

Parabacteroides

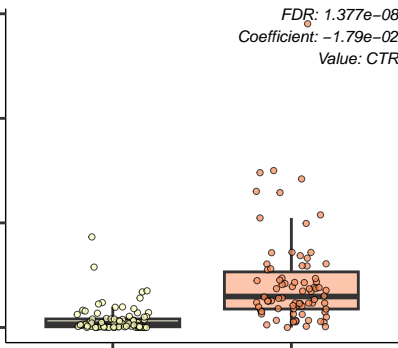
0.15
0.10
0.05
0.00

FDR: 1.377e-08
Coefficient: -1.79e-02
Value: CTR

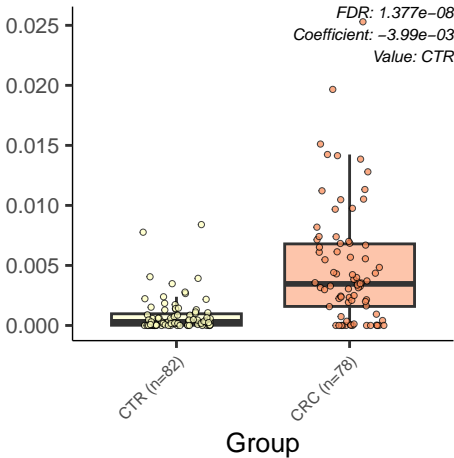
CTR (n=82)

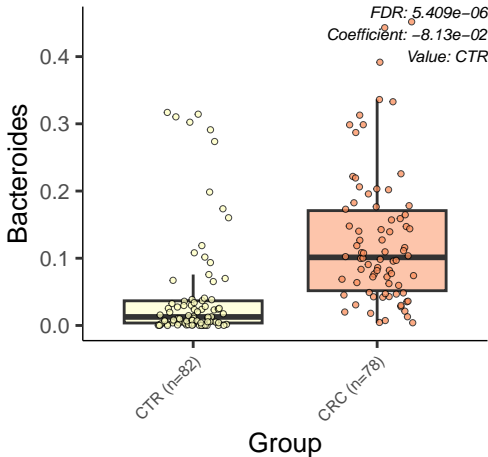
CRC (n=78)

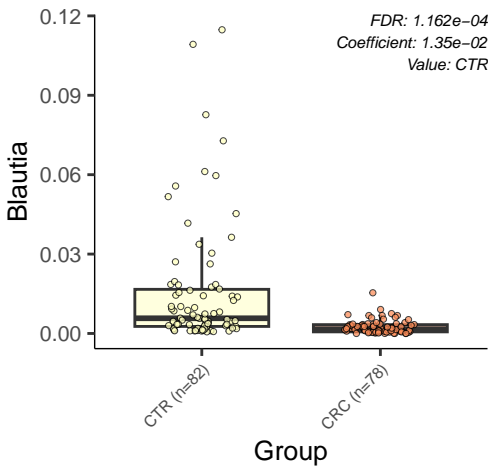
Group

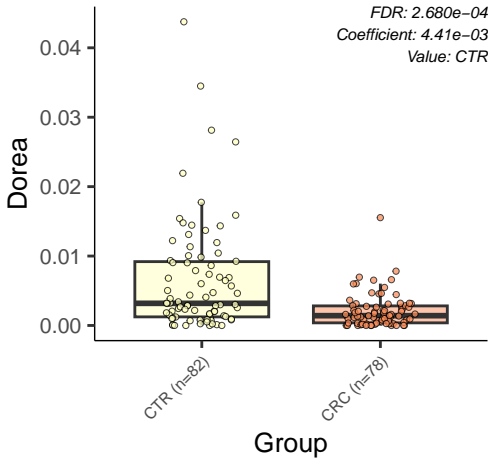


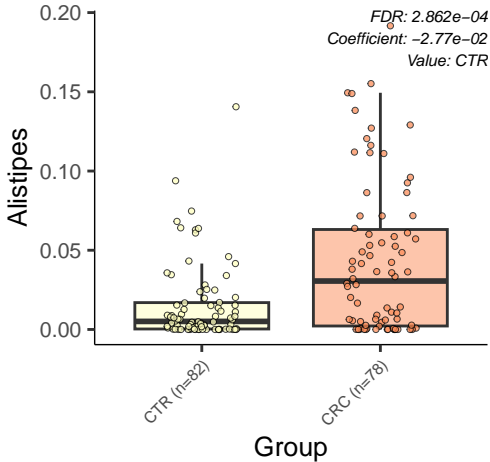
Bilophila

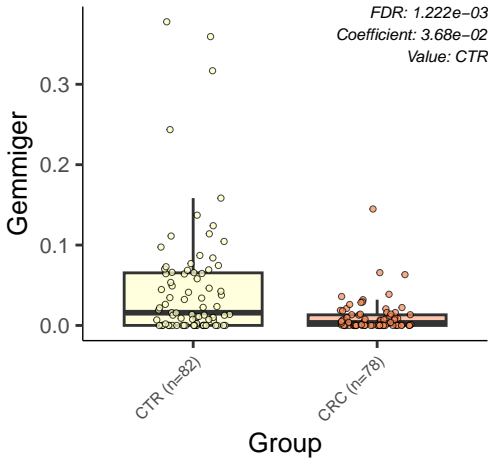




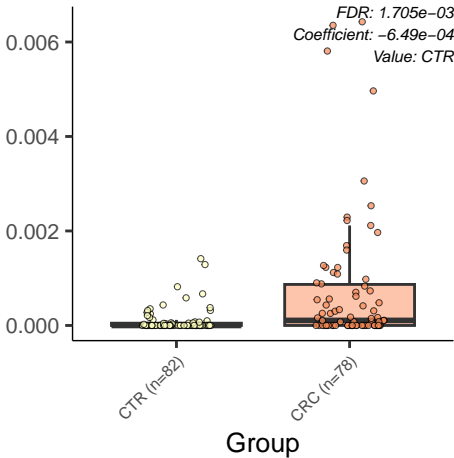






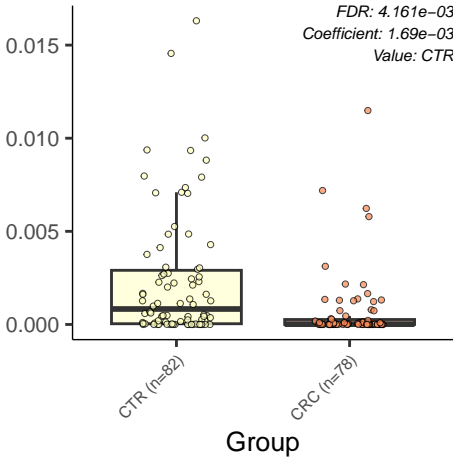


Lawsonibacter



Faecal catena

FDR: 4.161e-03
Coefficient: 1.69e-03
Value: CTR



Gordonibacter

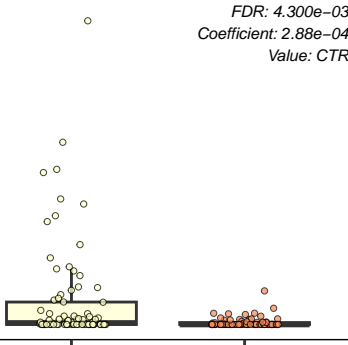
0.004
0.003
0.002
0.001
0.000

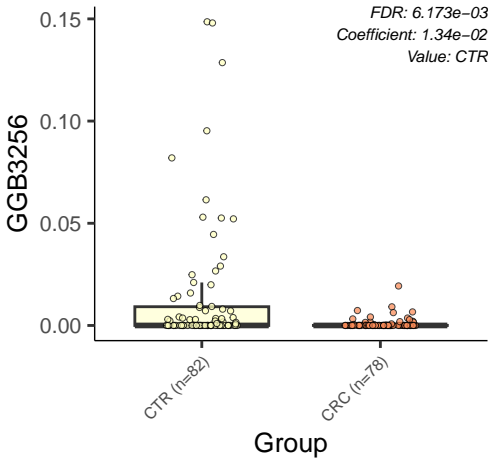
FDR: 4.300e-03
Coefficient: 2.88e-04
Value: CTR

CTR (n=82)

CRC (n=78)

Group





Phoce

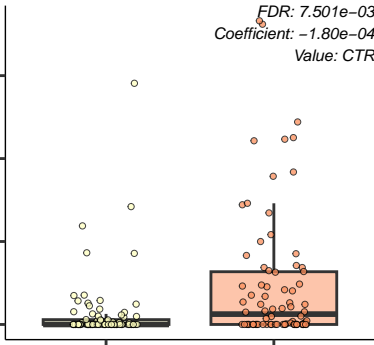
0.0015
0.0010
0.0005
0.0000

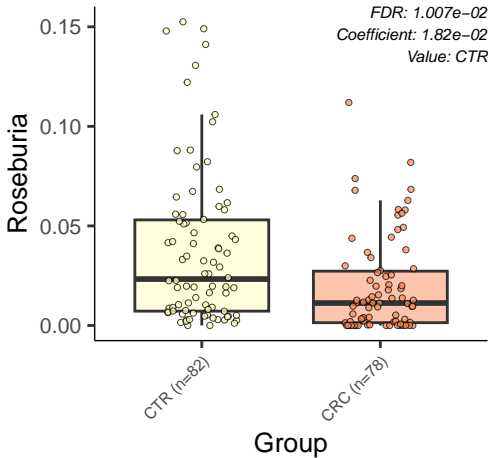
CTR (n=82)

CRC (n=78)

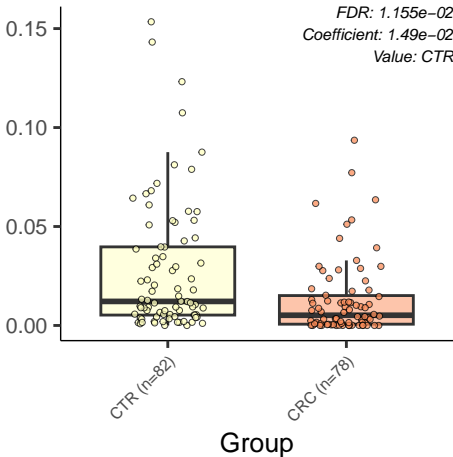
Group

FDR: $7.501e-03$
Coefficient: $-1.80e-04$
Value: CTR





Ruminococcus



Odoribacter

0.02

0.01

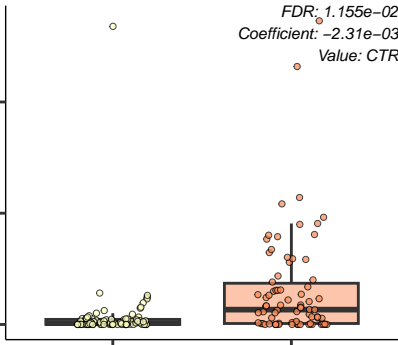
0.00

CTR (n=82)

CRC (n=78)

Group

FDR: $1.155e-02$
Coefficient: $-2.31e-03$
Value: CTR



Dysosmobacter

0.03

0.02

0.01

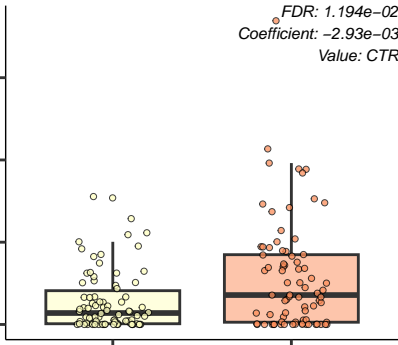
0.00

CTR (n=82)

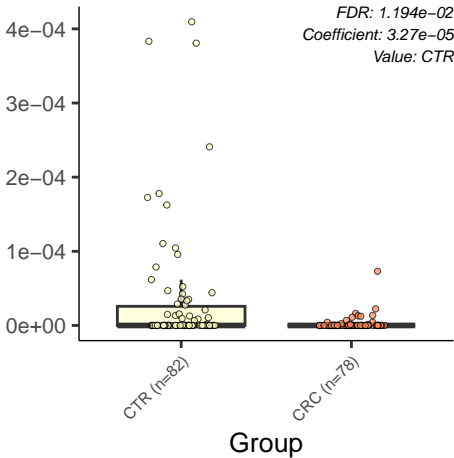
CRC (n=78)

Group

FDR: $1.194e-02$
Coefficient: $-2.93e-03$
Value: CTR



Massiliimalia



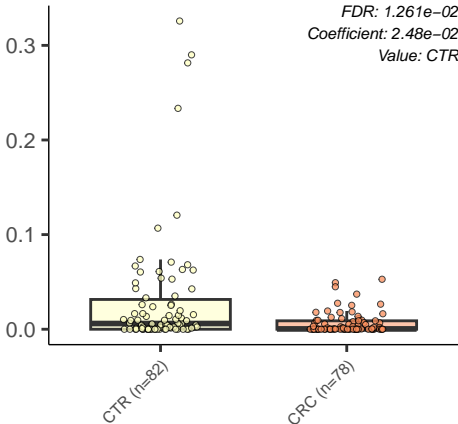
Candidatus_Cibionibacter

FDR: 1.261e-02
Coefficient: 2.48e-02
Value: CTR

CTR (n=82)

CRC (n=78)

Group



Butyricicoccus

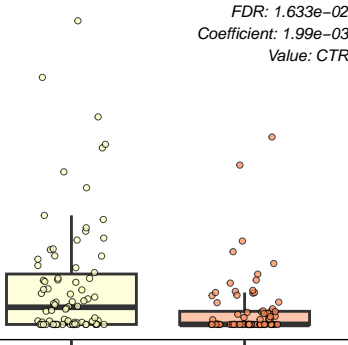
FDR: 1.633e-02
Coefficient: 1.99e-03
Value: CTR

0.025
0.020
0.015
0.010
0.005
0.000

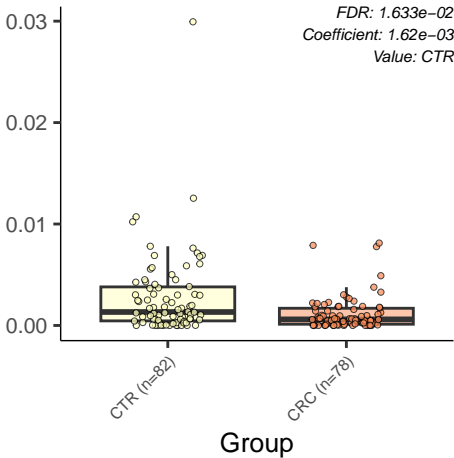
CTR (n=82)

CRC (n=78)

Group



Agathobaculum



Candidatus_Pararuminococcus

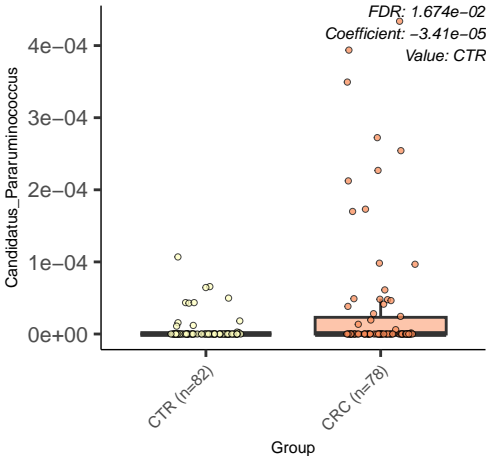
4e-04
3e-04
2e-04
1e-04
0e+00

FDR: 1.674e-02
Coefficient: -3.41e-05
Value: CTR

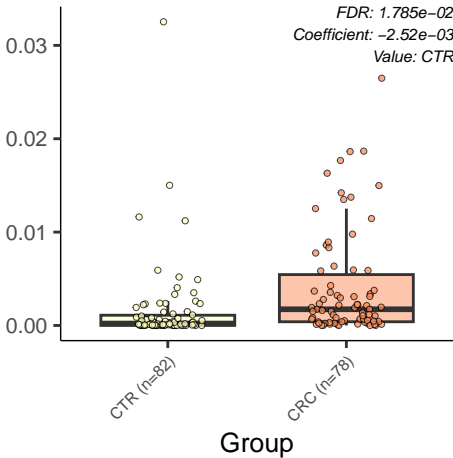
CTR (n=82)

CRC (n=78)

Group



Flavonifractor



Lachnoclostridium

0.020

0.015

0.010

0.005

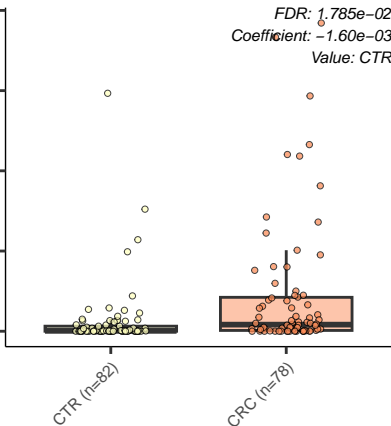
0.000

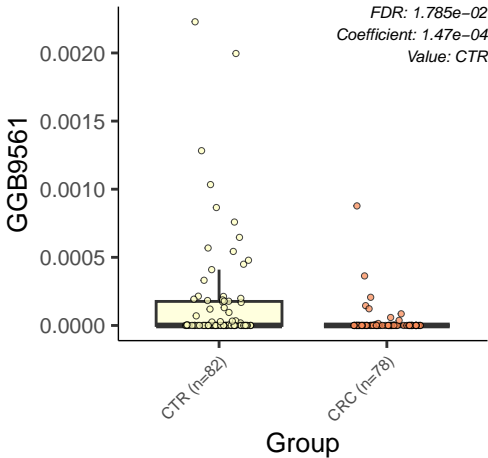
CTR (n=82)

CRC (n=78)

Group

FDR: $1.785e-02$
Coefficient: $-1.60e-03$
Value: CTR





Candidatus_Nanosynsacchari

FDR: 1.937e-02
Coefficient: 3.36e-04
Value: CTR

0.006

0.004

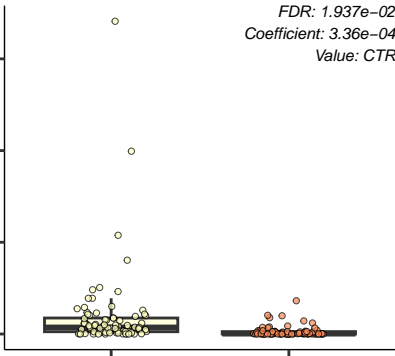
0.002

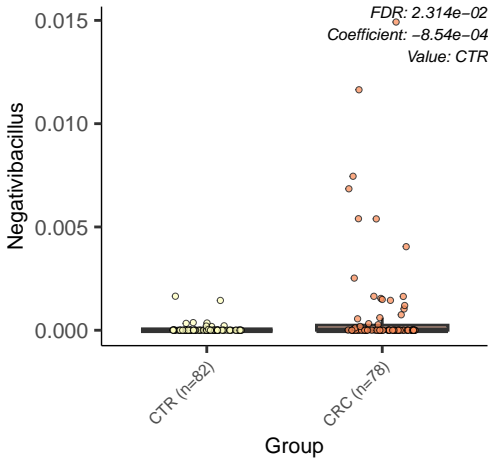
0.000

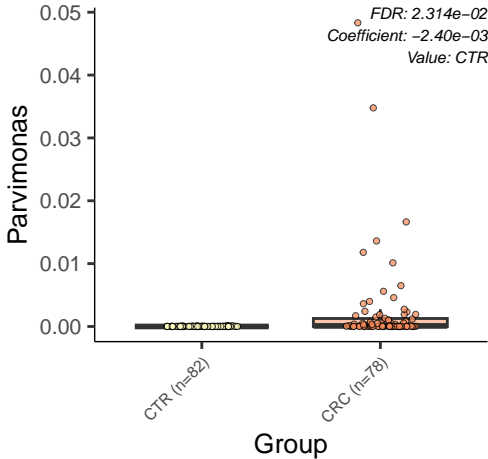
CTR (n=82)

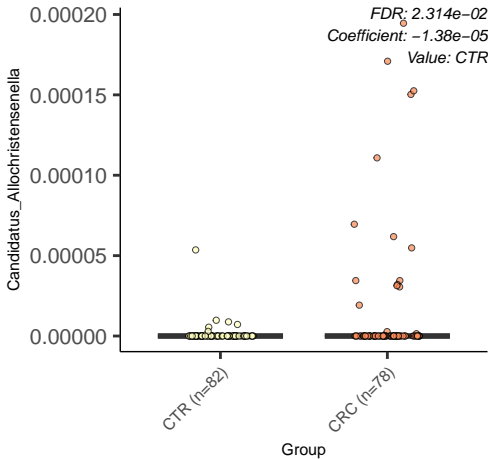
CRC (n=78)

Group









GGB79734

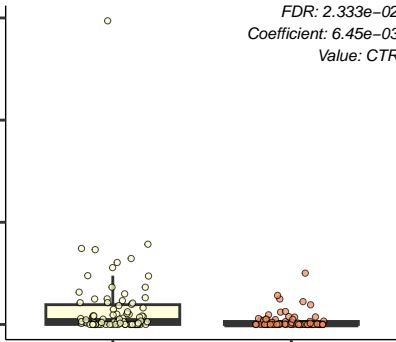
0.15
0.10
0.05
0.00

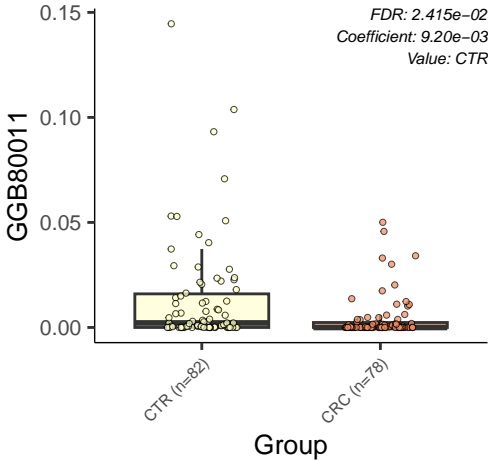
FDR: 2.333e-02
Coefficient: 6.45e-03
Value: CTR

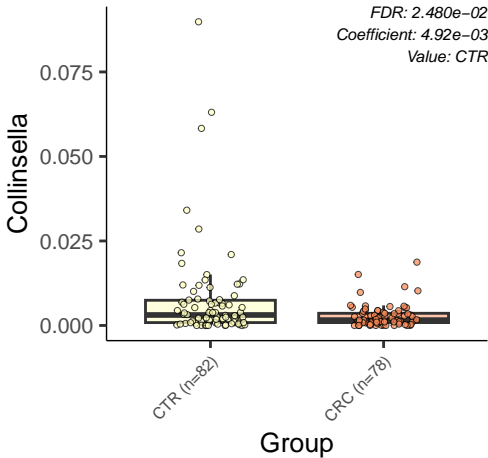
CTR (n=82)

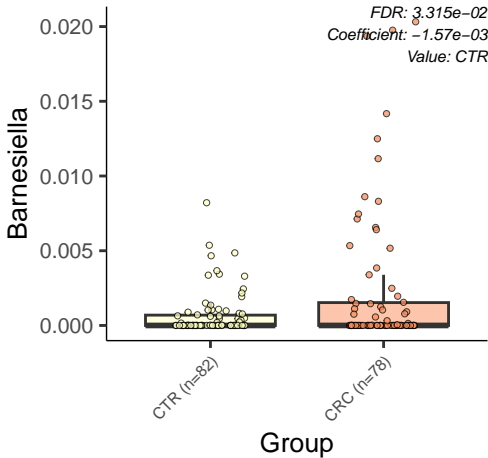
CRC (n=78)

Group









Anaerobutyricum

FDR: $3.758e-02$
Coefficient: $5.70e-04$
Value: CTR

0.0075

0.0050

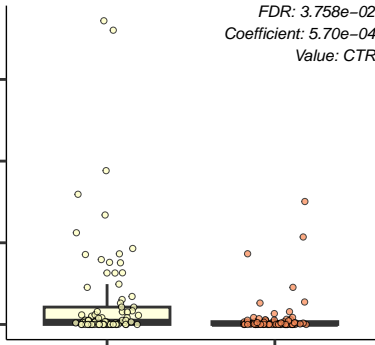
0.0025

0.0000

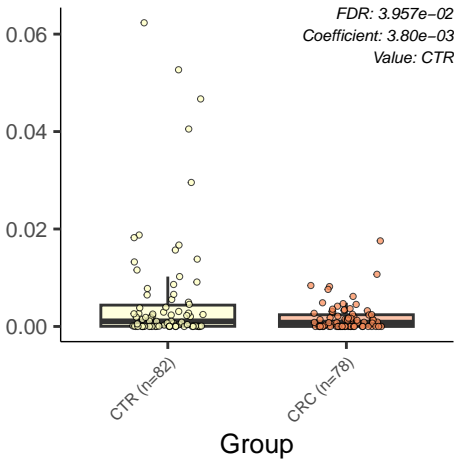
CTR (n=82)

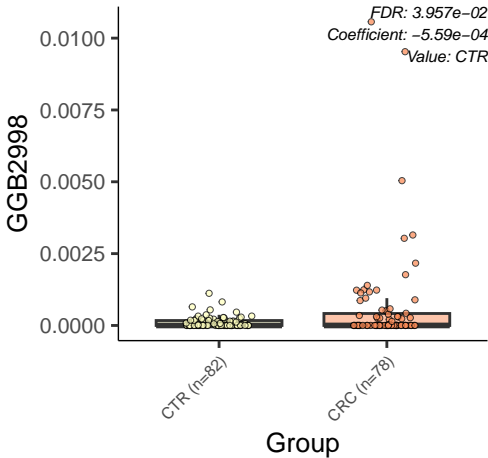
CRC (n=78)

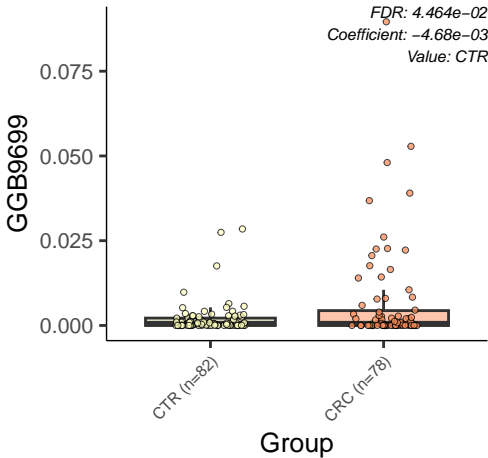
Group



Faecalibacillus







GGB9787

FDR: 4.464e-02
Coefficient: 1.82e-04
Value: CTR

0.004

0.003

0.002

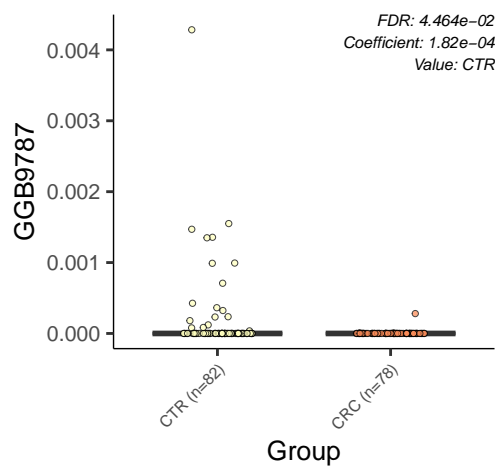
0.001

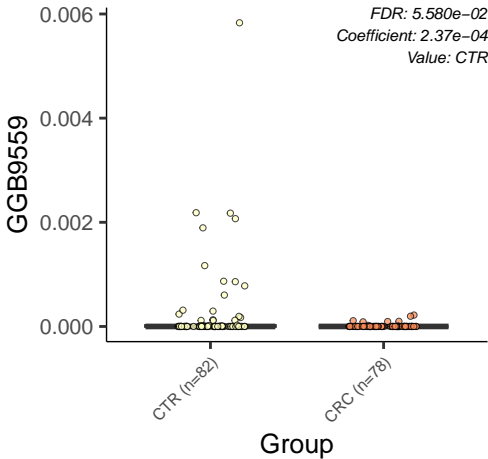
0.000

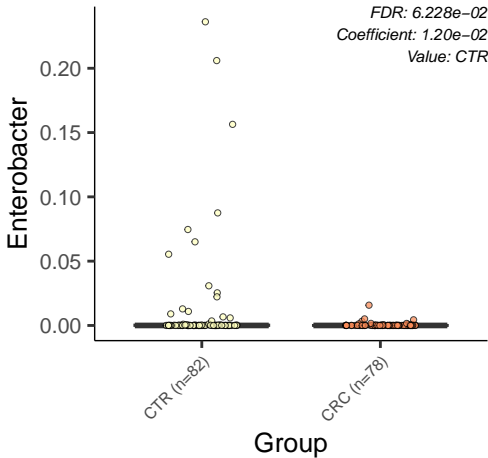
CTR (n=82)

CRC (n=78)

Group







Solobacterium

0.006

0.004

0.002

0.000

CTR (n=82)

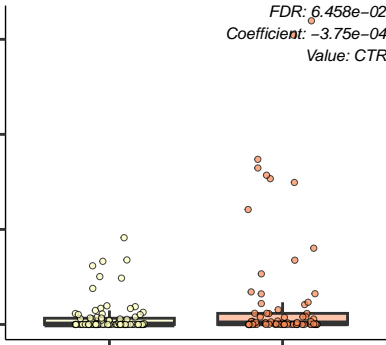
CRC (n=78)

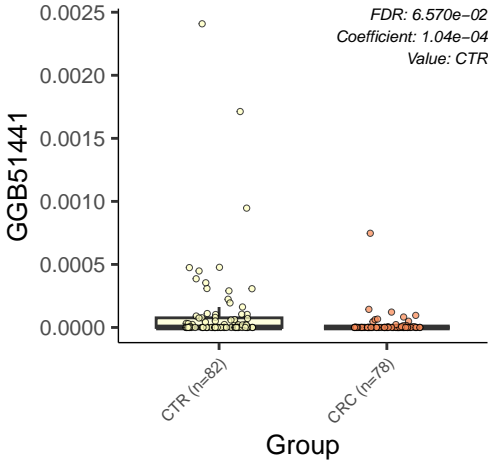
Group

FDR: $6.458e-02$

Coefficient: $-3.75e-04$

Value: CTR





Hydrogenoanaerobacterium

FDR: $6.757e-02$
Coefficient: $-1.61e-04$
Value: CTR

0.002

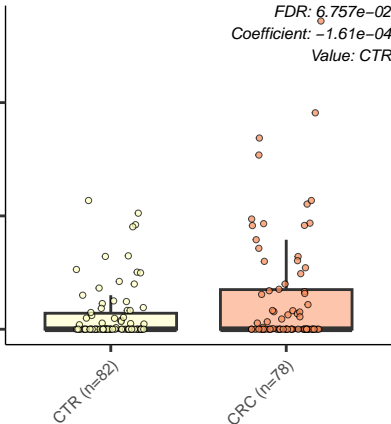
0.001

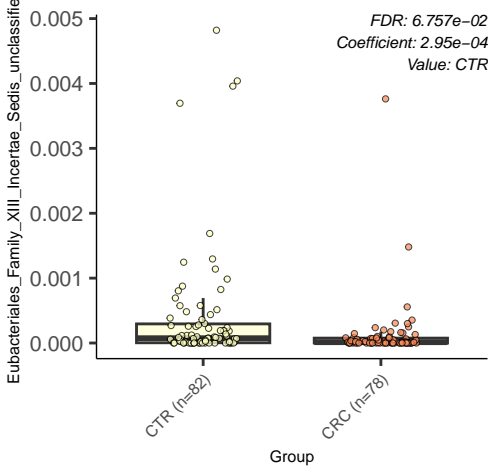
0.000

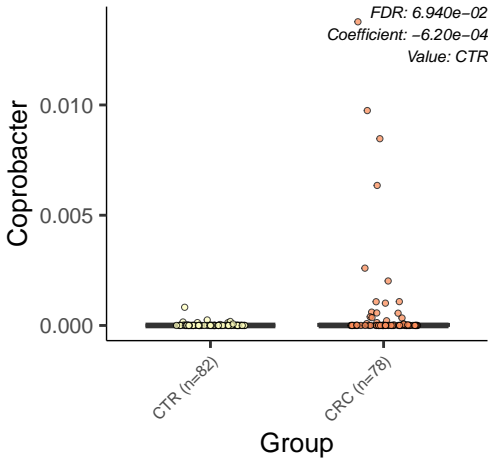
CTR (n=82)

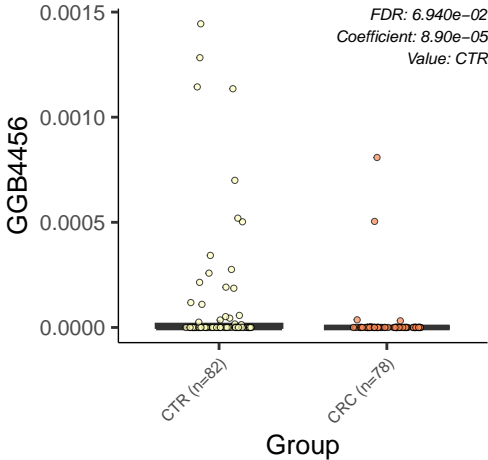
CRC (n=78)

Group









GGB3653

FDR: 8.264e-02
Coefficient: 1.14e-03
Value: CTR

0.02

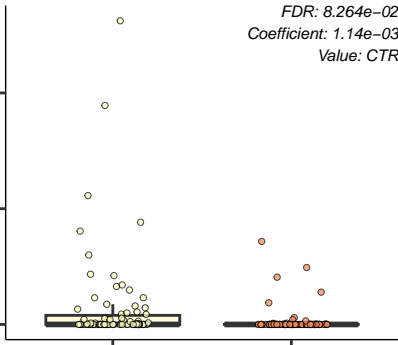
0.01

0.00

CTR (n=82)

CRC (n=78)

Group



Anaerotruncus

0.010

0.005

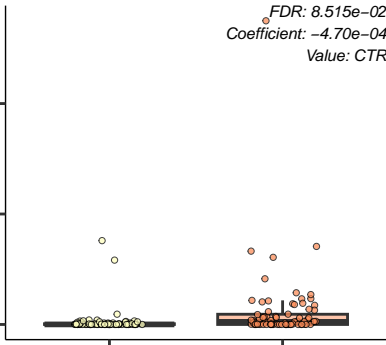
0.000

CTR (n=82)

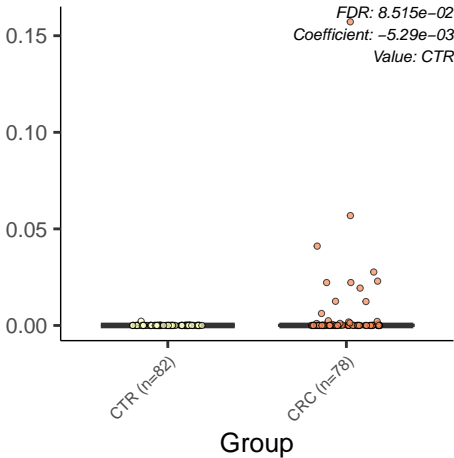
CRC (n=78)

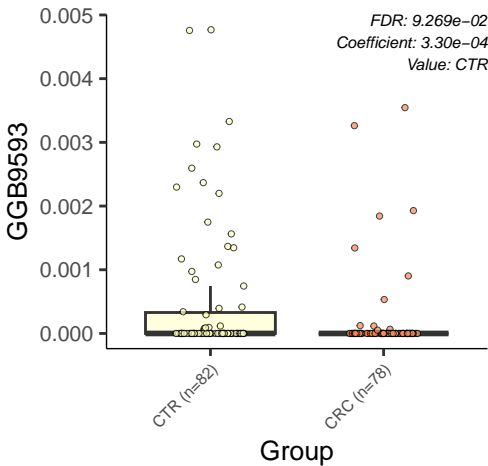
Group

FDR: $8.515e-02$
Coefficient: $-4.70e-04$
Value: CTR



Porphyromonas





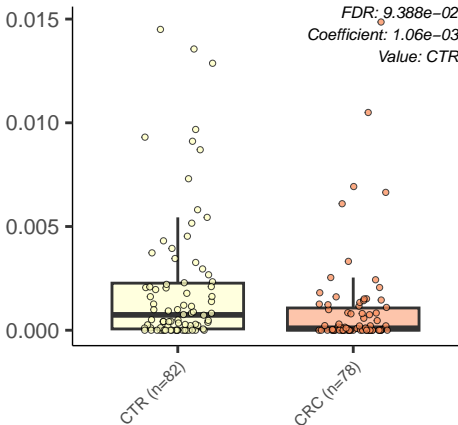
Firmicutes_unclassified

FDR: 9.388e-02
Coefficient: 1.06e-03
Value: CTR

CTR (n=82)

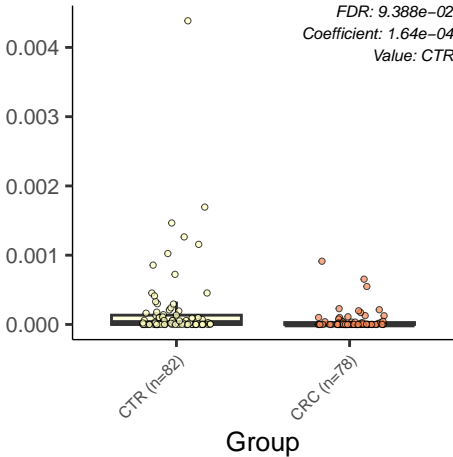
CRC (n=78)

Group

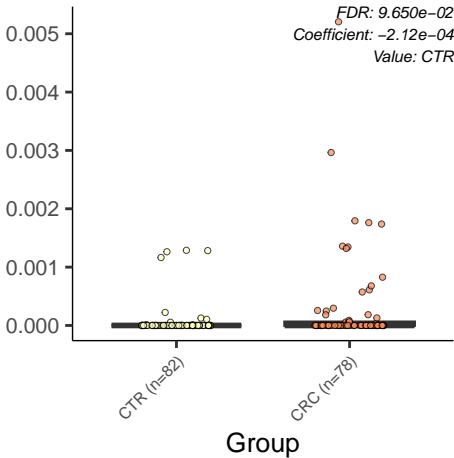


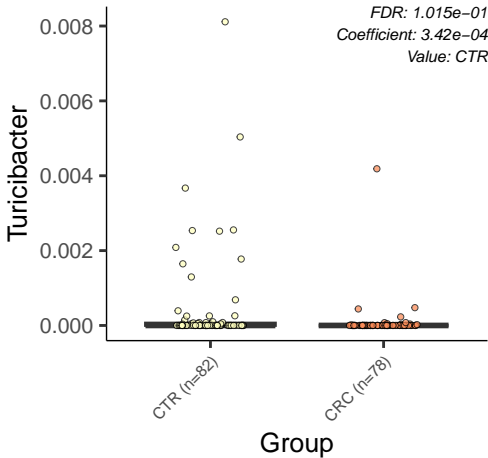
Mogibacterium

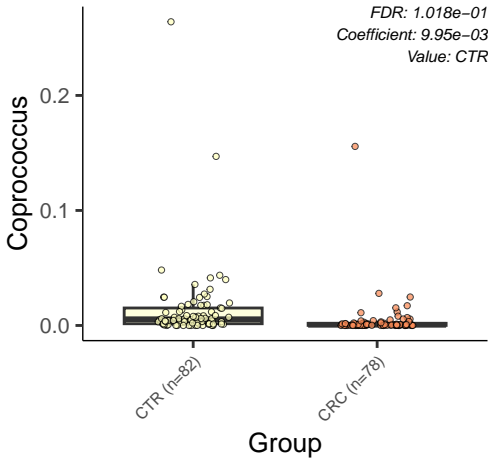
FDR: 9.388e-02
Coefficient: 1.64e-04
Value: CTR

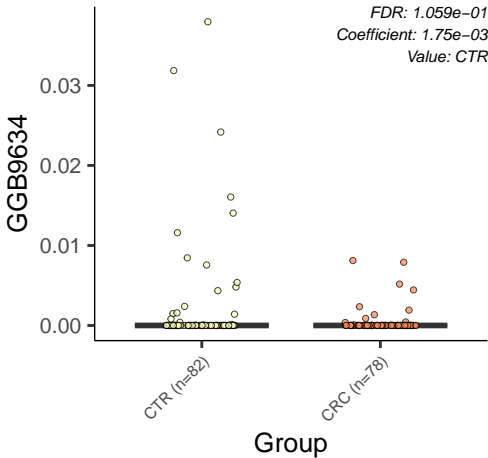


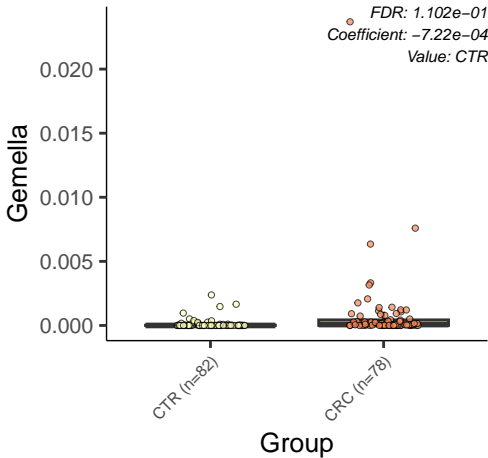
GGB3109

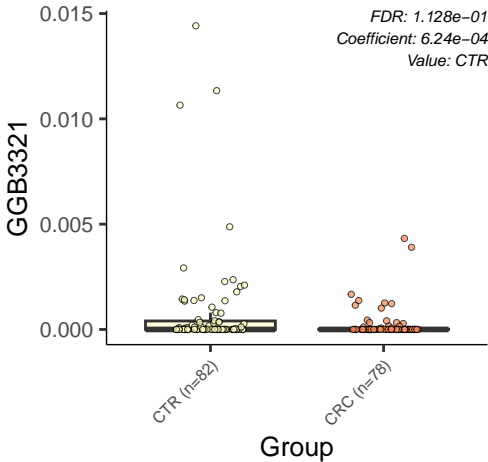


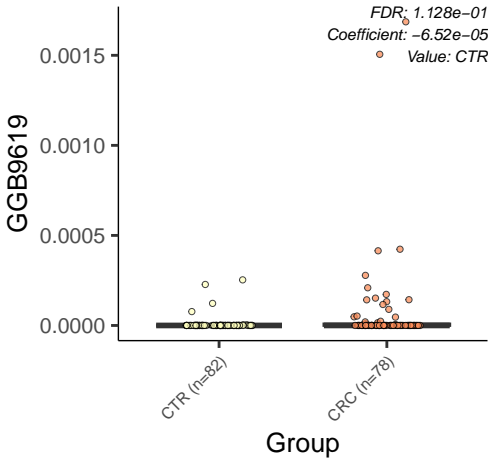


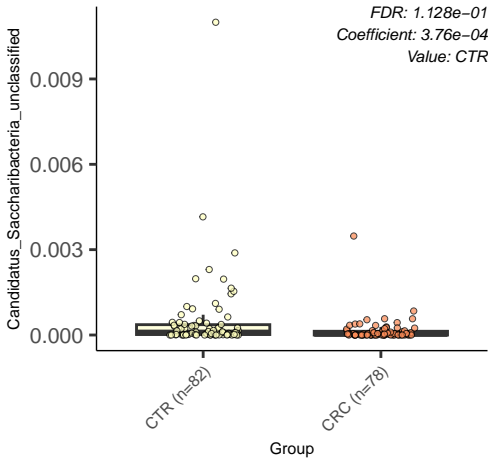


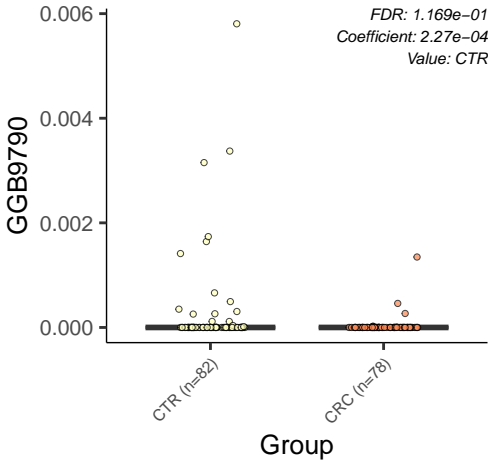


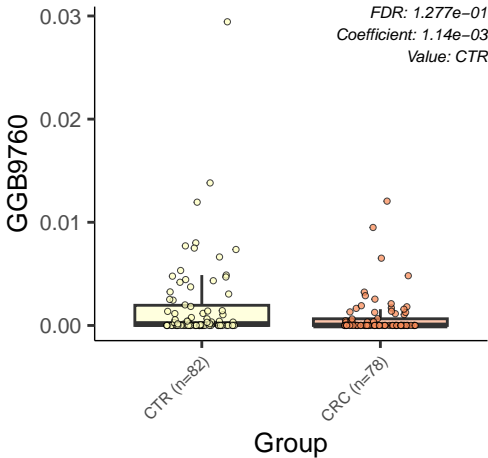




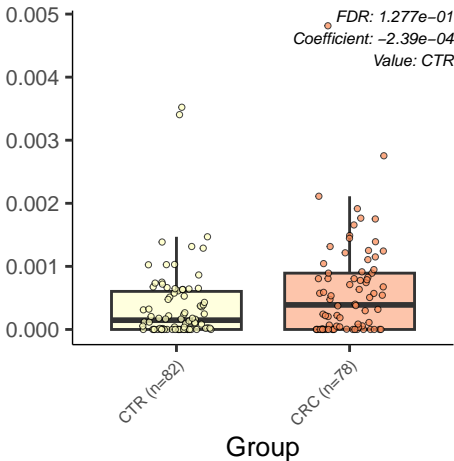








Rheinheimer



Raoutibacter

FDR: 1.277e-01
Coefficient: 4.43e-05
Value: CTR

1e-03

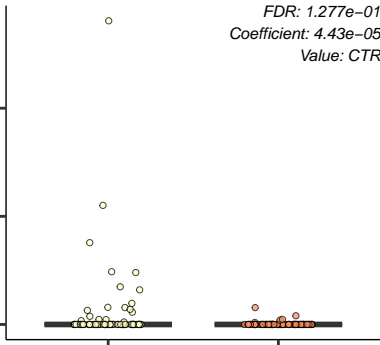
5e-04

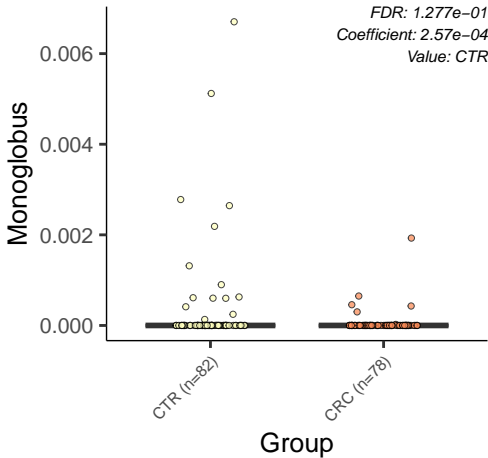
0e+00

CTR (n=82)

CRC (n=78)

Group





Mesosutterella

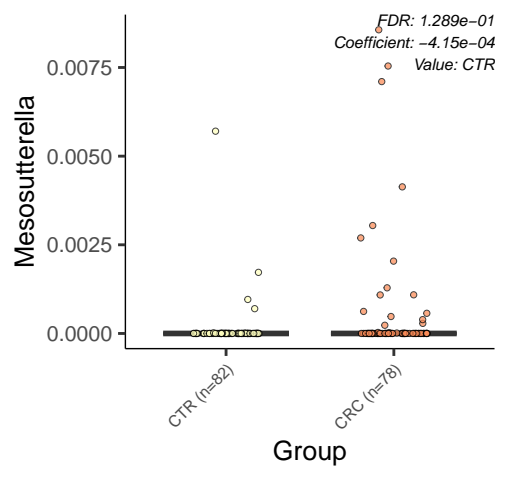
0.0075
0.0050
0.0025
0.0000

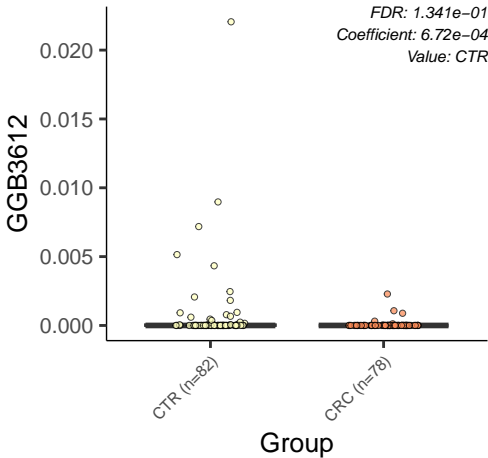
CTR (n=82)

CRC (n=78)

Group

FDR: 1.289e-01
Coefficient: -4.15e-04
Value: CTR





GGB2653

0.0020

0.0015

0.0010

0.0005

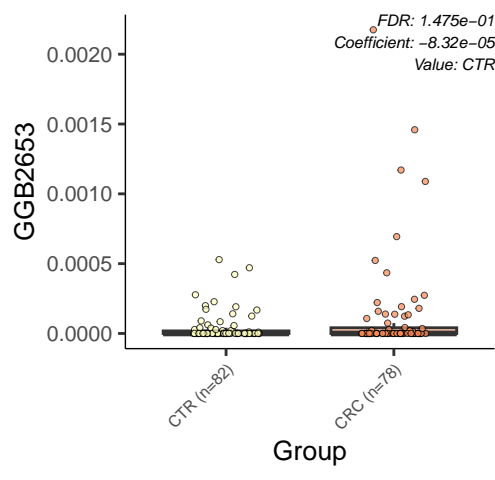
0.0000

FDR: 1.475e-01
Coefficient: -8.32e-05
Value: CTR

CTR (n=82)

CRC (n=78)

Group



Peptostreptococcus

0.04

0.02

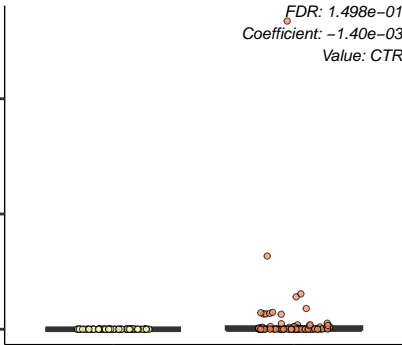
0.00

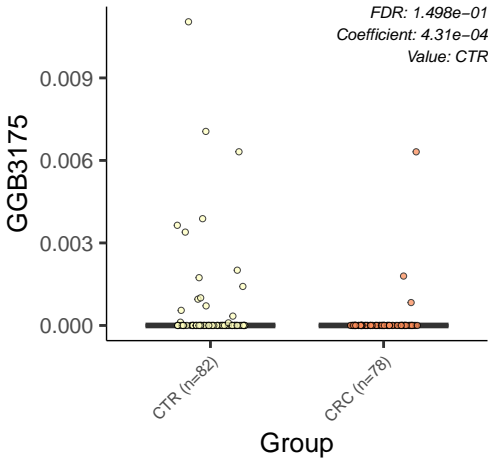
CTR (n=82)

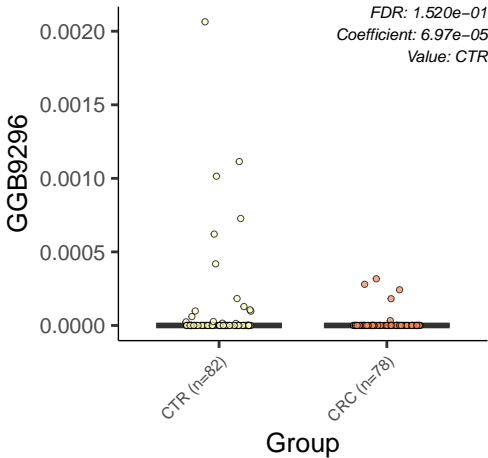
CRC (n=78)

Group

FDR: 1.498e-01
Coefficient: -1.40e-03
Value: CTR







Lachnospiraceae_unclassified

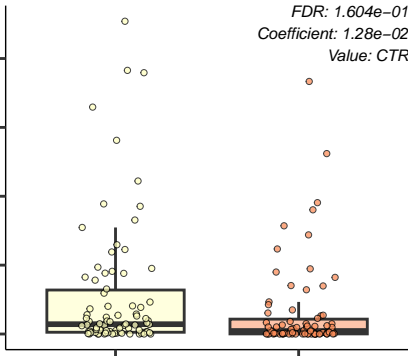
FDR: 1.604e-01
Coefficient: 1.28e-02
Value: CTR

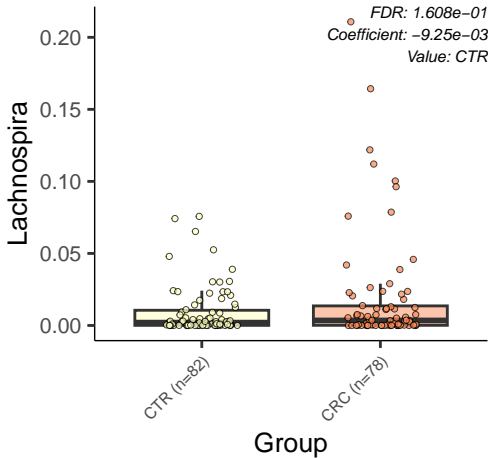
0.20
0.15
0.10
0.05
0.00

CTR (n=82)

CRC (n=78)

Group





Christensenella

FDR: 1.716e-01
Coefficient: 7.39e-05
Value: CTR

0.0020

0.0015

0.0010

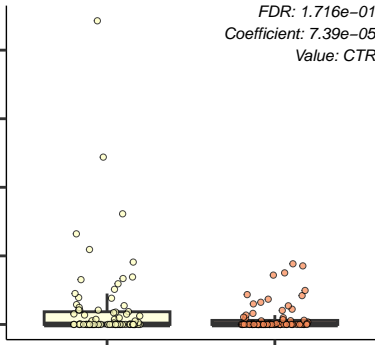
0.0005

0.0000

CTR (n=82)

CRC (n=78)

Group



Haemophilus

FDR: 1.717e-01
Coefficient: 2.55e-03
Value: CTR

0.075

0.050

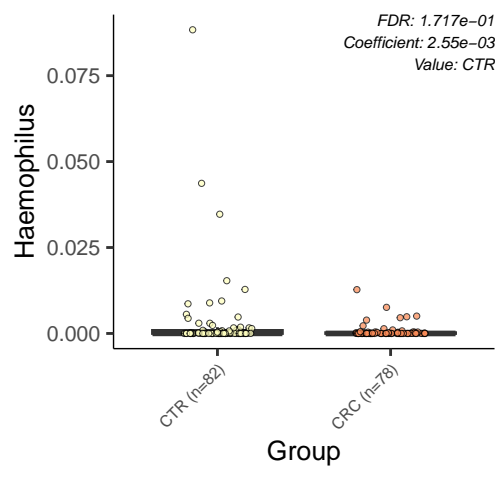
0.025

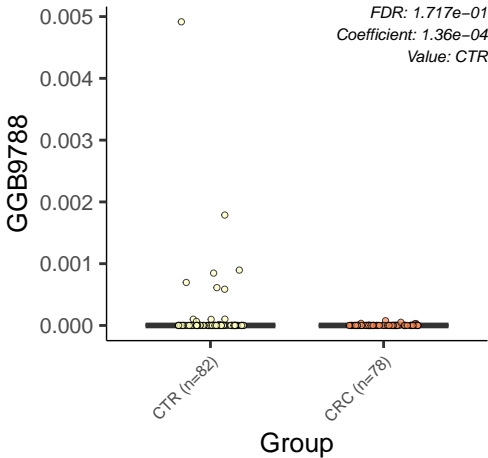
0.000

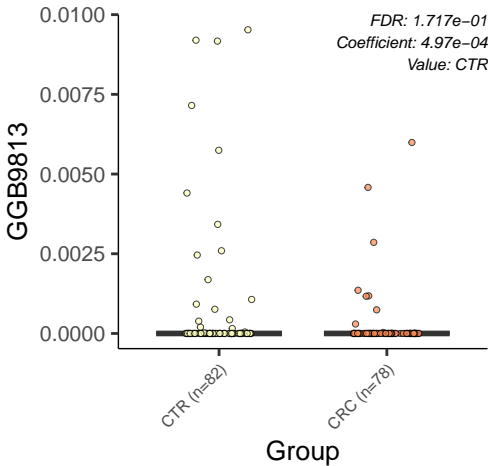
CTR (n=82)

CRC (n=78)

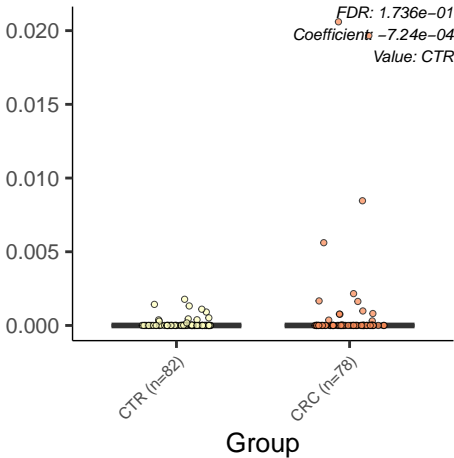
Group

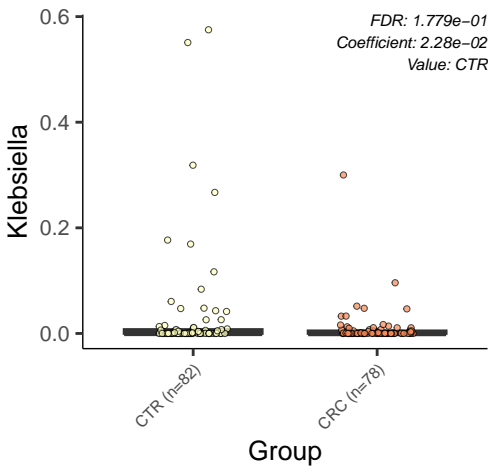


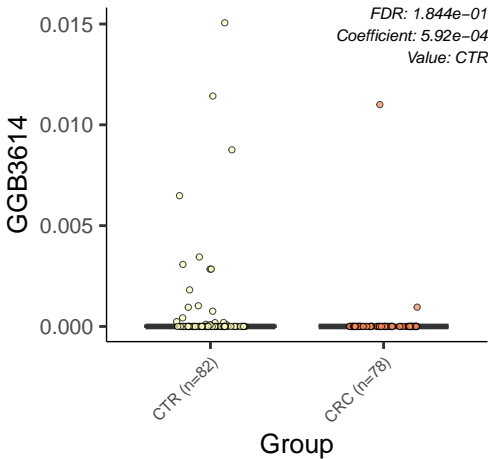


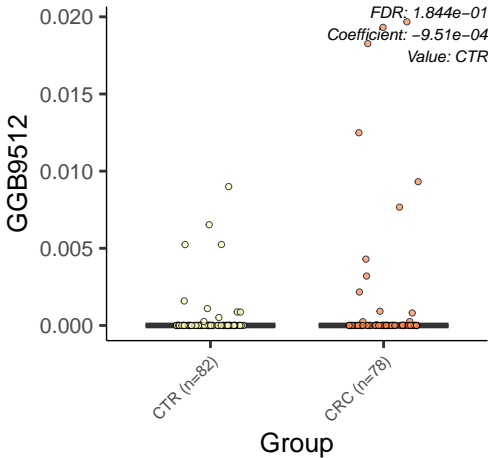


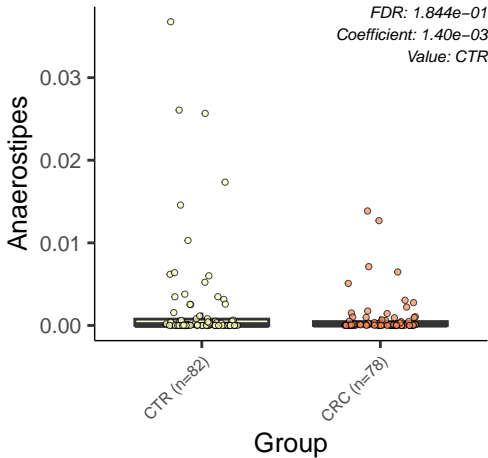
GGB9524

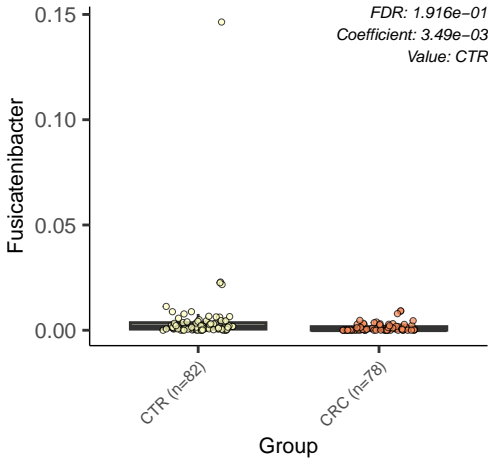












Value: CTR

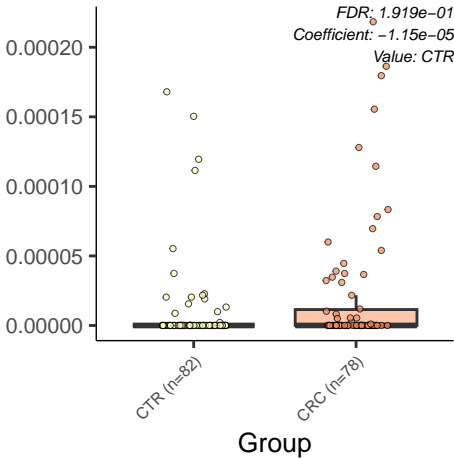


CTR (n=82)

CRC (n=78)

Group

GGB58233



Fusobacterium

FDR: 1.919e-01
Coefficient: -5.95e-03
Value: CTR

0.2

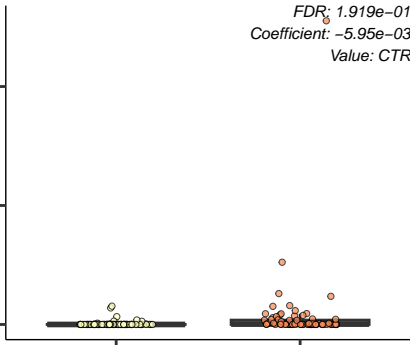
0.1

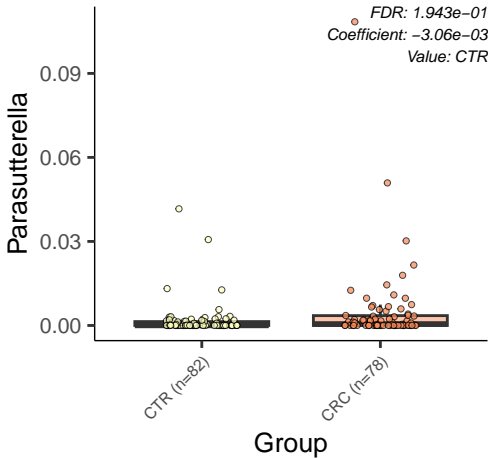
0.0

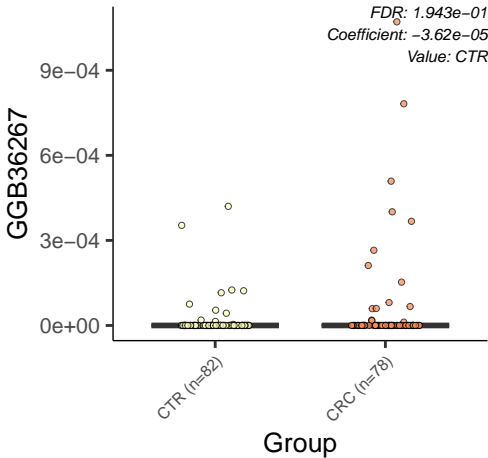
CTR (n=82)

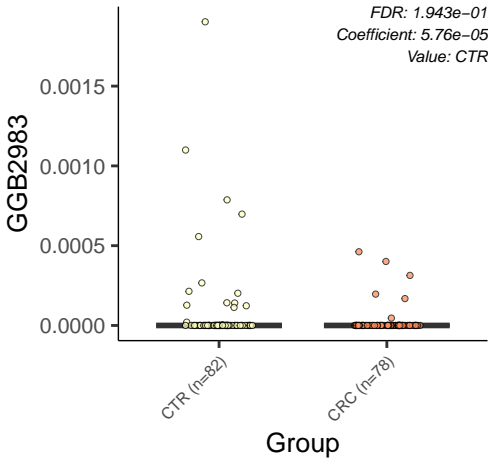
CRC (n=78)

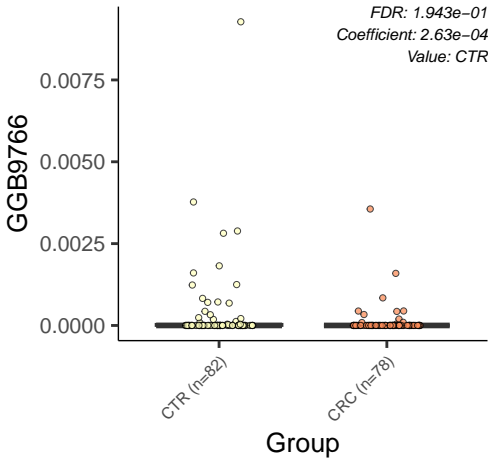
Group

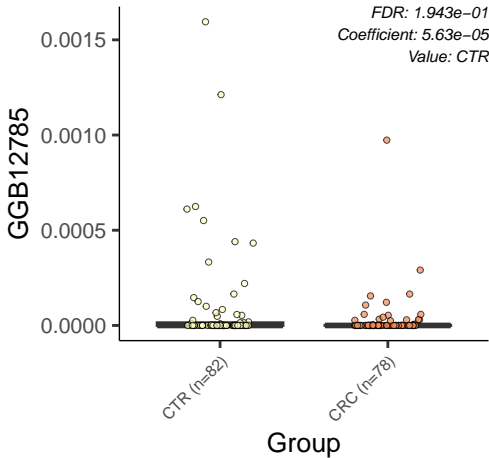


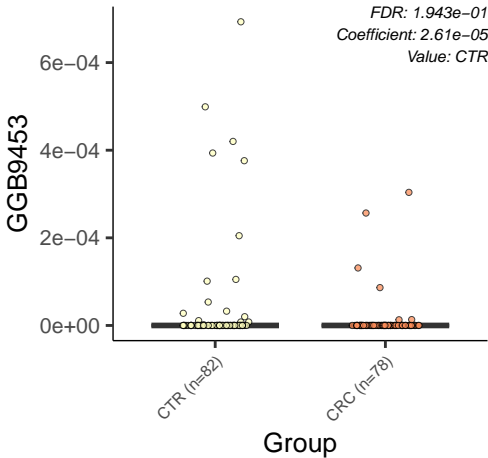




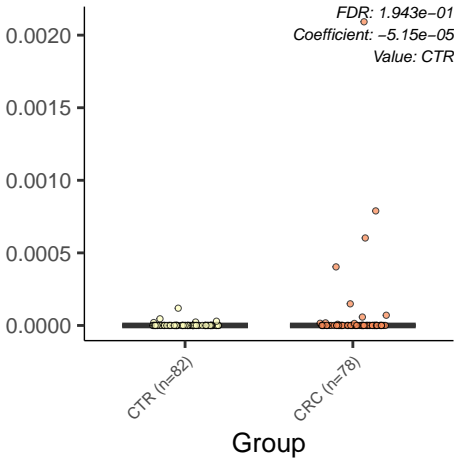


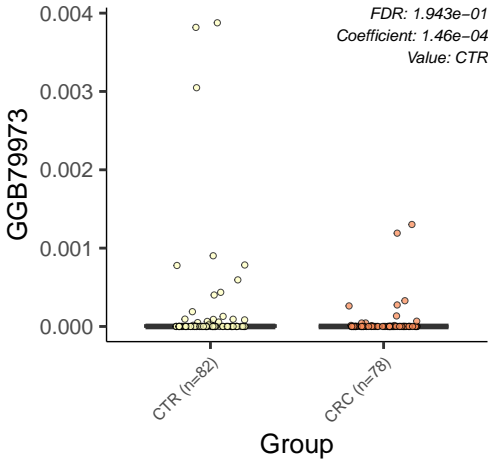






Campylobacter





GGB55957

FDR: 1.943e-01

Coefficient: -1.44e-05

Value: CTR

4e-04

3e-04

2e-04

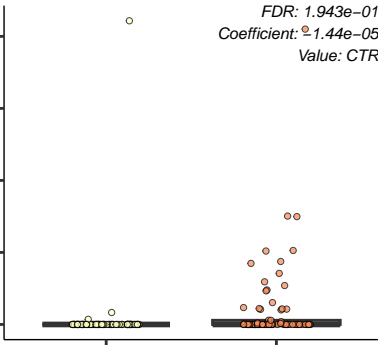
1e-04

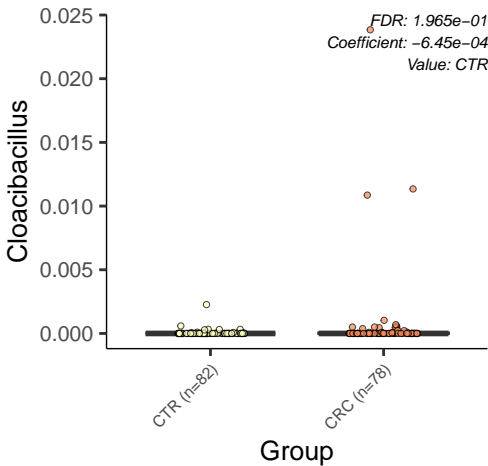
0e+00

CTR (n=82)

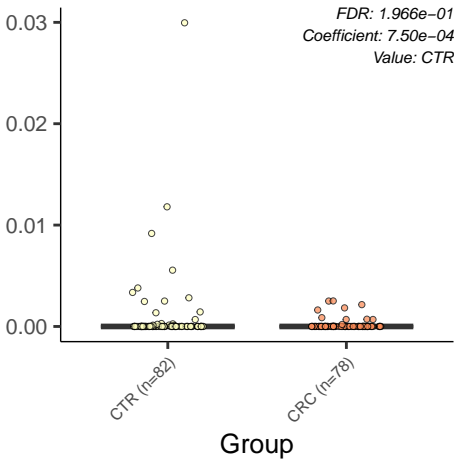
CRC (n=78)

Group





Faecalimonas



Eubacteriales_unclassified

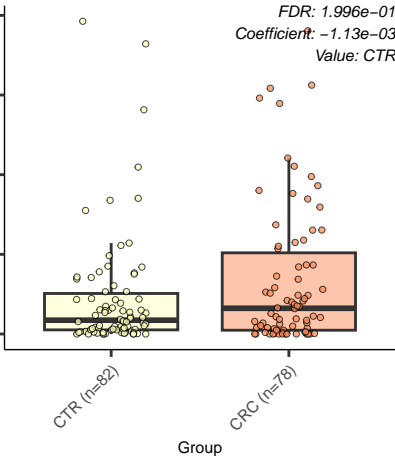
0.020
0.015
0.010
0.005
0.000

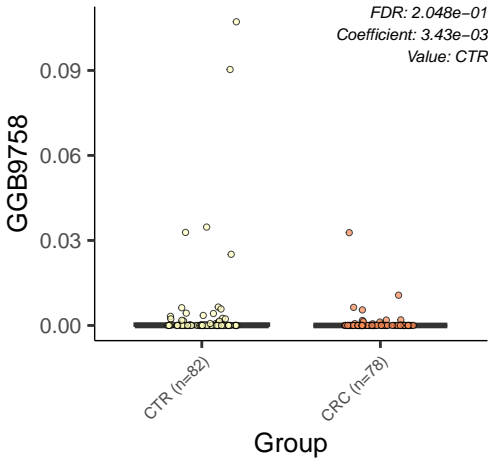
CTR (n=82)

CRC (n=78)

Group

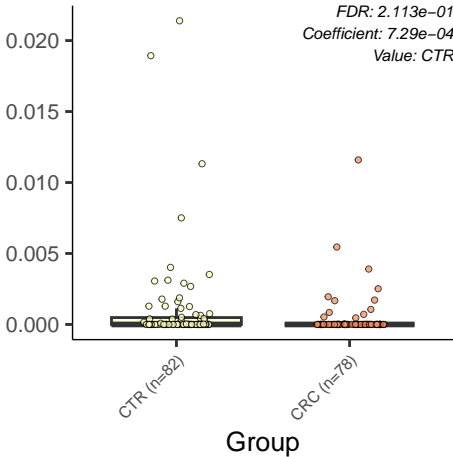
FDR: 1.996e-01
Coefficient: -1.13e-03
Value: CTR





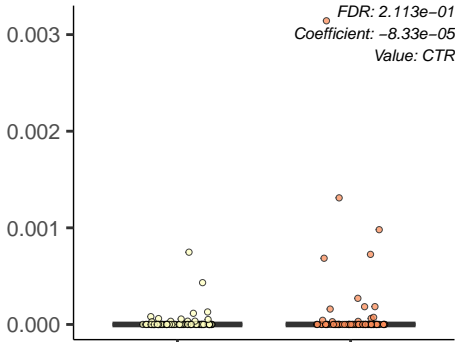
Catenibacterium

FDR: 2.113e-01
Coefficient: 7.29e-04
Value: CTR



Group

GGB34228

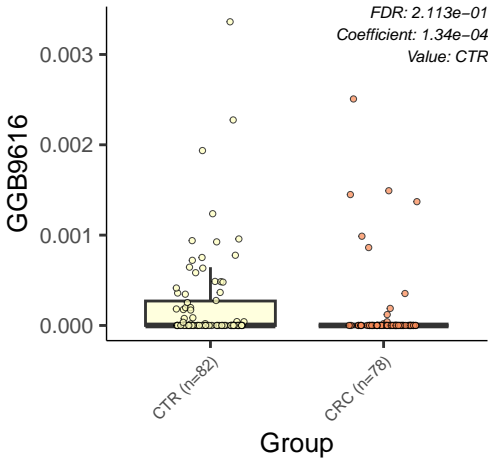


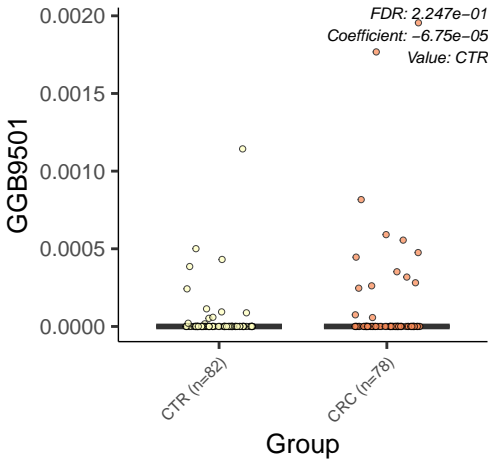
FDR: 2.113e-01
Coefficient: -8.33e-05
Value: CTR

CTR (n=82)

CRC (n=78)

Group





Anaerotignum

FDR: 2.289e-01

Coefficient: -1.91e-03

Value: CTR

0.04

0.03

0.02

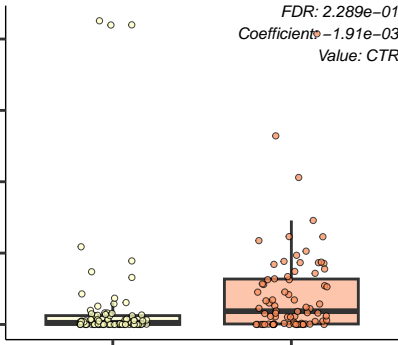
0.01

0.00

CTR (n=82)

CRC (n=78)

Group



GGB3815

0.0015

0.0010

0.0005

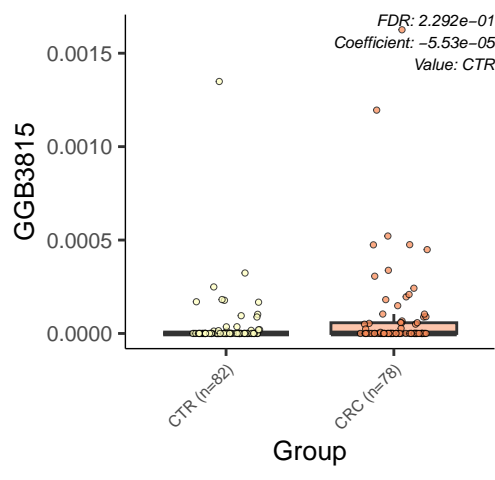
0.0000

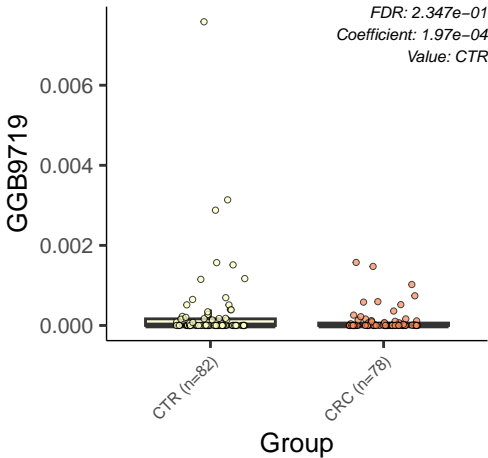
CTR (n=82)

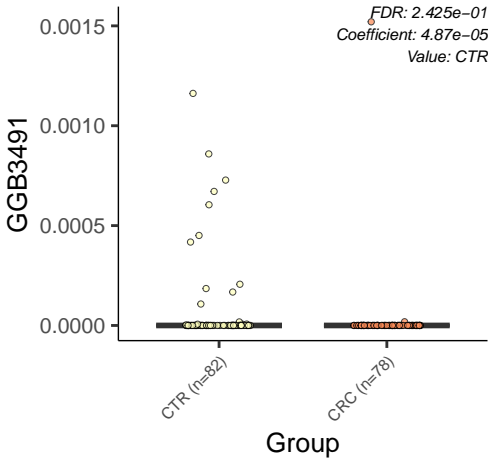
CRC (n=78)

Group

FDR: 2.292e-01
Coefficient: -5.53e-05
Value: CTR







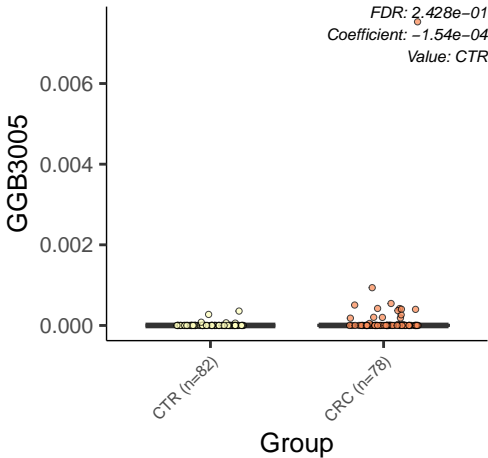
Value: CTR



CTR (n=82)

CRC (n=78)

Group



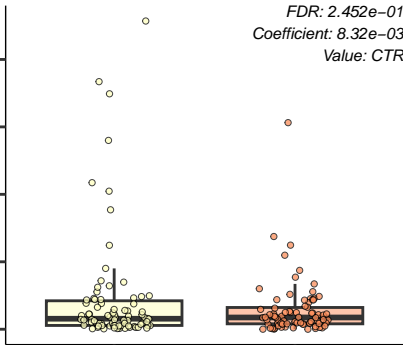
Mediterraneibacter

FDR: 2.452e-01
Coefficient: 8.32e-03
Value: CTR

CTR (n=82)

CRC (n=78)

Group



Intestinibacter

FDR: 2.452e-01
Coefficient: 1.07e-04
Value: CTR

0.002

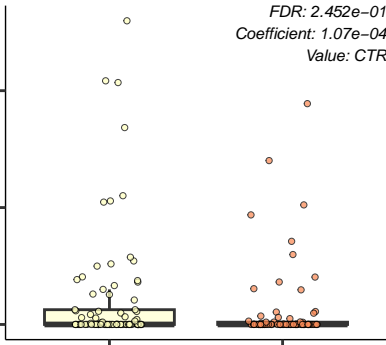
0.001

0.000

CTR (n=82)

CRC (n=78)

Group



Lancefieldella

1e-03

5e-04

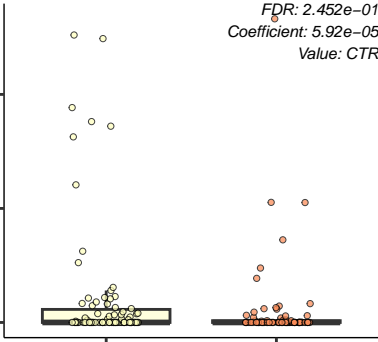
0e+00

CTR (n=82)

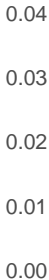
CRC (n=78)

Group

FDR: 2.452e-01
Coefficient: 5.92e-05
Value: CTR



Value: CTR



CTR (n=82)

CRC (n=78)

Group

