

Lachnoclostridium

FDR: 7.752e-02

Coefficient: -1.95e-03

Value: CTR

0.02

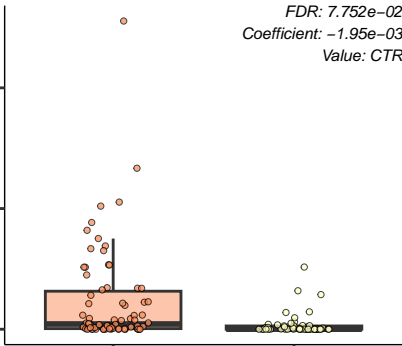
0.01

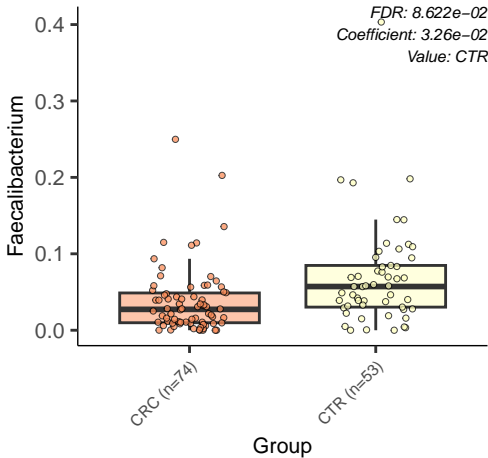
0.00

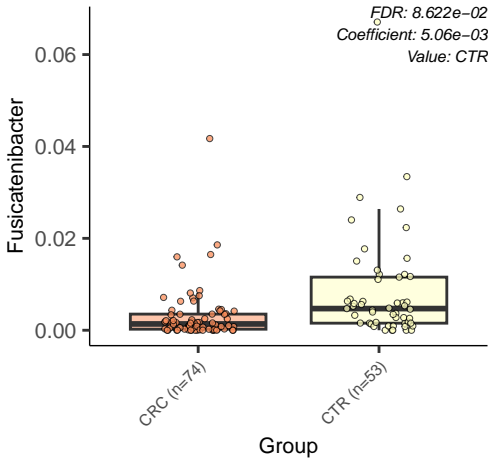
CRC (n=74)

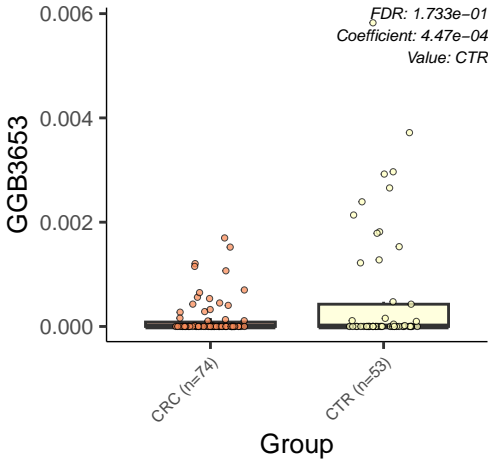
CTR (n=53)

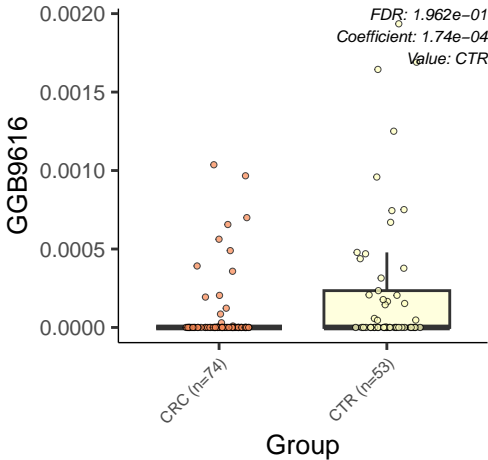
Group











Erysipelatoclostridium

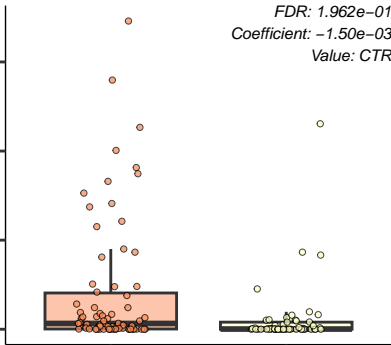
0.015
0.010
0.005
0.000

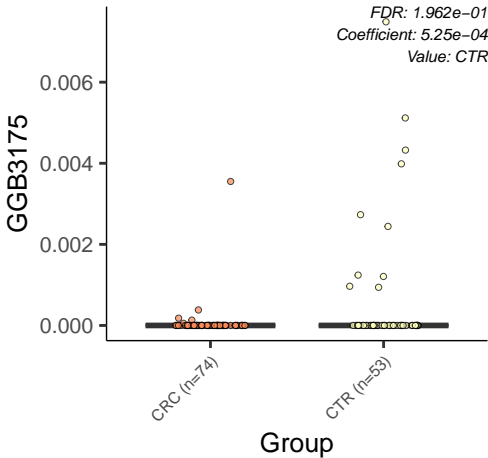
CRC (n=74)

CTR (n=53)

Group

FDR: 1.962e-01
Coefficient: -1.50e-03
Value: CTR





Mediterraneibacter

0.20

0.15

0.10

0.05

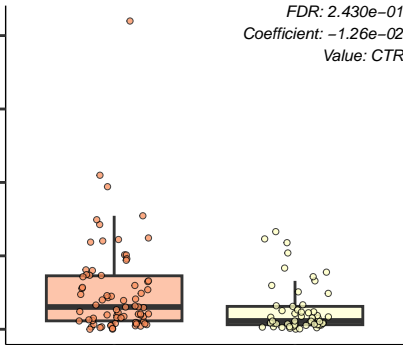
0.00

CRC (n=74)

CTR (n=53)

Group

FDR: 2.430e-01
Coefficient: -1.26e-02
Value: CTR



Mogibacterium

FDR: 2.430e-01

Coefficient: -2.65e-04

Value: CTR

0.004

0.003

0.002

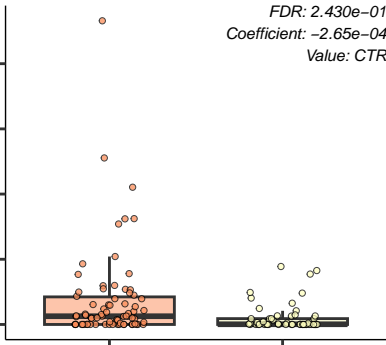
0.001

0.000

CRC (n=74)

CTR (n=53)

Group



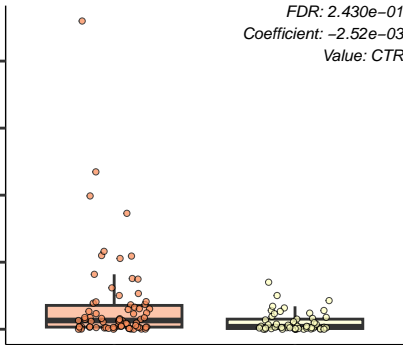
Ruthenibacterium

FDR: 2.430e-01
Coefficient: -2.52e-03
Value: CTR

CRC (n=74)

CTR (n=53)

Group



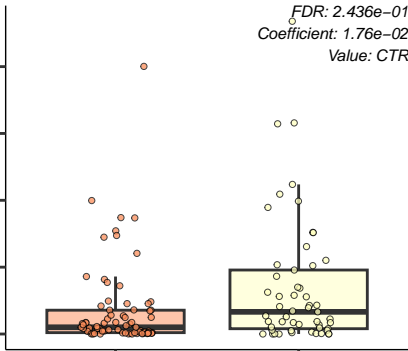
Lachnospiraceae_unclassified

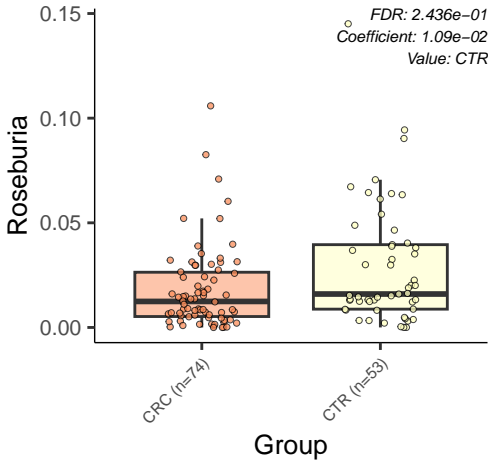
FDR: 2.436e-01
Coefficient: 1.76e-02
Value: CTR

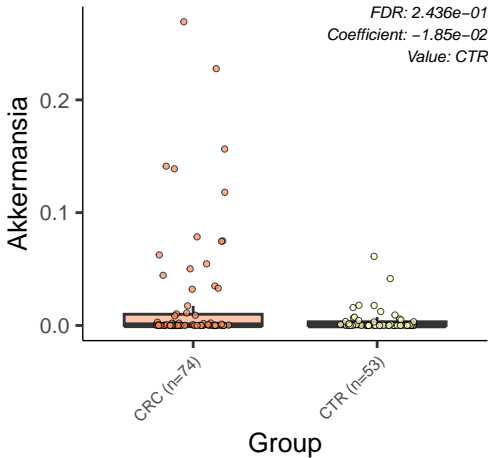
CRC (n=74)

CTR (n=53)

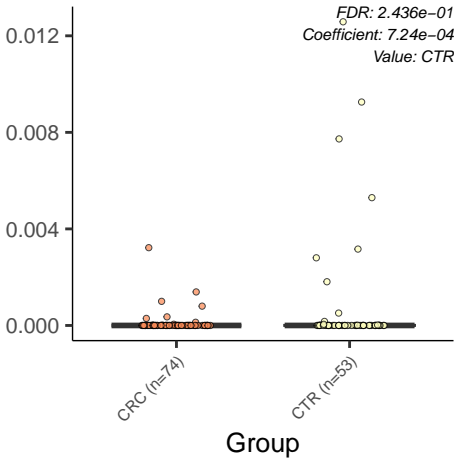
Group







Corynebacterium



Firmicutes_unclassified

FDR: 2.436e-01
Coefficient: 3.91e-03
Value: CTR

0.075

0.050

0.025

0.000

CRC (n=74)

CTR (n=53)

Group

