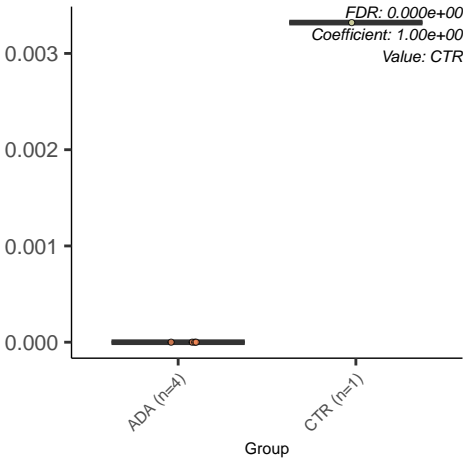


Streptococcus_thermophilus



Clostridium_sp_AF20_17LB

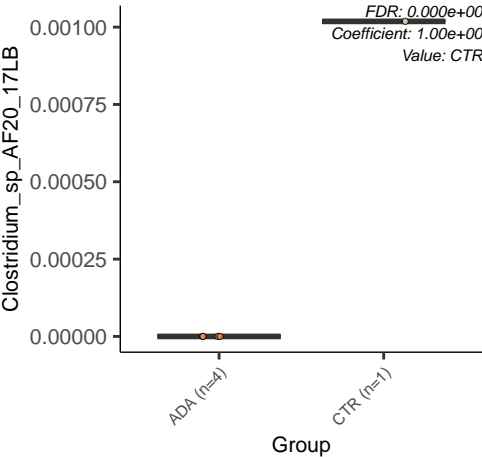
0.00100
0.00075
0.00050
0.00025
0.00000

ADA (n=4)

CTR (n=1)

Group

FDR: 0.000e+00
Coefficient: 1.00e+00
Value: CTR



Flavonifractor_plautii

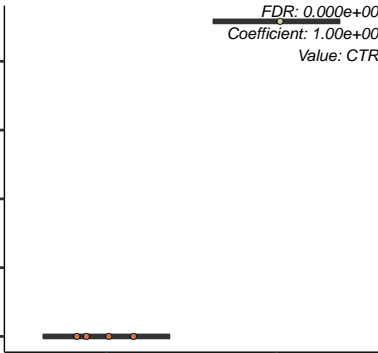
0.0020
0.0015
0.0010
0.0005
0.0000

ADA (n=4)

CTR (n=1)

Group

FDR: 0.000e+00
Coefficient: 1.00e+00
Value: CTR



Lacrimispora_saccharolytica

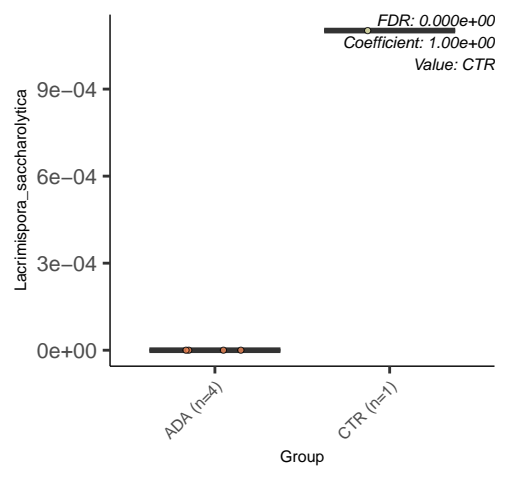
9e-04
6e-04
3e-04
0e+00

ADA (n=4)

CTR (n=1)

Group

FDR: 0.000e+00
Coefficient: 1.00e+00
Value: CTR



Evtepia_gabavorous

0.002

0.001

0.000

ADA (n=4)

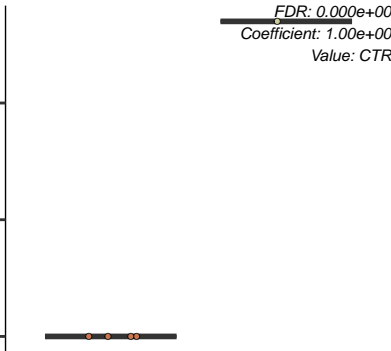
CTR (n=1)

Group

FDR: 0.000e+00

Coefficient: 1.00e+00

Value: CTR



Faecalimonas_umbilicata

6e-04

4e-04

2e-04

0e+00

ADA (n=4)

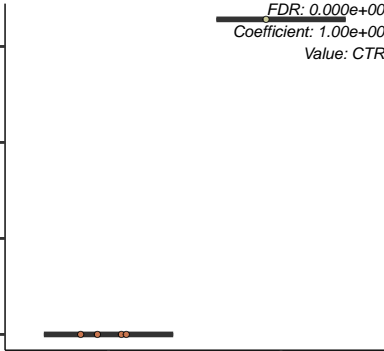
CTR (n=1)

Group

FDR: 0.000e+00

Coefficient: 1.00e+00

Value: CTR



Bacteroides_stercoris

0.2

0.1

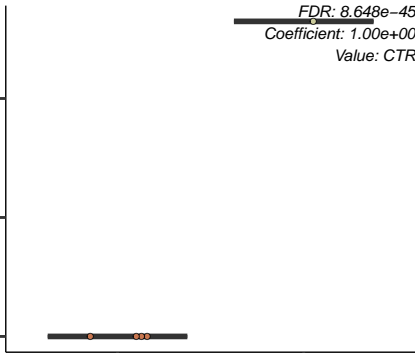
0.0

ADA (n=4)

CTR (n=1)

Group

FDR: 8.648e-45
Coefficient: 1.00e+00
Value: CTR



Dysosmobacter_welbionis

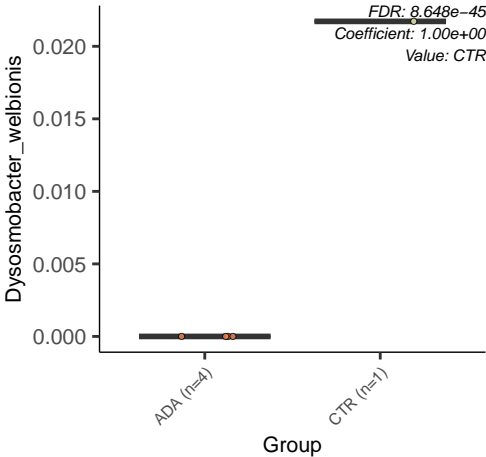
0.020
0.015
0.010
0.005
0.000

ADA (n=4)

CTR (n=1)

Group

FDR: 8.648e-45
Coefficient: 1.00e+00
Value: CTR



GGB3109_SGB4119

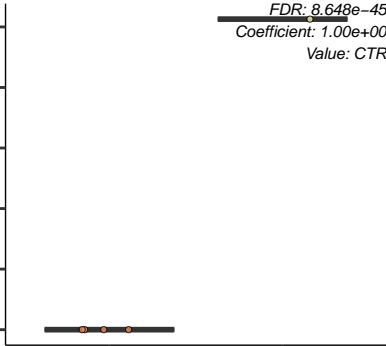
0.005
0.004
0.003
0.002
0.001
0.000

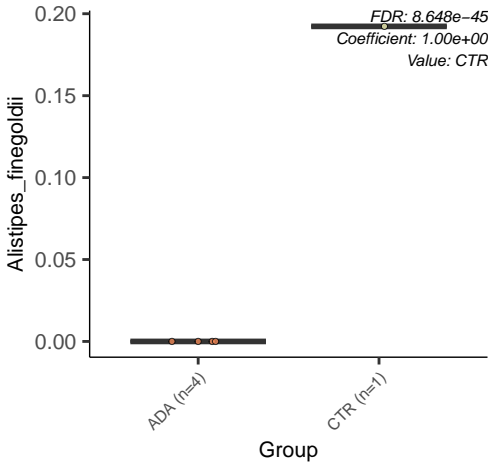
ADA (n=4)

CTR (n=1)

Group

FDR: 8.648e-45
Coefficient: 1.00e+00
Value: CTR





Bacteroides_caccae

0.2

0.1

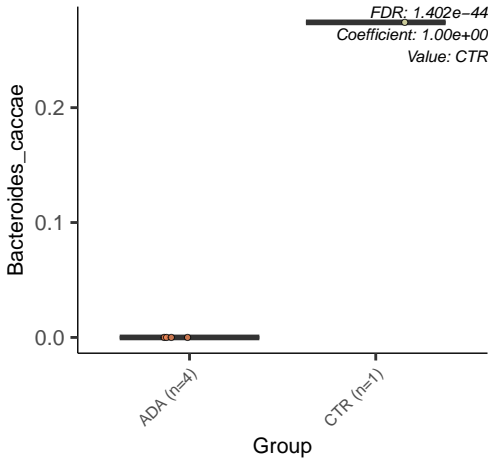
0.0

ADA (n=4)

CTR (n=1)

Group

FDR: 1.402e-44
Coefficient: 1.00e+00
Value: CTR



Odoribacter_splanchnicus

0.00

0.02

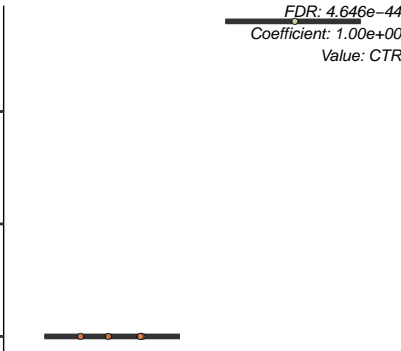
0.04

ADA (n=4)

CTR (n=1)

Group

FDR: 4.646e-44
Coefficient: 1.00e+00
Value: CTR



Parasutterella_SGB9260

0.00

0.02

0.04

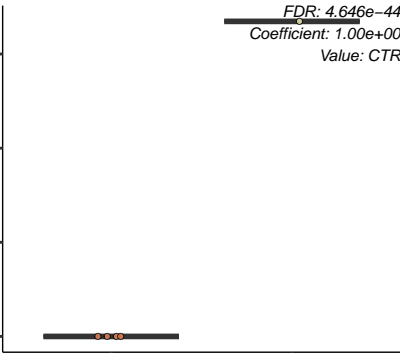
0.06

ADA (n=4)

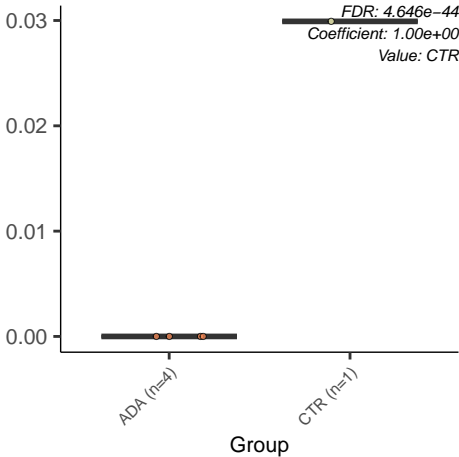
CTR (n=1)

Group

FDR: 4.646e-44
Coefficient: 1.00e+00
Value: CTR



Alistipes_communis



Clostridiales_bacterium

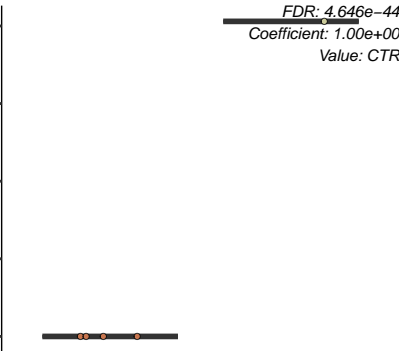
0.04
0.03
0.02
0.01
0.00

ADA (n=4)

CTR (n=1)

Group

FDR: 4.646e-44
Coefficient: 1.00e+00
Value: CTR



Faecalibacterium_SGB15315

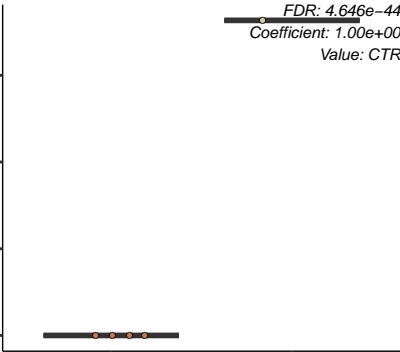
0.06
0.04
0.02
0.00

ADA (n=4)

CTR (n=1)

Group

FDR: 4.646e-44
Coefficient: 1.00e+00
Value: CTR



Lawsonibacter_asaccharolyticus

0.02

0.01

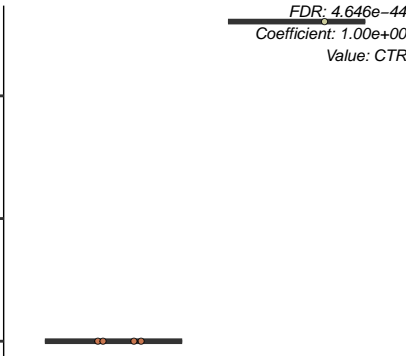
0.00

ADA (n=4)

CTR (n=1)

Group

FDR: 4.646e-44
Coefficient: 1.00e+00
Value: CTR



GGB9619_SGB15067

2e-04

1e-04

0e+00

ADA (n=4)

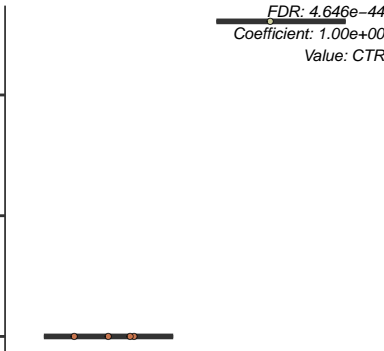
CTR (n=1)

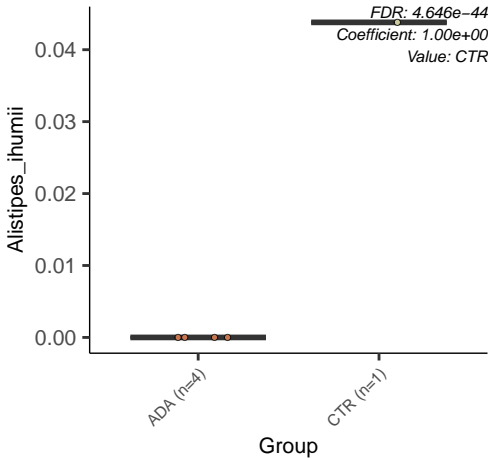
Group

FDR: 4.646e-44

Coefficient: 1.00e+00

Value: CTR





Ruminococcus_lactaris

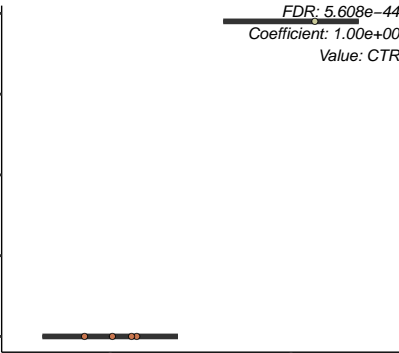
0.12
0.09
0.06
0.03
0.00

ADA (n=4)

CTR (n=1)

Group

FDR: 5.608e-44
Coefficient: 1.00e+00
Value: CTR



Bacteroides_finegoldii

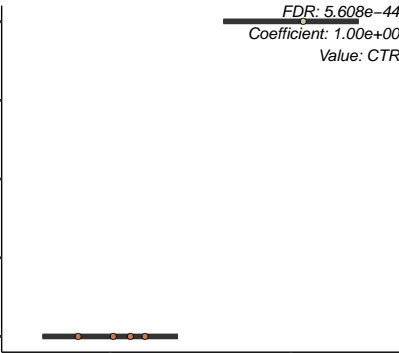
0.08
0.06
0.04
0.02
0.00

ADA (n=4)

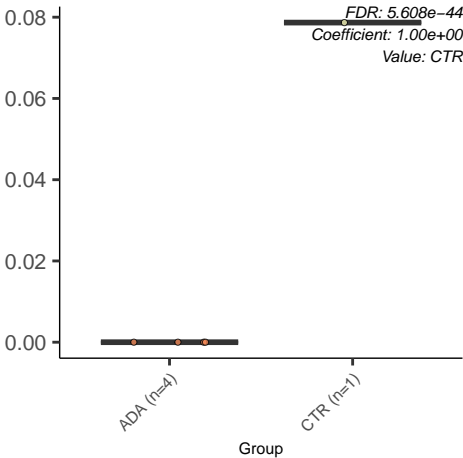
CTR (n=1)

Group

FDR: 5.608e-44
Coefficient: 1.00e+00
Value: CTR



Bacteroides_thetaiotaomicron



Clostridium_fessum

3e-04

2e-04

1e-04

0e+00

FDR: 2.522e-43

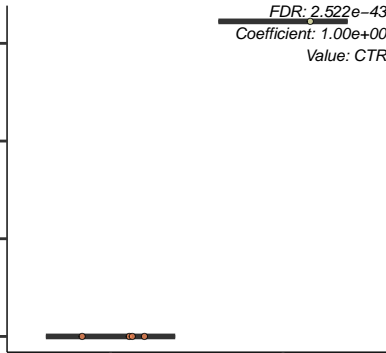
Coefficient: 1.00e+00

Value: CTR

ADA (n=4)

CTR (n=1)

Group



Parasutterella_excrementihominis

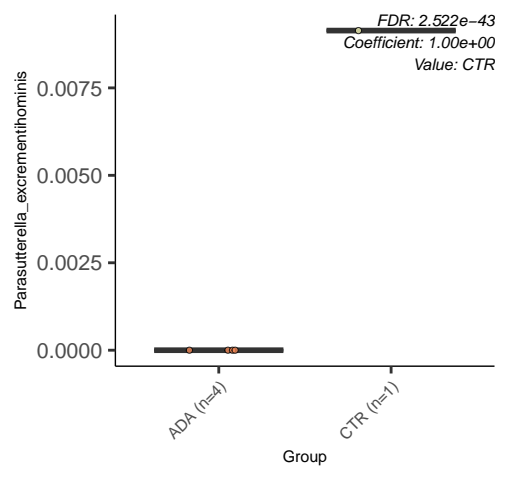
0.0075
0.0050
0.0025
0.0000

ADA (n=4)

CTR (n=1)

Group

FDR: 2.522e-43
Coefficient: 1.00e+00
Value: CTR



GGB2653_SGB3574

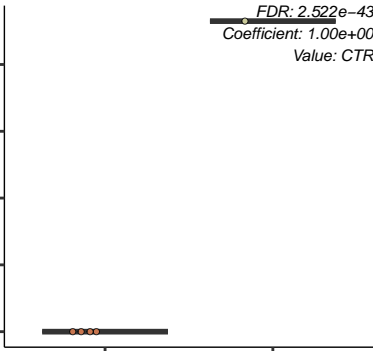
0.0020
0.0015
0.0010
0.0005
0.0000

ADA (n=4)

CTR (n=1)

Group

$FDR: 2.522e-43$
 $Coefficient: 1.00e+00$
 $Value: CTR$



Lacrimispora_amygdalina

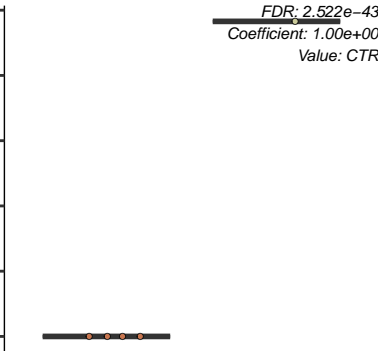
0.0025
0.0020
0.0015
0.0010
0.0005
0.0000

ADA (n=4)

CTR (n=1)

Group

FDR: 2.522e-43
Coefficient: 1.00e+00
Value: CTR



GGB9627_SGB15081

0.003

0.002

0.001

0.000

ADA (n=4)

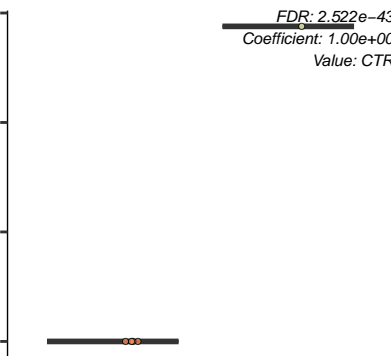
CTR (n=1)

Group

FDR: 2.522e-43

Coefficient: 1.00e+00

Value: CTR



GGB2998_SGB3988

0.00075
0.00050
0.00025
0.00000

ADA (n=4)

CTR (n=1)

Group

FDR: 2.522e-43
Coefficient: 1.00e+00
Value: CTR



Clostridium_SGB4750

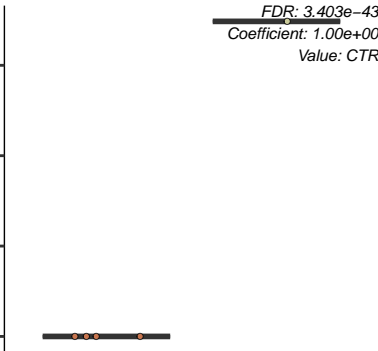
0.0075
0.0050
0.0025
0.0000

ADA (n=4)

CTR (n=1)

Group

FDR: 3.403e-43
Coefficient: 1.00e+00
Value: CTR



Blautia_sp_MSX_20_85

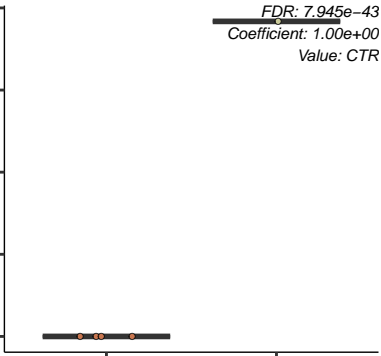
0.0100
0.0075
0.0050
0.0025
0.0000

ADA (n=4)

CTR (n=1)

Group

FDR: 7.945e-43
Coefficient: 1.00e+00
Value: CTR



Erysipelatoclostridium_ramosum

0e+00

1e-04

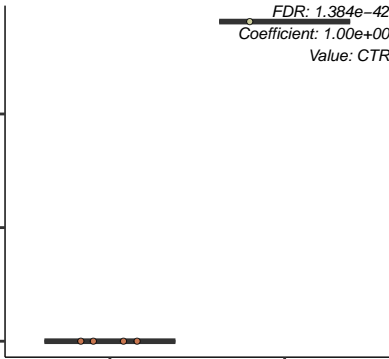
2e-04

ADA (n=4)

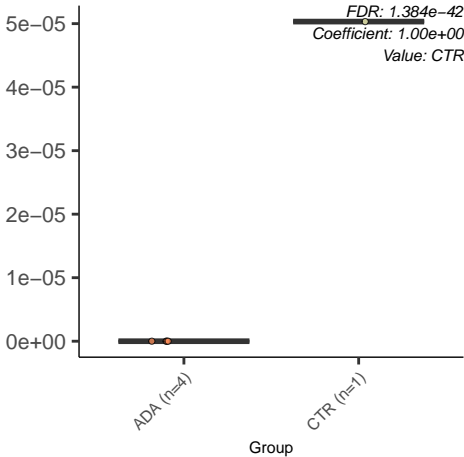
CTR (n=1)

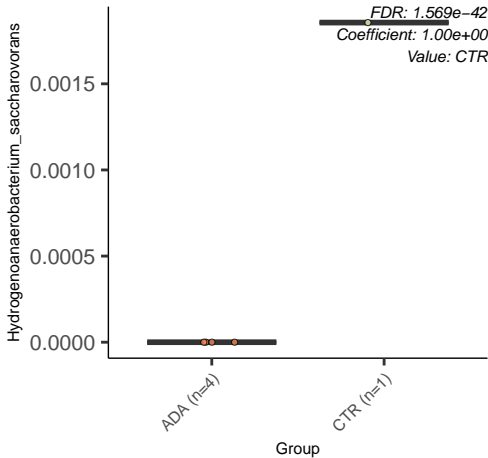
Group

FDR: $1.384e-42$
Coefficient: $1.00e+00$
Value: CTR



Eisenbergiella_massiliensis





Bilophila_wadsworthia

0.0015

0.0010

0.0005

0.0000

ADA (n=4)

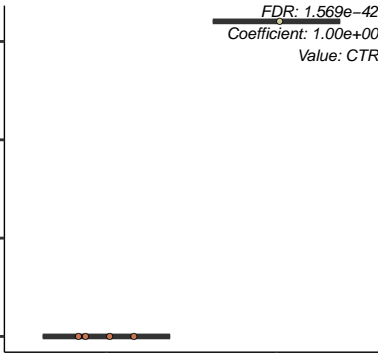
CTR (n=1)

Group

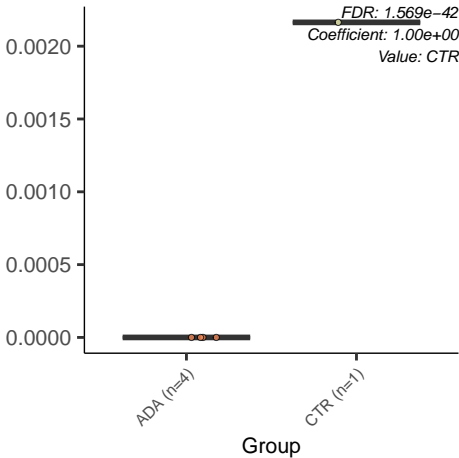
FDR: 1.569e-42

Coefficient: 1.00e+00

Value: CTR



Rheinheimer_SGB14999



Lachnospira_pectinoschiza

6e-04

4e-04

2e-04

0e+00

ADA (n=4)

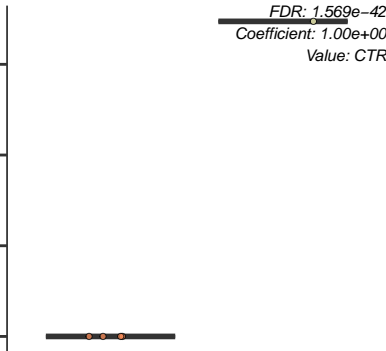
CTR (n=1)

Group

FDR: 1.569e-42

Coefficient: 1.00e+00

Value: CTR



Enterocloster_bolteae

0e+00

2e-04

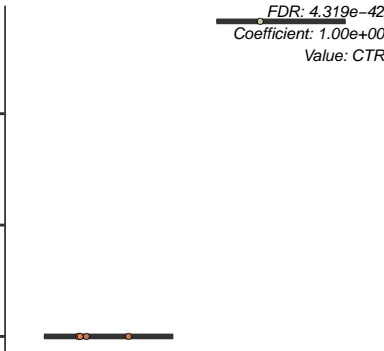
4e-04

ADA (n=4)

CTR (n=1)

Group

FDR: 4.319e-42
Coefficient: 1.00e+00
Value: CTR



Blautia_massiliensis

0.10

0.05

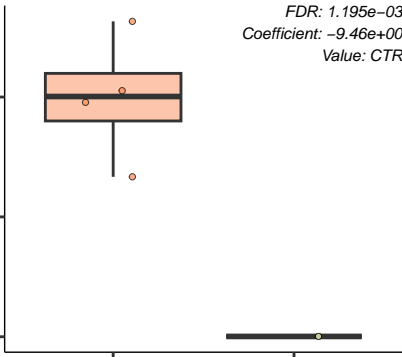
0.00

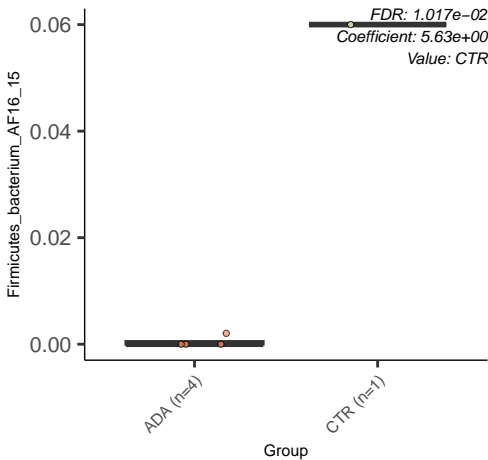
ADA (n=4)

CTR (n=1)

Group

FDR: 1.195e-03
Coefficient: -9.46e+00
Value: CTR





Phocaeicola_vulgatus

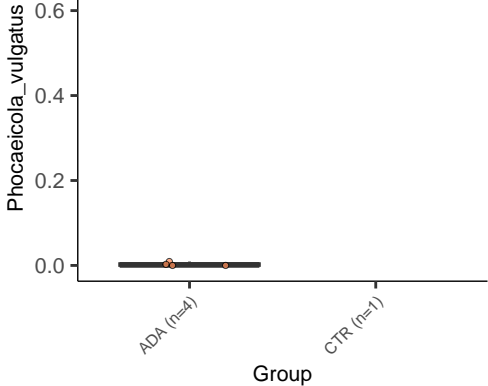
0.6
0.4
0.2
0.0

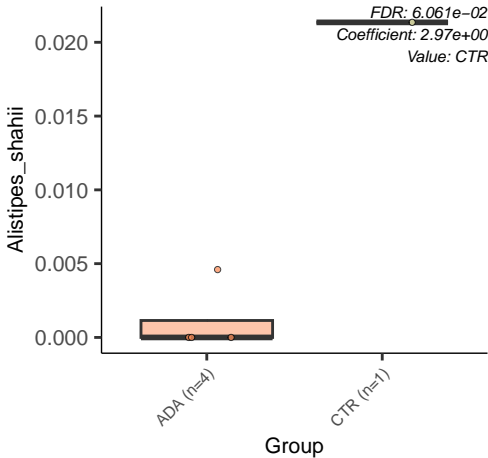
ADA (n=4)

CTR (n=1)

Group

FDR: 5.739e-02
Coefficient: 8.24e+00
Value: CTR





Fusicatenibacter_saccharivorans

FDR: 7.549e-02
Coefficient: -4.41e+00
Value: CTR

0.10

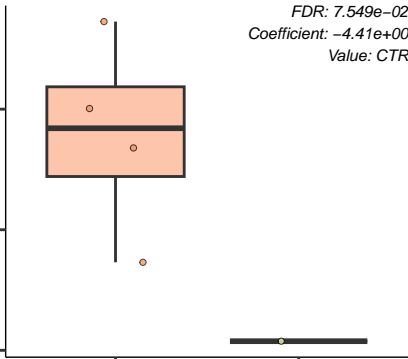
0.05

0.00

ADA (n=4)

CTR (n=1)

Group



Clostridiaceae_bacterium_AF18_31LB

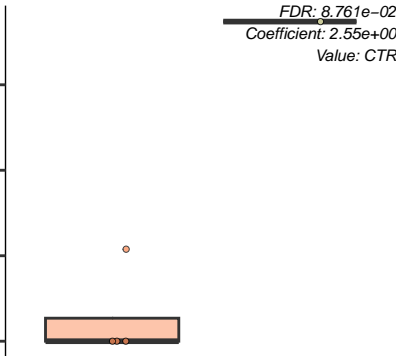
0.015
0.010
0.005
0.000

ADA (n=4)

CTR (n=1)

Group

FDR: 8.761e-02
Coefficient: 2.55e+00
Value: CTR



Bacteroides_ovatus

0.03

0.02

0.01

0.00

ADA (n=4)

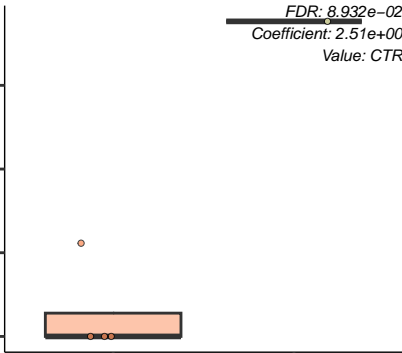
CTR (n=1)

Group

FDR: 8.932e-02

Coefficient: 2.51e+00

Value: CTR



Clostridiaceae_bacterium

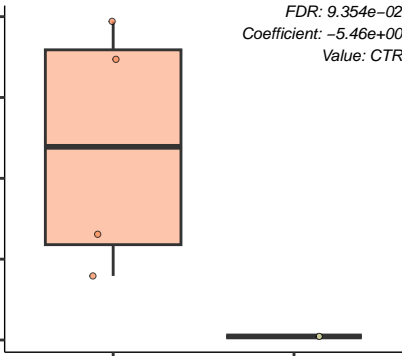
0.08
0.06
0.04
0.02
0.00

FDR: 9.354e-02
Coefficient: -5.46e+00
Value: CTR

ADA (n=4)

CTR (n=1)

Group



Clostridium_sp_AF36_4

0.003
0.002
0.001
0.000

ADA (n=4)

CTR (n=1)

Group

FDR: 1.525e-01
Coefficient: 2.01e+00
Value: CTR

