e-04

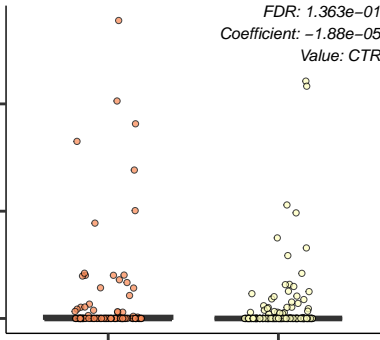
0e+00

CRC (n=108)

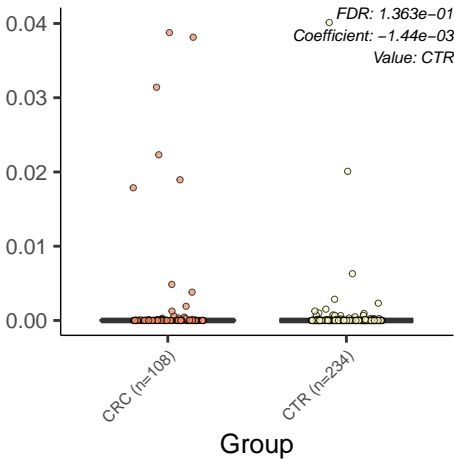
CTR (n=234)

Group

FDR: 1.363e-01  
Coefficient: -1.88e-05  
Value: CTR



GGB33512





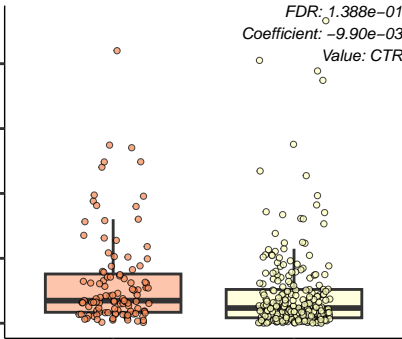
Mediterraneibacter

*FDR: 1.388e-01*  
*Coefficient: -9.90e-03*  
*Value: CTR*

CRC (n=108)

CTR (n=234)

Group



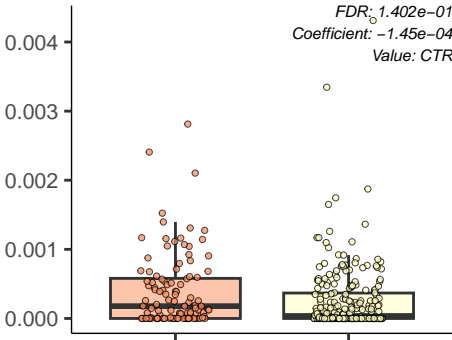
Rheinheimer

FDR:  $1.402e-01$   
Coefficient:  $-1.45e-04$   
Value: CTR

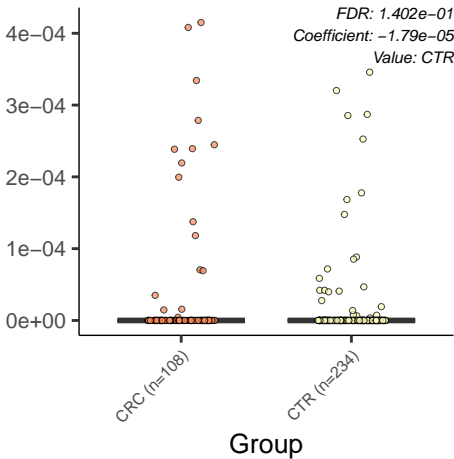
CRC (n=108)

CTR (n=234)

Group



Angelakisella



Faecalibacterium

*FDR: 1.589e-01*  
*Coefficient: 1.05e-02*  
*Value: CTR*

0.15

0.10

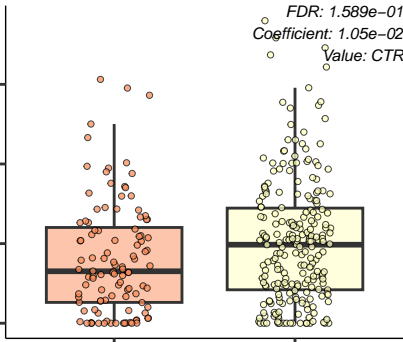
0.05

0.00

CRC (n=108)

CTR (n=234)

Group



Merdimonas

FDR: 1.589e-01  
Coefficient: -3.69e-04  
Value: CTR

0.0100

0.0075

0.0050

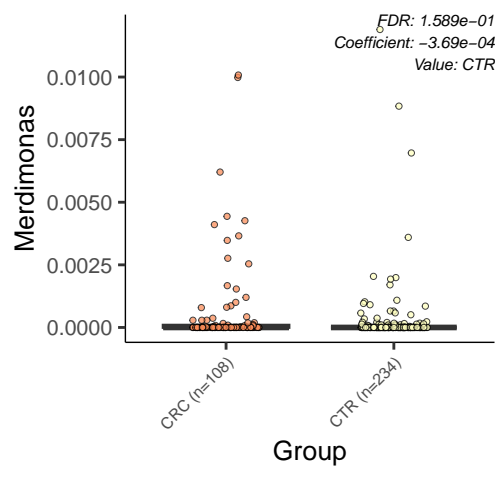
0.0025

0.0000

CRC (n=108)

CTR (n=234)

Group



GGB3034

*FDR: 1.589e-01*

*Coefficient: -4.95e-05*

*Value: CTR*

0.0020

0.0015

0.0010

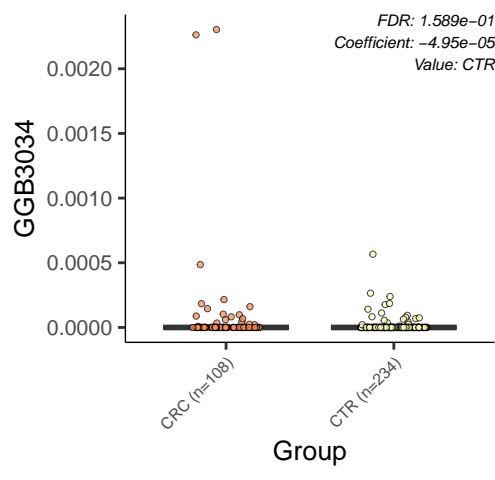
0.0005

0.0000

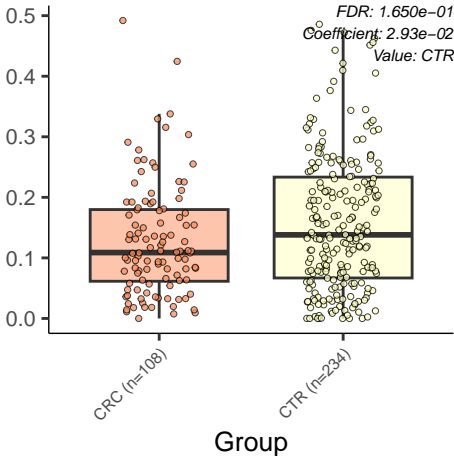
CRC (n=108)

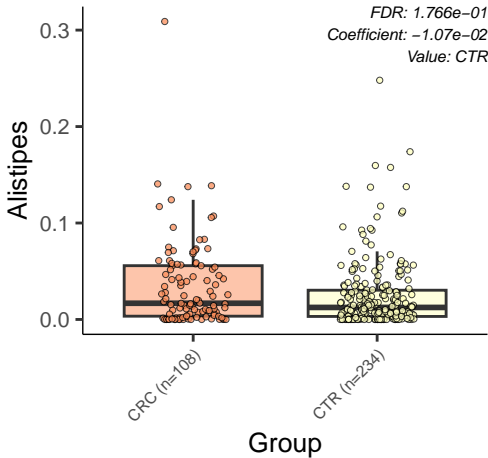
CTR (n=234)

Group



Phocaeicola







Anaerotruncus

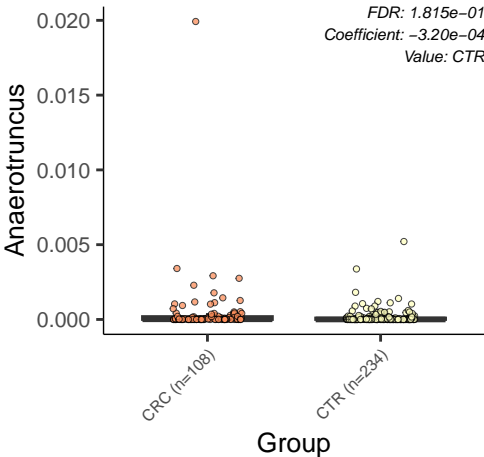
0.020  
0.015  
0.010  
0.005  
0.000

*FDR: 1.815e-01*  
*Coefficient: -3.20e-04*  
*Value: CTR*

CRC (n=108)

CTR (n=234)

Group



Candidatus\_Metaruminococcus

1e-03

5e-04

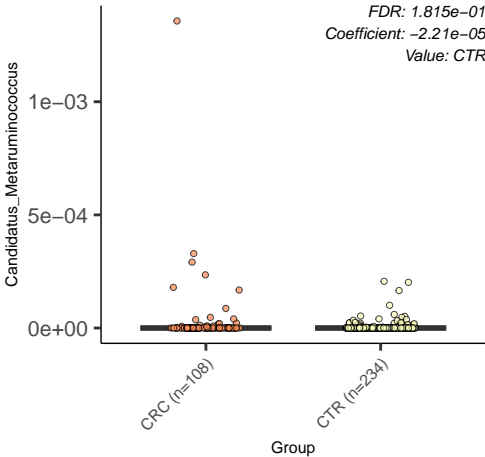
0e+00

CRC (n=108)

CTR (n=234)

Group

FDR: 1.815e-01  
Coefficient: -2.21e-05  
Value: CTR



GGB9619

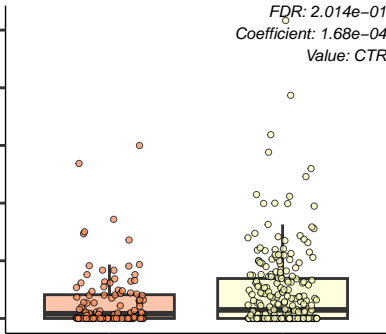
*FDR: 2.014e-01*  
*Coefficient: 1.68e-04*  
*Value: CTR*

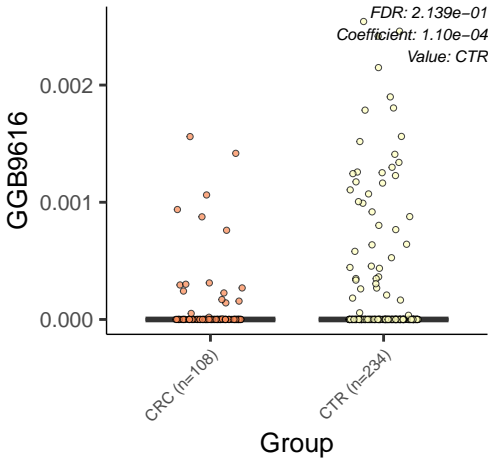
CRC (n=108)

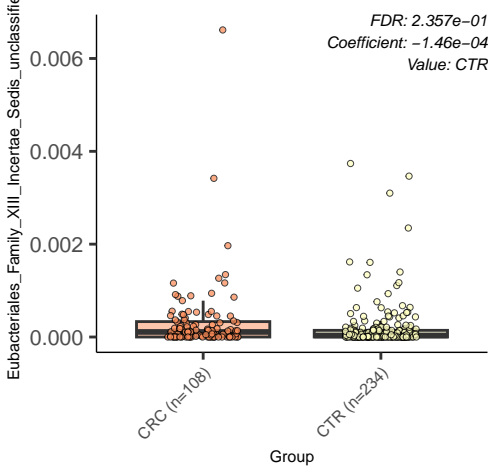
CTR (n=234)

Group

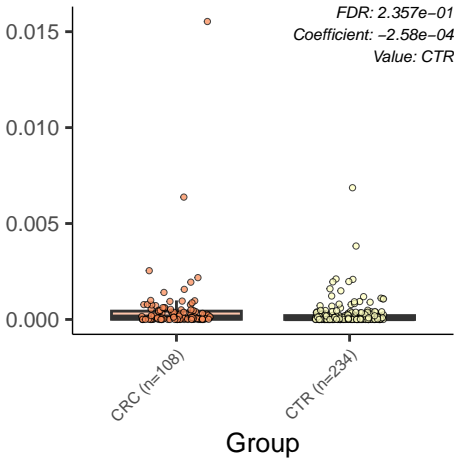
0.005  
0.004  
0.003  
0.002  
0.001  
0.000

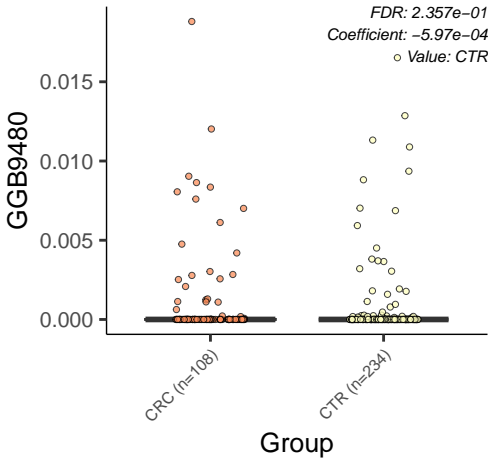






Mogibacterium





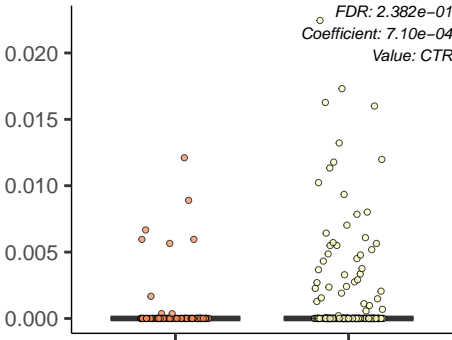
Catenibacterium

FDR: 2.382e-01  
Coefficient: 7.10e-04  
Value: CTR

CRC (n=108)

CTR (n=234)

Group





Clostridia\_unclassified

FDR: 2.423e-01  
Coefficient: -5.62e-04  
Value: CTR

0.015

0.010

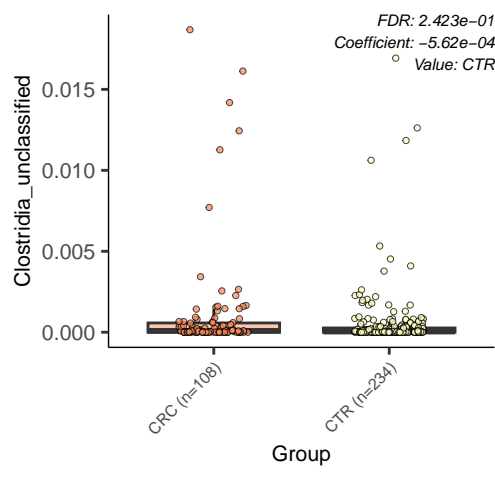
0.005

0.000

CRC (n=108)

CTR (n=234)

Group



GGB42994

*FDR: 2.427e-01*

*Coefficient: -5.96e-04*

*Value: CTR*

0.03

0.02

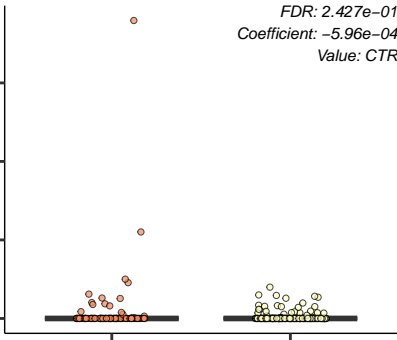
0.01

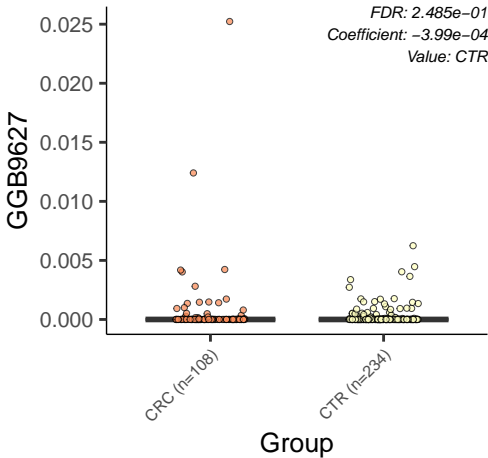
0.00

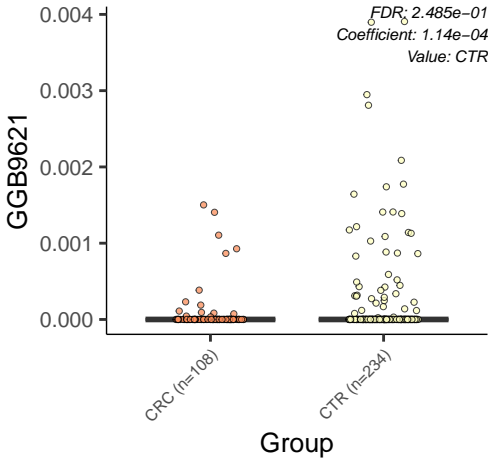
CRC (n=108)

CTR (n=234)

Group







Ruthenibacterium

FDR: 2.485e-01  
Coefficient: -7.47e-04  
Value: CTR

CRC (n=108)

CTR (n=234)

Group

