

GGB9619_SGB15067

1e-03

5e-04

0e+00

ADA (n=3)

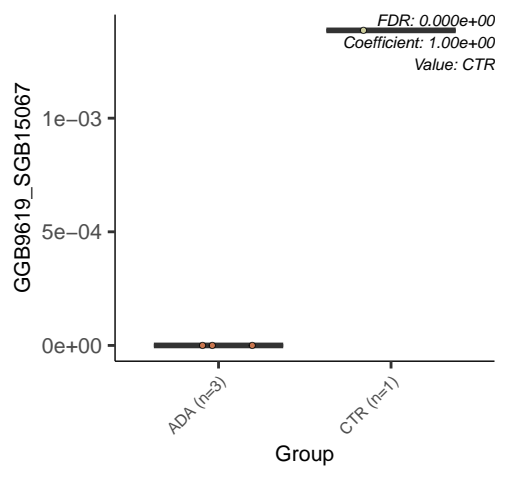
CTR (n=1)

Group

FDR: 0.000e+00

Coefficient: 1.00e+00

Value: CTR



Faecalibacillus_intestinalis

0.04

0.02

0.00

ADA (n=3)

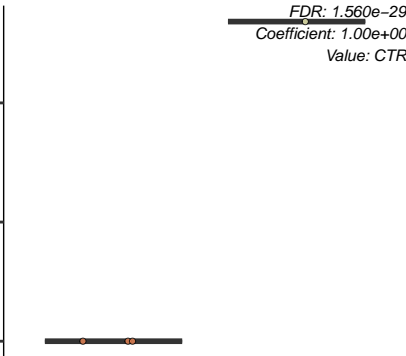
CTR (n=1)

Group

FDR: 1.560e-29

Coefficient: 1.00e+00

Value: CTR



Staphylococcus_SGB6340

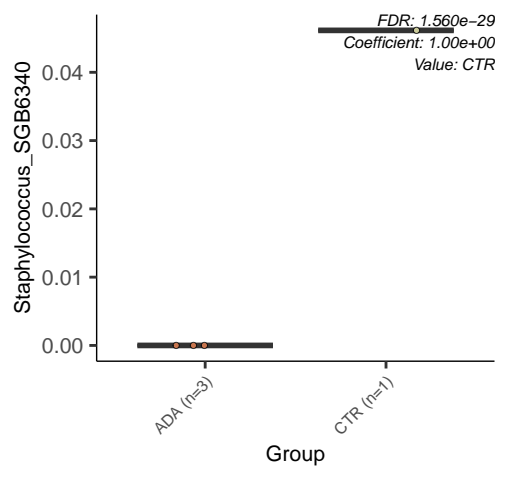
0.04
0.03
0.02
0.01
0.00

ADA (n=3)

CTR (n=1)

Group

FDR: 1.560e-29
Coefficient: 1.00e+00
Value: CTR



Clostridium_sp_AF34_13

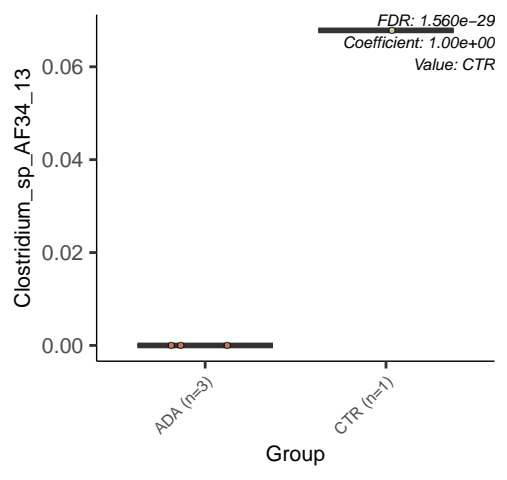
0.06
0.04
0.02
0.00

ADA (n=3)

CTR (n=1)

Group

FDR: 1.560e-29
Coefficient: 1.00e+00
Value: CTR



Klebsiella_pneumoniae

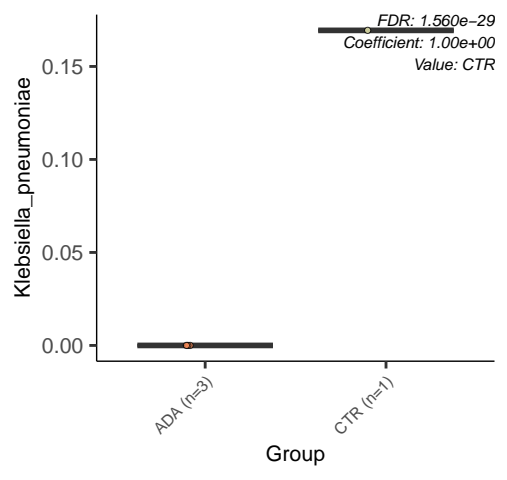
0.15
0.10
0.05
0.00

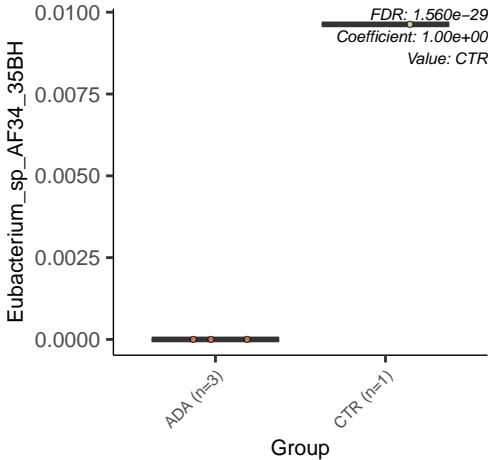
ADA (n=3)

CTR (n=1)

Group

FDR: 1.560e-29
Coefficient: 1.00e+00
Value: CTR





Bacteroides_cellulosilyticus

0.3

0.2

0.1

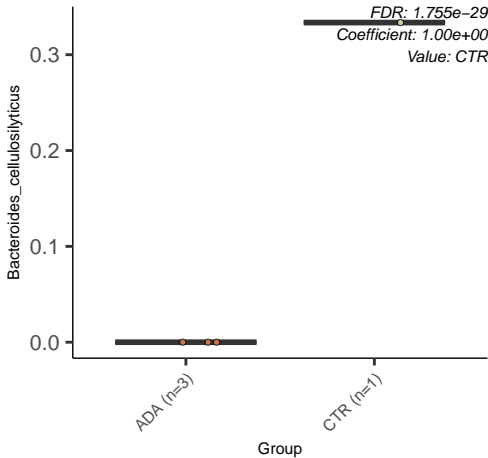
0.0

ADA (n=3)

CTR (n=1)

Group

FDR: 1.755e-29
Coefficient: 1.00e+00
Value: CTR



GGB9453_SGB14845

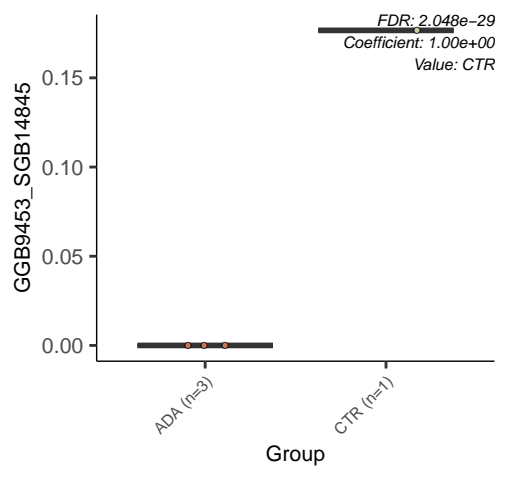
0.15
0.10
0.05
0.00

ADA (n=3)

CTR (n=1)

Group

FDR: 2.048e-29
Coefficient: 1.00e+00
Value: CTR



Bacteroides_xylanisolvens

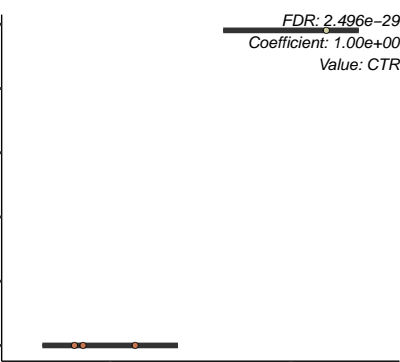
0.05
0.04
0.03
0.02
0.01
0.00

ADA (n=3)

CTR (n=1)

Group

FDR: 2.496e-29
Coefficient: 1.00e+00
Value: CTR



Lachnospira_sp_NSJ_43

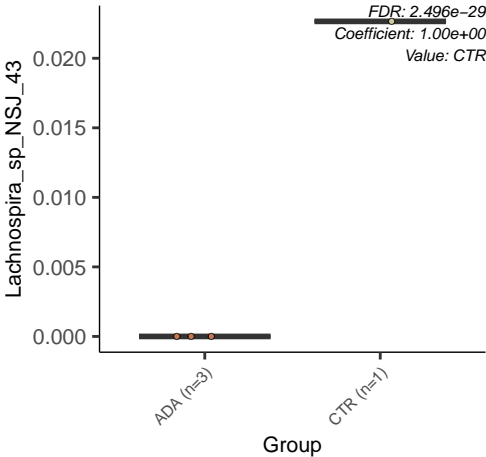
0.020
0.015
0.010
0.005
0.000

ADA (n=3)

CTR (n=1)

Group

FDR: 2.496e-29
Coefficient: 1.00e+00
Value: CTR



GGB3612_SGB4882

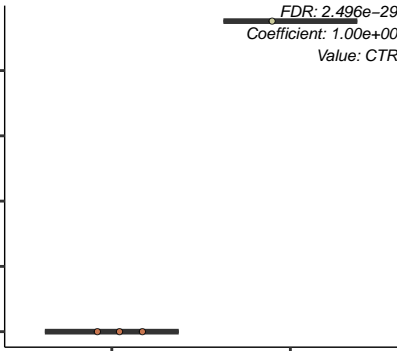
0.04
0.03
0.02
0.01
0.00

ADA (n=3)

CTR (n=1)

Group

FDR: 2.496e-29
Coefficient: 1.00e+00
Value: CTR



GGB2653_SGB3574

0.010

0.005

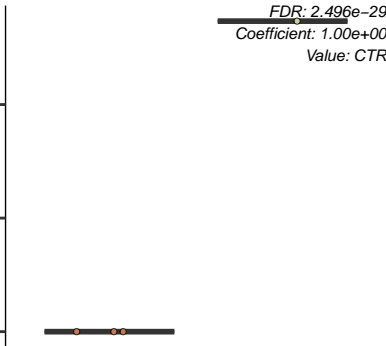
0.000

ADA (n=3)

CTR (n=1)

Group

FDR: 2.496e-29
Coefficient: 1.00e+00
Value: CTR



GGB9760_SGB15373

0.02

0.01

0.00

ADA (n=3)

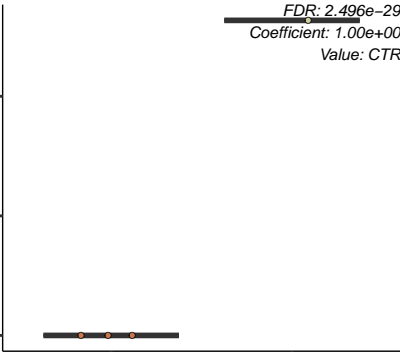
CTR (n=1)

Group

FDR: 2.496e-29

Coefficient: 1.00e+00

Value: CTR



GGB9708_SGB15234

0.06

0.04

0.02

0.00

FDR: 2.496e-29

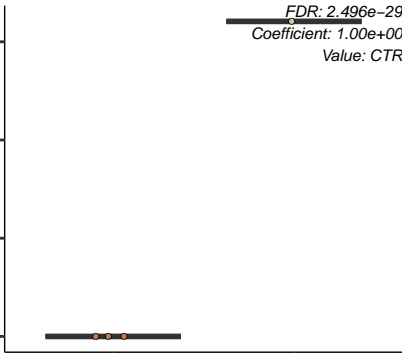
Coefficient: 1.00e+00

Value: CTR

ADA (n=3)

CTR (n=1)

Group



Vescimonas_SGB15087

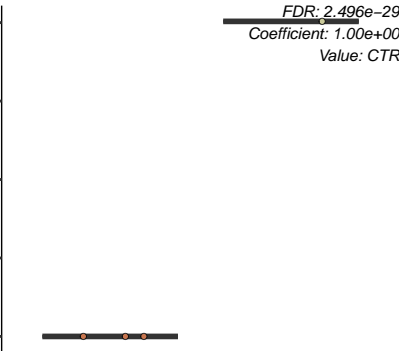
0.08
0.06
0.04
0.02
0.00

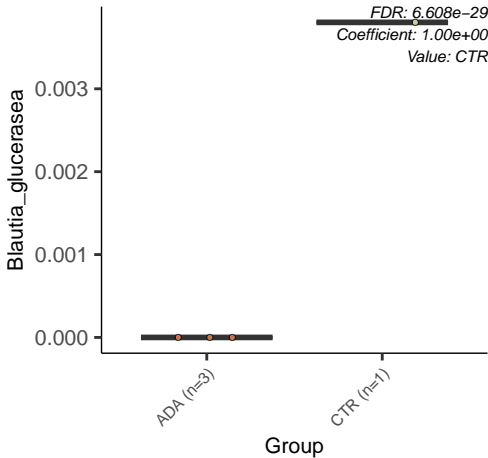
ADA (n=3)

CTR (n=1)

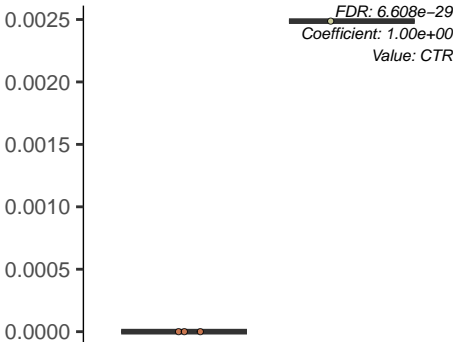
Group

FDR: 2.496e-29
Coefficient: 1.00e+00
Value: CTR





GGB4566_SGB6305



Group

Coprococcus_catus

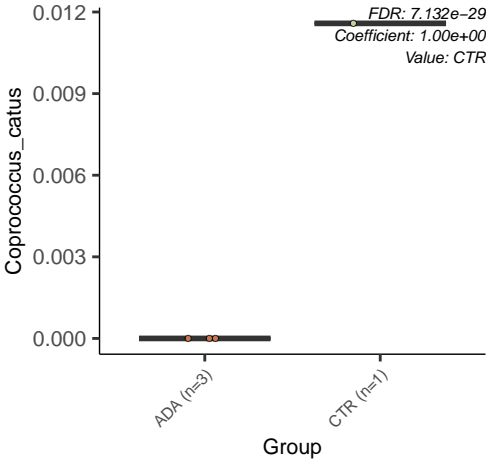
0.012
0.009
0.006
0.003
0.000

FDR: 7.132e-29
Coefficient: 1.00e+00
Value: CTR

ADA (n=3)

CTR (n=1)

Group



Clostridium_sp_AF27_2AA

0.002

0.001

0.000

ADA (n=3)

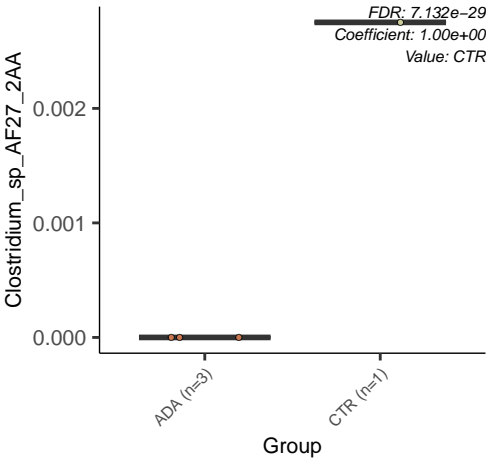
CTR (n=1)

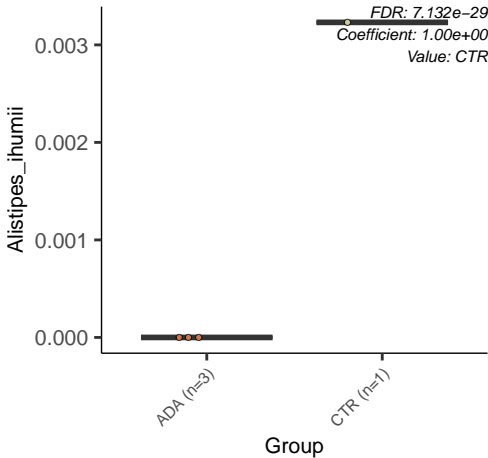
Group

FDR: 7.132e-29

Coefficient: 1.00e+00

Value: CTR





GGB9453_SGB14844

0.003

0.002

0.001

0.000

ADA (n=3)

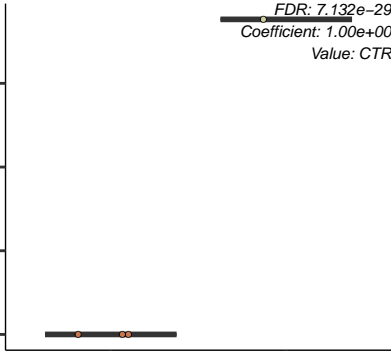
CTR (n=1)

Group

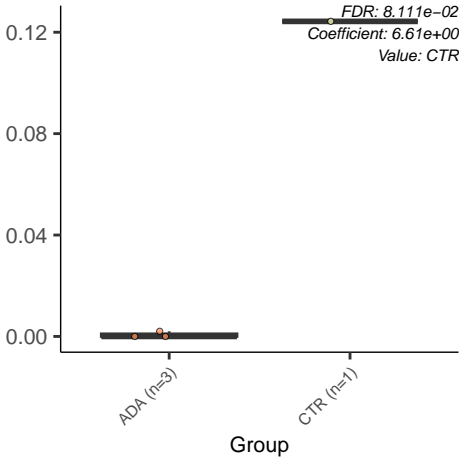
FDR: 7.132×10^{-29}

Coefficient: 1.00×10^0

Value: CTR



Faecal catena fissicatena



Lachnospiraceae_bacterium

0.15

0.10

0.05

0.00

ADA (n=3)

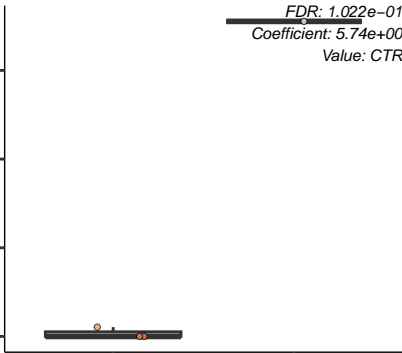
CTR (n=1)

Group

FDR: 1.022e-01

Coefficient: 5.74e+00

Value: CTR



Clostridiales_bacterium_KLE1615

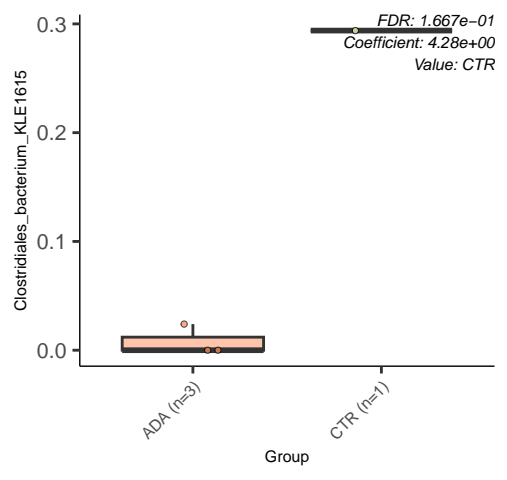
0.3
0.2
0.1
0.0

ADA (n=3)

CTR (n=1)

Group

FDR: 1.667e-01
Coefficient: 4.28e+00
Value: CTR



Collinsella_SGB14861

0.100
0.075
0.050
0.025
0.000

ADA (n=3)

CTR (n=1)

Group

FDR: 1.667e-01
Coefficient: 4.33e+00
Value: CTR

