

Peptostreptococcus

*FDR: 2.239e-04*

*Coefficient: -1.41e-03*

*Value: CTR*

0.03

0.02

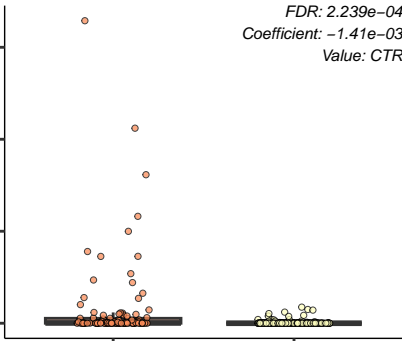
0.01

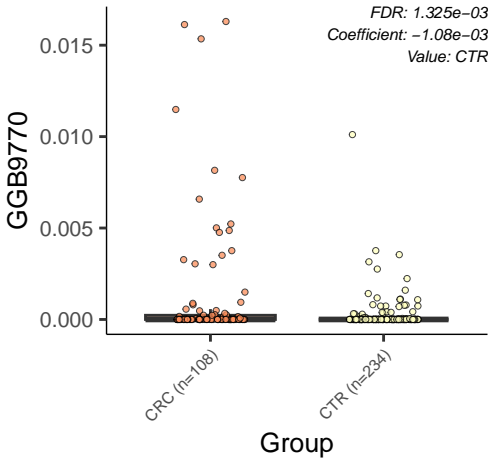
0.00

CRC (n=108)

CTR (n=234)

Group







*FDR: 4.383e-03*

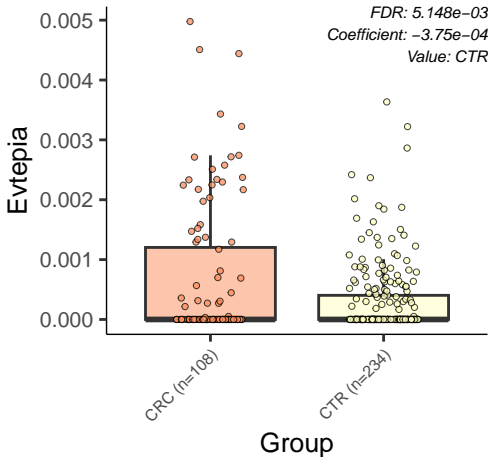
Coefficient:  $-3.59e-04$

Value: CTR

CRC (n=108)

CTR (n=234)

Group



Anaerotignum

FDR:  $7.489\text{e-}03$   
Coefficient:  $-2.31\text{e-}03$   
Value: CTR

0.03

0.02

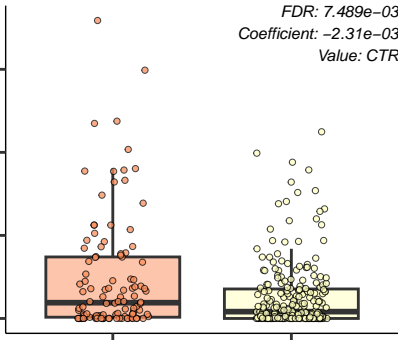
0.01

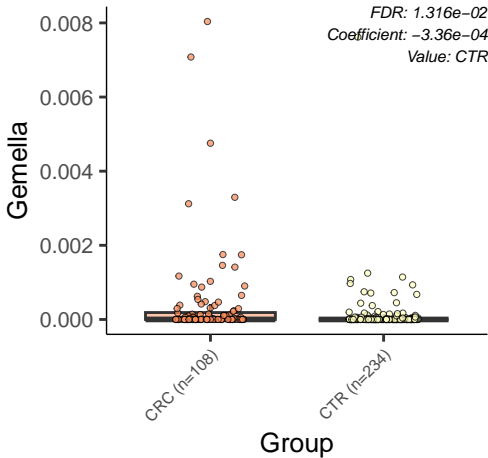
0.00

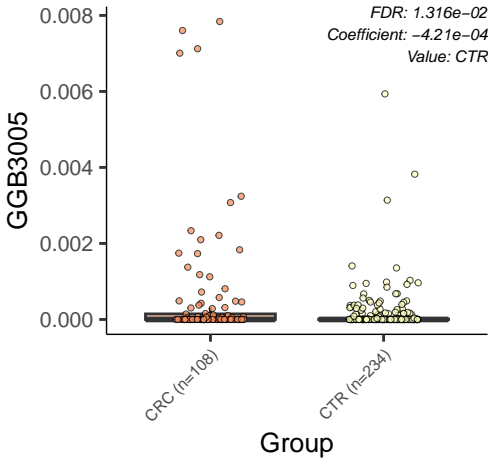
CRC (n=108)

CTR (n=234)

Group









Odoribacter

*FDR: 1.497e-02*

*Coefficient: -2.88e-03*

*Value: CTR*

0.075

0.050

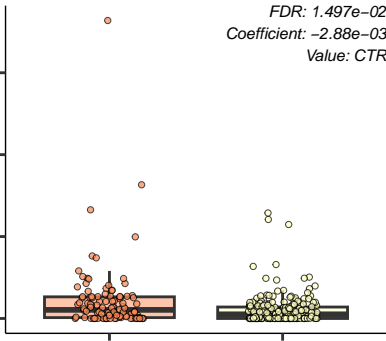
0.025

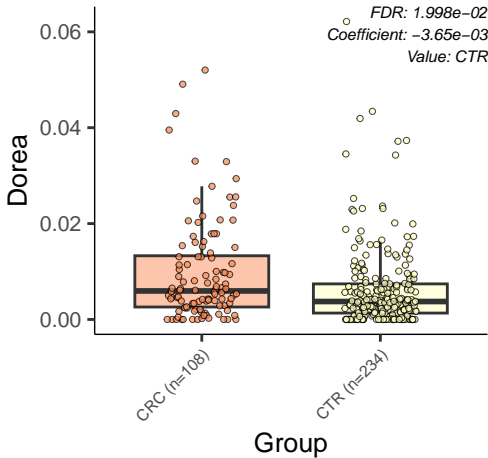
0.000

CRC (n=108)

CTR (n=234)

Group





Sanguibacter

*FDR: 2.267e-02*

*Coefficient: -3.24e-04*

*Value: CTR*

0.006

0.004

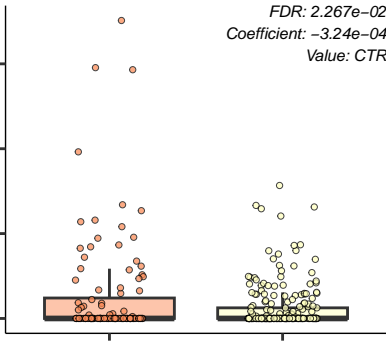
0.002

0.000

CRC (n=108)

CTR (n=234)

Group



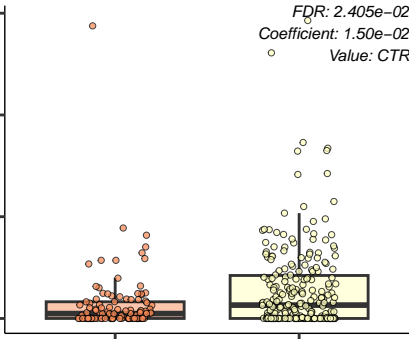
Lachnospira

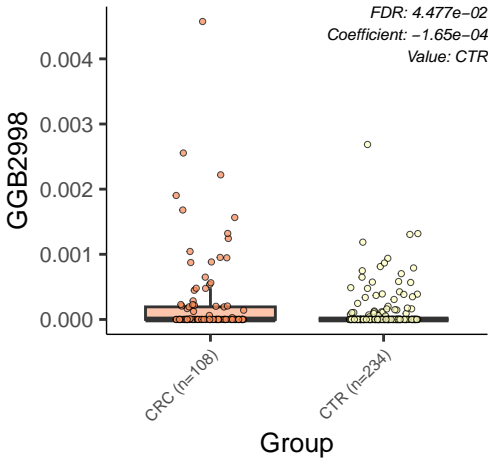
FDR:  $2.405e-02$   
Coefficient:  $1.50e-02$   
Value: CTR

CRC (n=108)

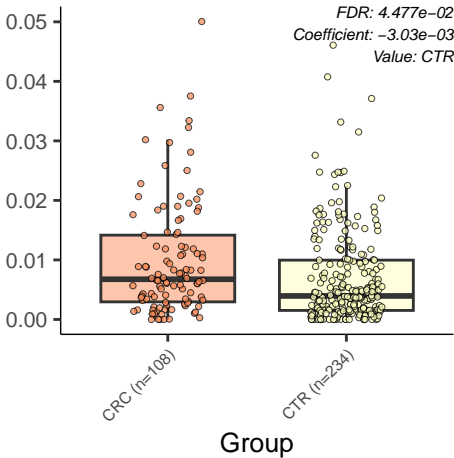
CTR (n=234)

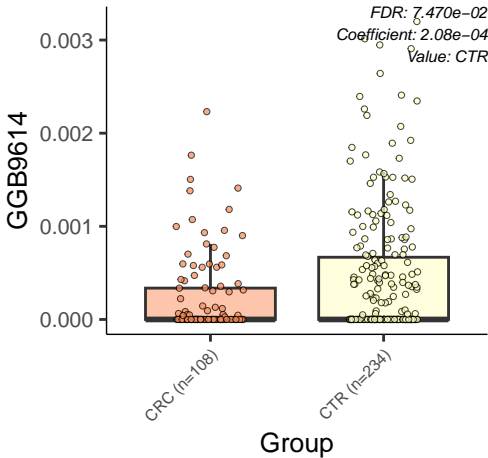
Group

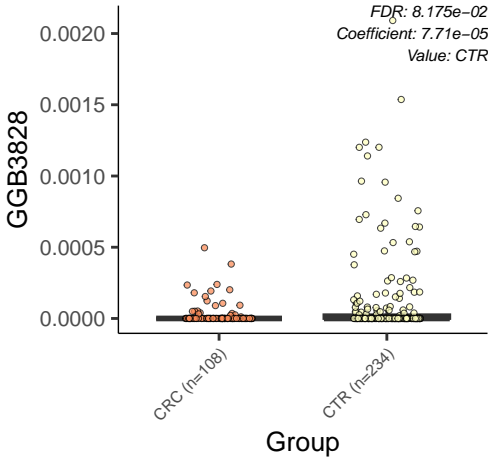




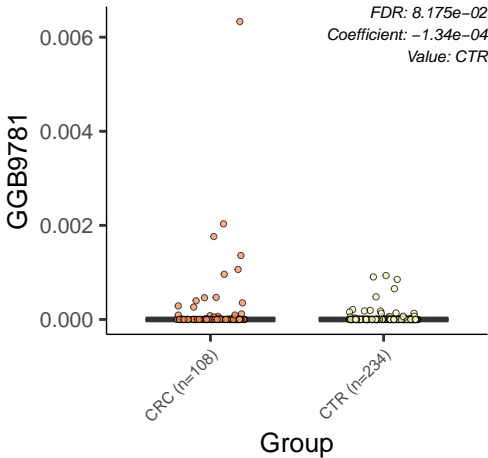
Collinsella

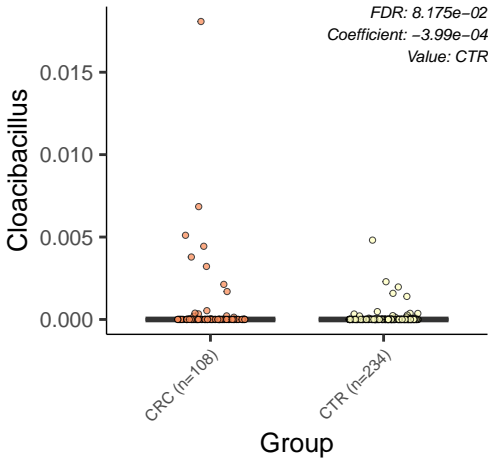












Christensenella

0.0012

0.0009

0.0006

0.0003

0.0000

*FDR: 8.880e-02*

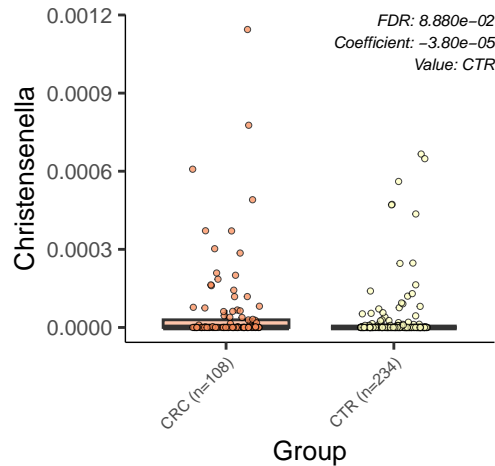
*Coefficient: -3.80e-05*

*Value: CTR*

CRC (n=108)

CTR (n=234)

Group



Porphyrromonas

FDR: 1.091e-01

Coefficient:  $-3.07e-03$

Value: CTR

0.10

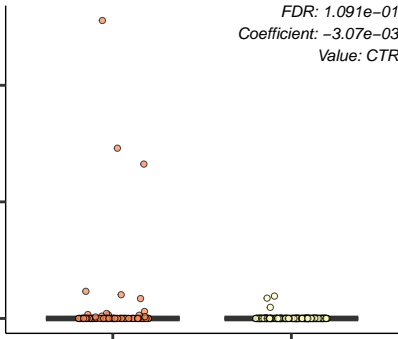
0.05

0.00

CRC (n=108)

CTR (n=234)

Group



Candidatus\_Pararuminococcus

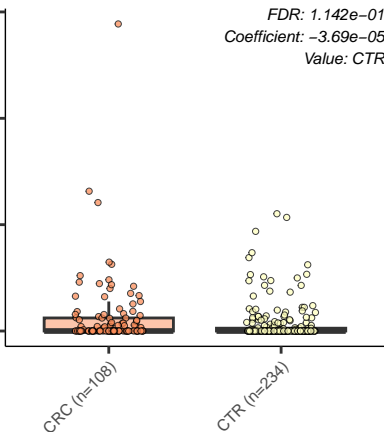
0.0015  
0.0010  
0.0005  
0.0000

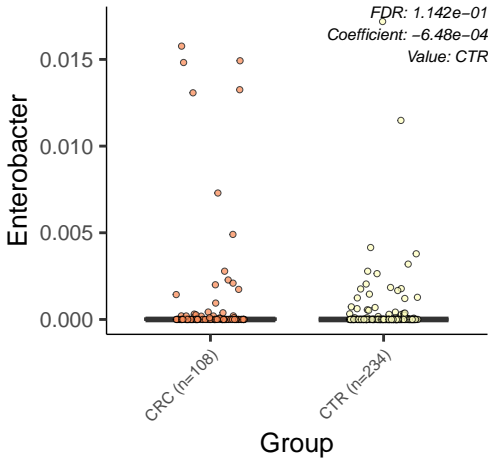
*FDR: 1.142e-01*  
*Coefficient: -3.69e-05*  
*Value: CTR*

CRC (n=108)

CTR (n=234)

Group





Rheinheimer

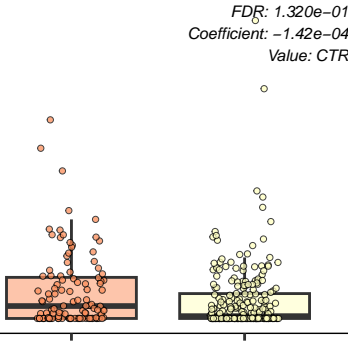
0.004  
0.003  
0.002  
0.001  
0.000

CRC (n=108)

CTR (n=234)

Group

FDR: 1.320e-01  
Coefficient: -1.42e-04  
Value: CTR



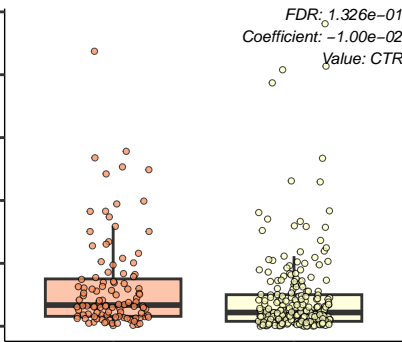
Mediterraneibacter

*FDR: 1.326e-01*  
*Coefficient: -1.00e-02*  
*Value: CTR*

CRC (n=108)

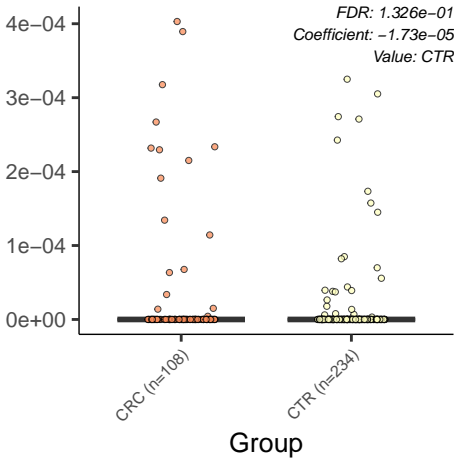
CTR (n=234)

Group





Angelakisella



Harryflintia

*FDR: 1.326e-01*  
*Coefficient: -1.81e-05*  
*Value: CTR*

4e-04

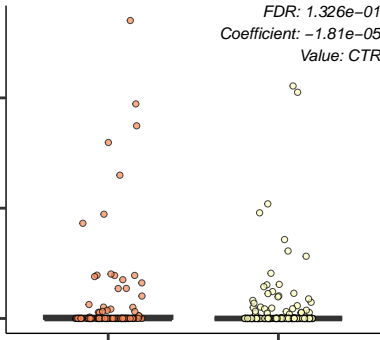
2e-04

0e+00

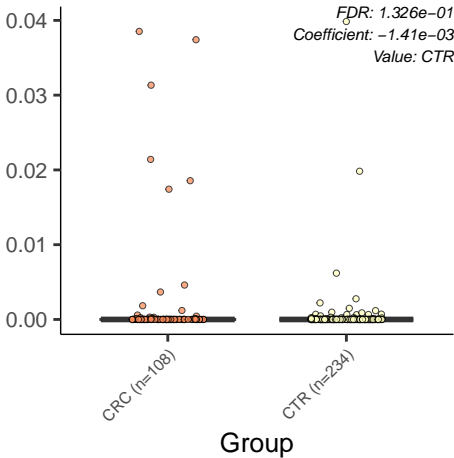
CRC (n=108)

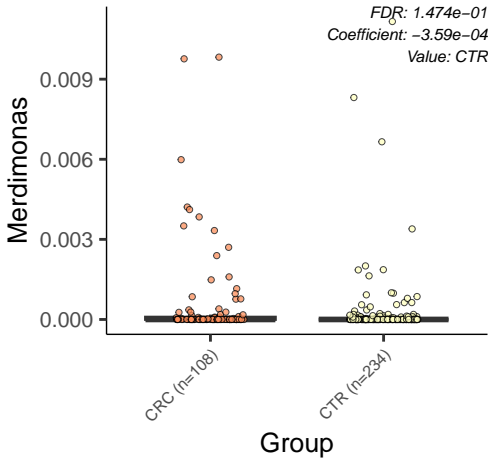
CTR (n=234)

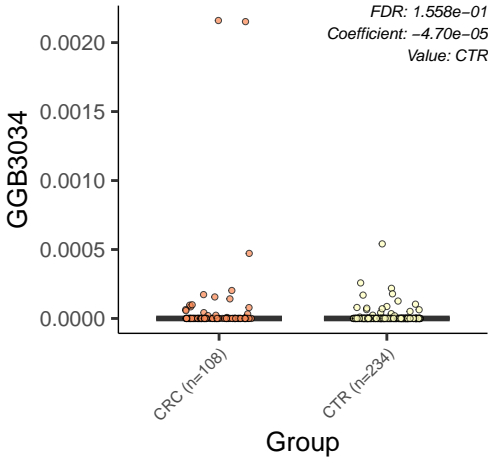
Group



GGB33512







Phocaeicola

0.4

0.2

0.0

CRC (n=108)

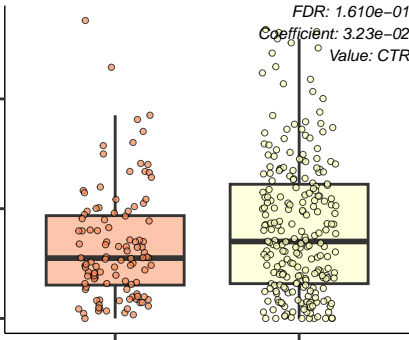
CTR (n=234)

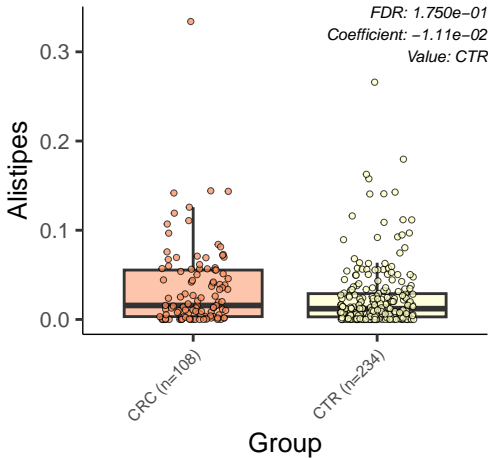
Group

FDR: 1.610e-01

Coefficient: 3.23e-02

Value: CTR





Faecalibacterium

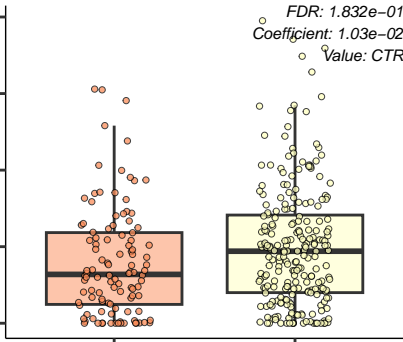
0.20  
0.15  
0.10  
0.05  
0.00

CRC (n=108)

CTR (n=234)

Group

FDR: 1.832e-01  
Coefficient: 1.03e-02  
Value: CTR





Candidatus\_Metaruminococcus

1e-03

5e-04

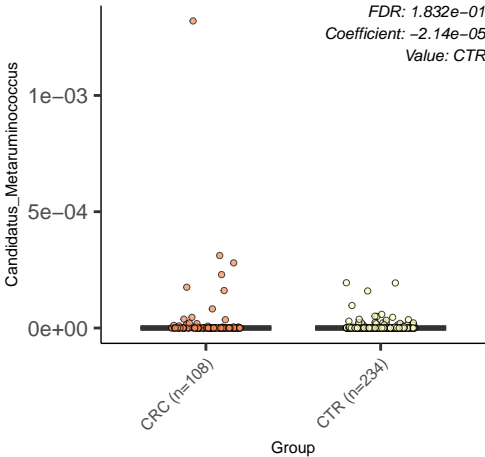
0e+00

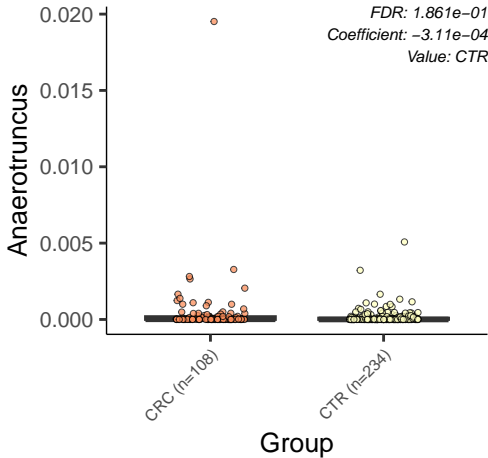
CRC (n=108)

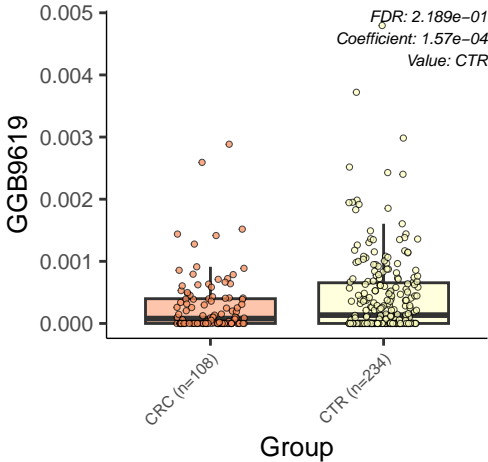
CTR (n=234)

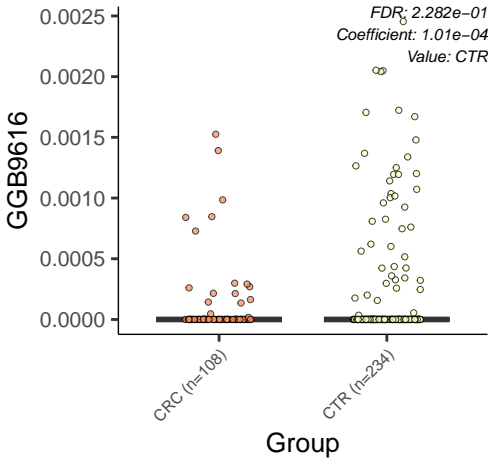
Group

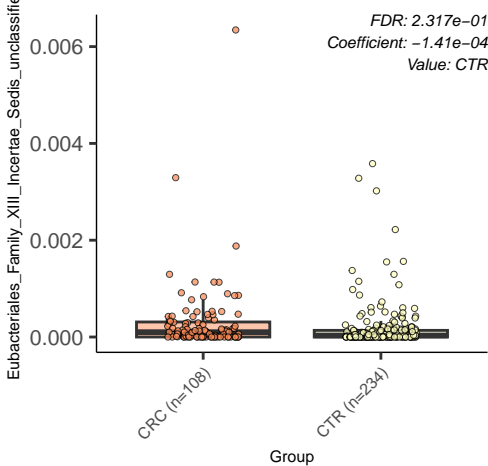
FDR: 1.832e-01  
Coefficient: -2.14e-05  
Value: CTR



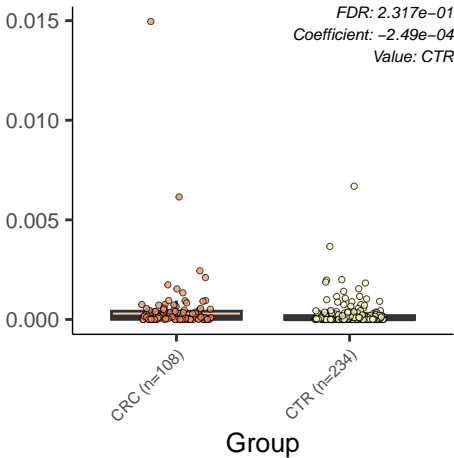








Mogibacterium



Clostridia\_unclassified

FDR: 2.317e-01  
Coefficient: -5.49e-04  
Value: CTR

0.015  
0.010  
0.005  
0.000

CRC (n=108)

CTR (n=234)

Group

