

Clostridium_sp_AF20_17LB

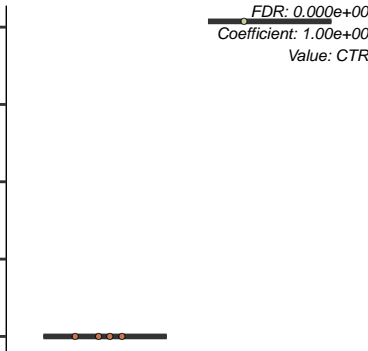
0.00100
0.00075
0.00050
0.00025
0.00000

ADA (n=4)

CTR (n=1)

Group

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



Lacrimispora_amygdalina

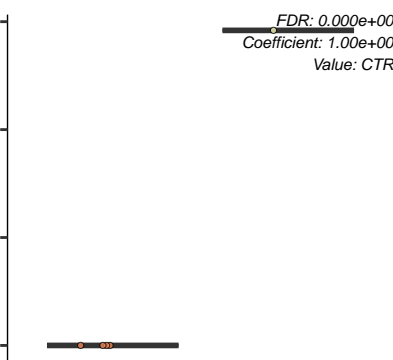
0.003
0.002
0.001
0.000

ADA (n=4)

CTR (n=1)

Group

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



GGB2998_SGB3988

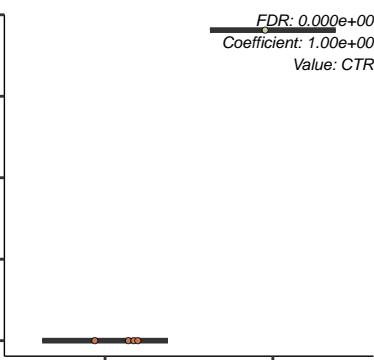
0.0012
0.0009
0.0006
0.0003
0.0000

ADA (n=4)

CTR (n=1)

Group

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



Rheinheimera_SGB14999

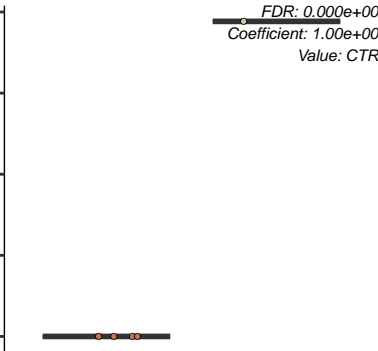
0.0020
0.0015
0.0010
0.0005
0.0000

ADA (n=4)

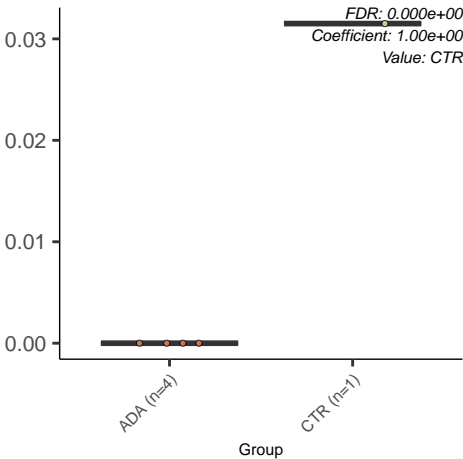
CTR (n=1)

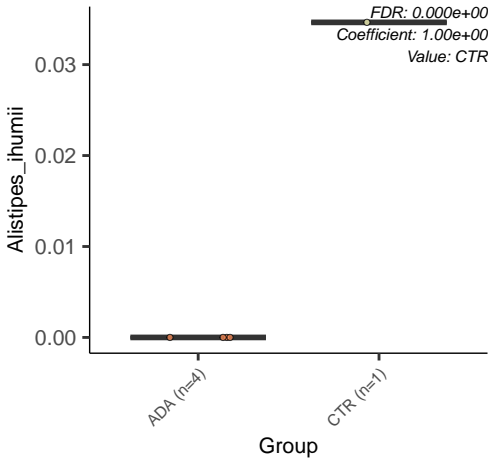
Group

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



Lawsonibacter_asaccharolyticus





Eisenbergiella_massiliensis

4e-05

3e-05

2e-05

1e-05

0e+00

ADA (n=4)

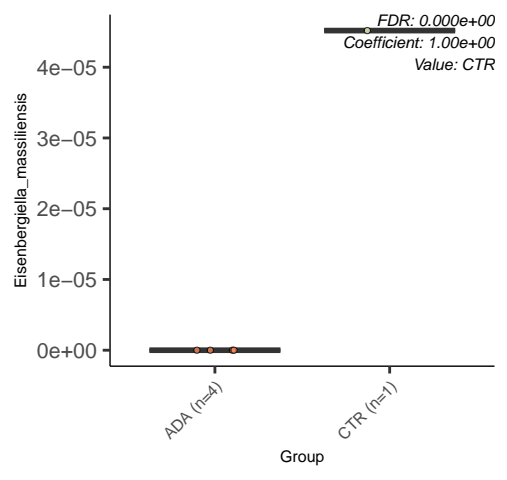
CTR (n=1)

Group

FDR: 0.000e+00

Coefficient: 1.00e+00

Value: CTR



Bacteroides_caccae

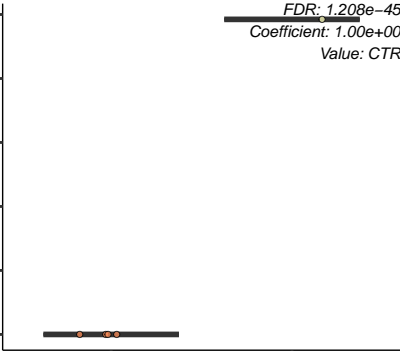
0.25
0.20
0.15
0.10
0.05
0.00

ADA (n=4)

CTR (n=1)

Group

FDR: 1.208e-45
Coefficient: 1.00e+00
Value: CTR



Ruminococcus_lactaris

0.10

0.05

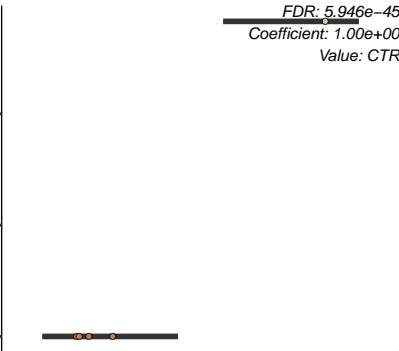
0.00

ADA (n=4)

CTR (n=1)

Group

FDR: 5.946e-45
Coefficient: 1.00e+00
Value: CTR



Parasutterella_SGB9260

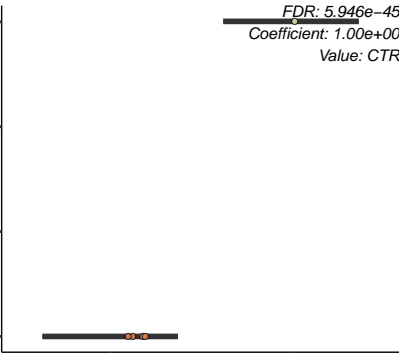
0.06
0.04
0.02
0.00

ADA (n=4)

CTR (n=1)

Group

FDR: 5.946e-45
Coefficient: 1.00e+00
Value: CTR



Alistipes_communis

0.020

0.015

0.010

0.005

0.000

ADA (n=4)

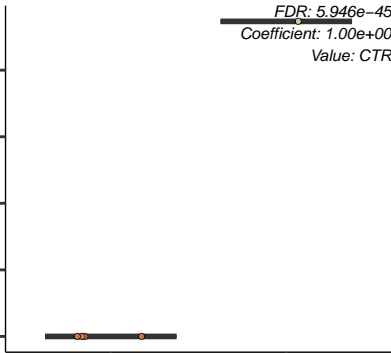
CTR (n=1)

Group

FDR: 5.946e-45

Coefficient: 1.00e+00

Value: CTR



Dysosmobacter_welbionis

0.00

0.01

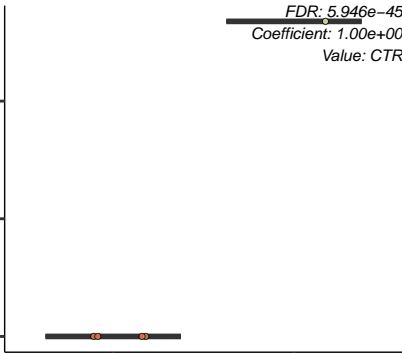
0.02

ADA (n=4)

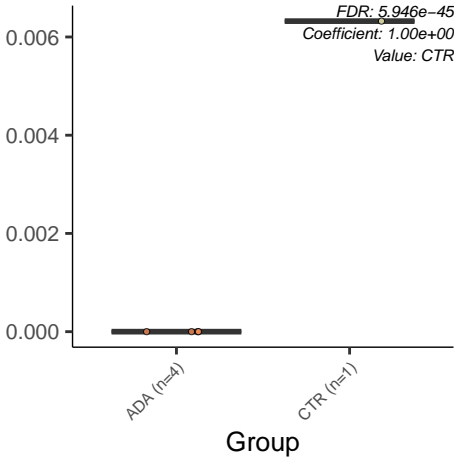
CTR (n=1)

Group

FDR: 5.946e-45
Coefficient: 1.00e+00
Value: CTR



GGB3109_SGB4119



Alistipes_finegoldii

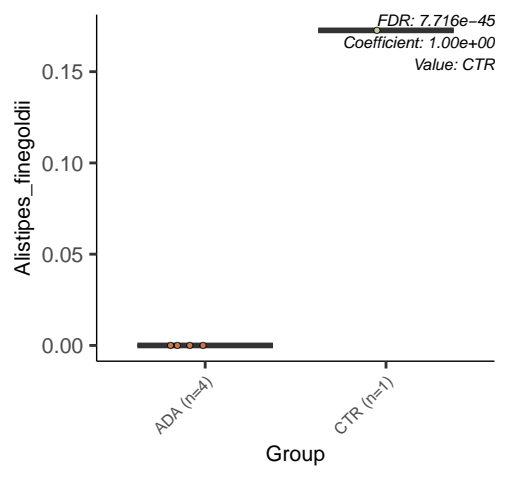
0.15
0.10
0.05
0.00

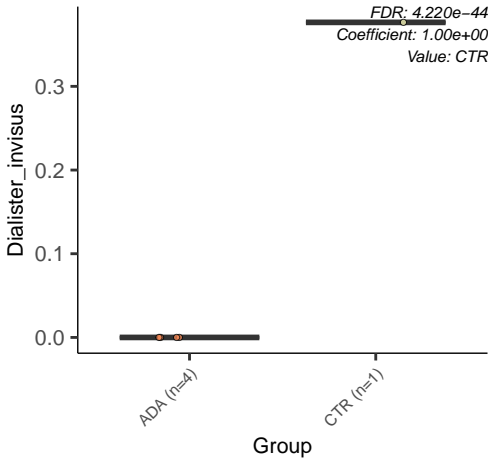
ADA (n=4)

CTR (n=1)

Group

FDR: 7.716e-45
Coefficient: 1.00e+00
Value: CTR





Odoribacter_splanchnicus

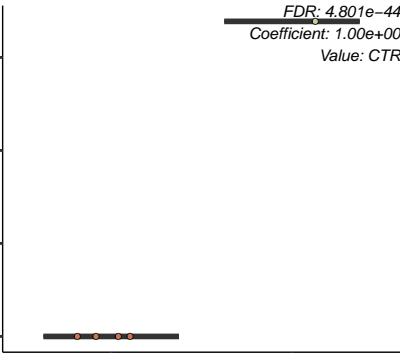
0.06
0.04
0.02
0.00

ADA (n=4)

CTR (n=1)

Group

FDR: 4.801e-44
Coefficient: 1.00e+00
Value: CTR



Clostridiales_bacterium

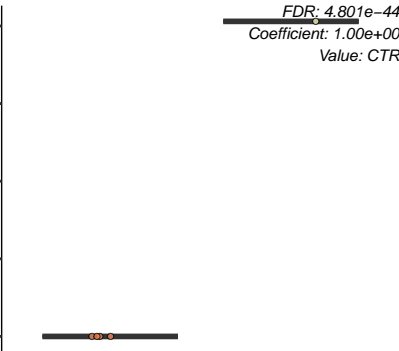
0.04
0.03
0.02
0.01
0.00

ADA (n=4)

CTR (n=1)

Group

FDR: 4.801e-44
Coefficient: 1.00e+00
Value: CTR



Faecalibacterium_SGB15315

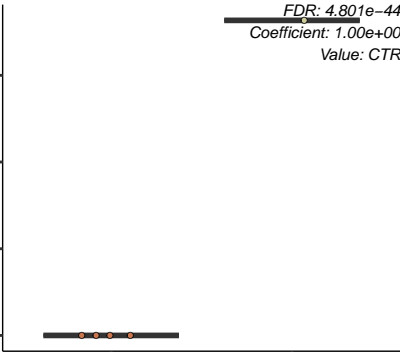
0.06
0.04
0.02
0.00

ADA (n=4)

CTR (n=1)

Group

FDR: 4.801e-44
Coefficient: 1.00e+00
Value: CTR



Bacteroides_finegoldii

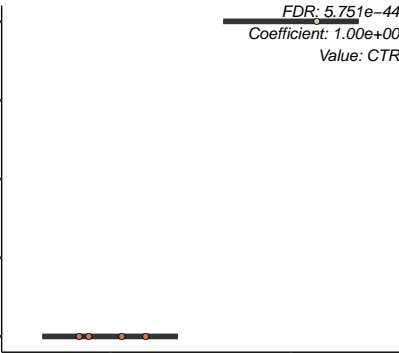
0.08
0.06
0.04
0.02
0.00

ADA (n=4)

CTR (n=1)

Group

FDR: 5.751e-44
Coefficient: 1.00e+00
Value: CTR



Bacteroides_stercoris

0.20

0.15

0.10

0.05

0.00

ADA (n=4)

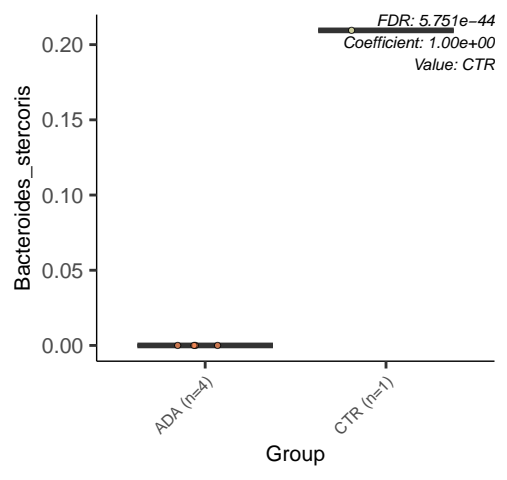
CTR (n=1)

Group

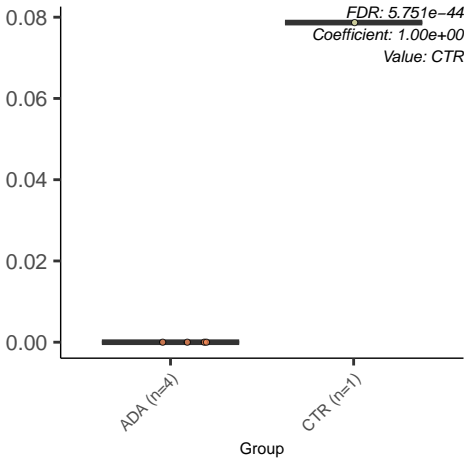
FDR: 5.751e-44

Coefficient: 1.00e+00

Value: CTR



Bacteroides_thetaiotaomicron



Clostridium_fessum

2e-04

1e-04

0e+00

ADA (n=4)

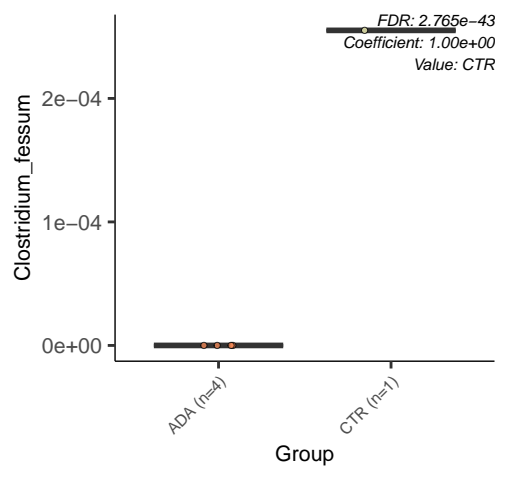
CTR (n=1)

Group

FDR: 2.765e-43

Coefficient: 1.00e+00

Value: CTR



Parasutterella_excrementihominis

0.009

0.006

0.003

0.000

ADA (n=4)

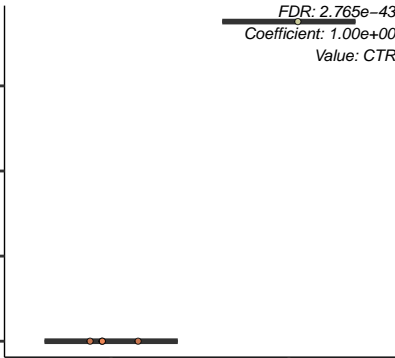
CTR (n=1)

Group

FDR: 2.765e-43

Coefficient: 1.00e+00

Value: CTR



GGB2653_SGB3574

0.002

0.001

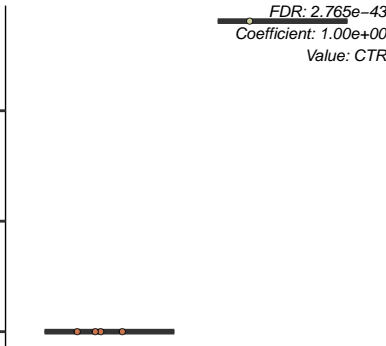
0.000

ADA (n=4)

CTR (n=1)

Group

FDR: 2.765e-43
Coefficient: 1.00e+00
Value: CTR



Lachnospira_pectinoschiza

8e-04

6e-04

4e-04

2e-04

0e+00

FDR: 2.765e-43

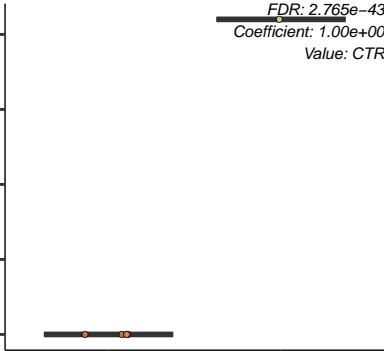
Coefficient: 1.00e+00

Value: CTR

ADA (n=4)

CTR (n=1)

Group



Blautia_sp_MSX_20_85

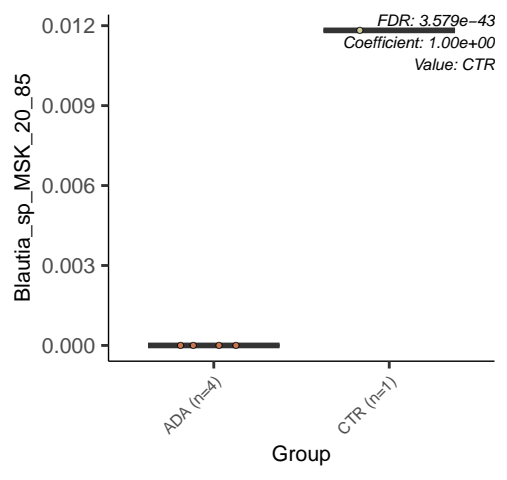
0.012
0.009
0.006
0.003
0.000

ADA (n=4)

CTR (n=1)

Group

FDR: 3.579e-43
Coefficient: 1.00e+00
Value: CTR



Clostridium_SGB4750

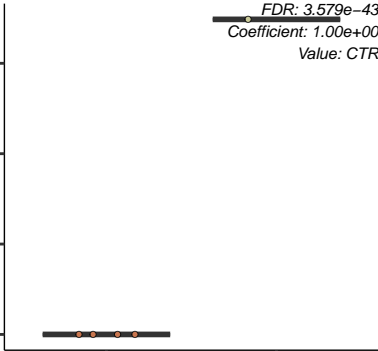
0.0075
0.0050
0.0025
0.0000

ADA (n=4)

CTR (n=1)

Group

FDR: 3.579e-43
Coefficient: 1.00e+00
Value: CTR



GGB9627_SGB15081

0.000

0.001

0.002

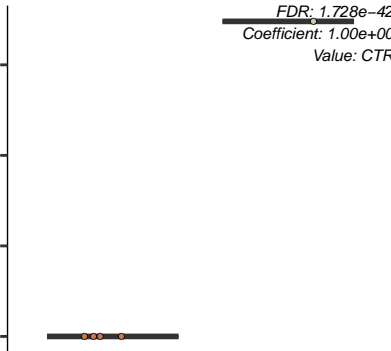
0.003

ADA (n=4)

CTR (n=1)

Group

FDR: 1.728e-42
Coefficient: 1.00e+00
Value: CTR



Flavonifractor_plautii

0.002

0.001

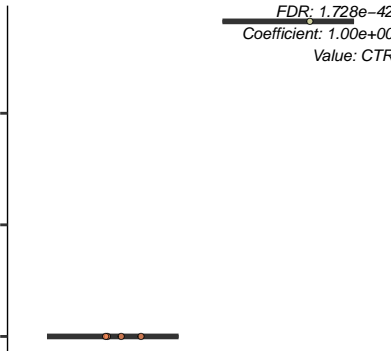
0.000

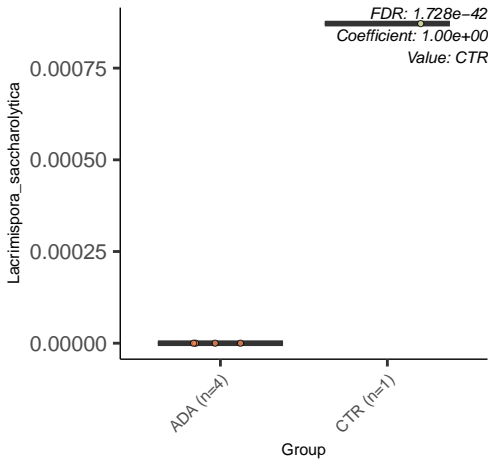
ADA (n=4)

CTR (n=1)

Group

FDR: $1.728e-42$
Coefficient: $1.00e+00$
Value: CTR





Erysipelatoclostridium_ramosum

0.00020

0.00015

0.00010

0.00005

0.00000

ADA (n=4)

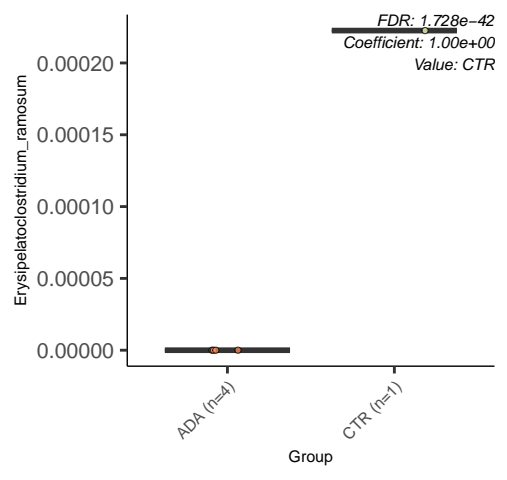
CTR (n=1)

Group

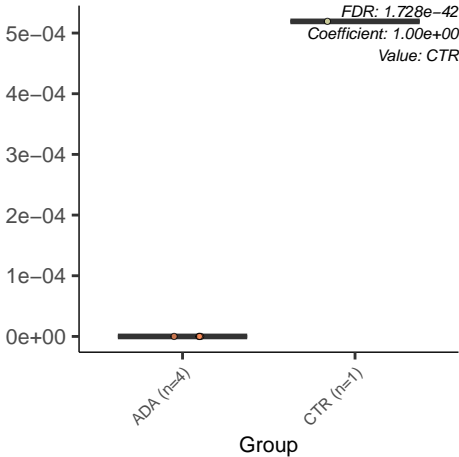
FDR: 1.728e-42

Coefficient: 1.00e+00

Value: CTR



Faecalimonas_umbilicata



Enterocloster_bolteae

0e+00

2e-04

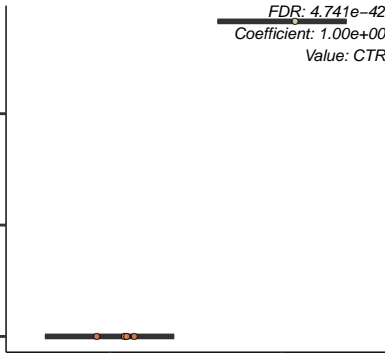
4e-04

ADA (n=4)

CTR (n=1)

Group

FDR: 4.741e-42
Coefficient: 1.00e+00
Value: CTR



GGB9619_SGB15067

3e-04

2e-04

1e-04

0e+00

FDR: 1.134e-41

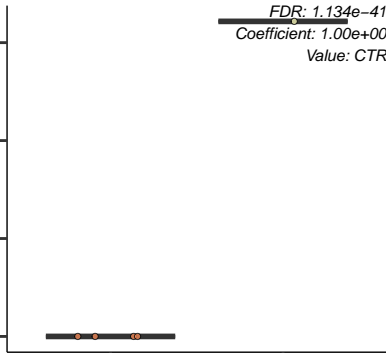
Coefficient: 1.00e+00

Value: CTR

ADA (n=4)

CTR (n=1)

Group



Blautia_massiliensis

0.10

0.05

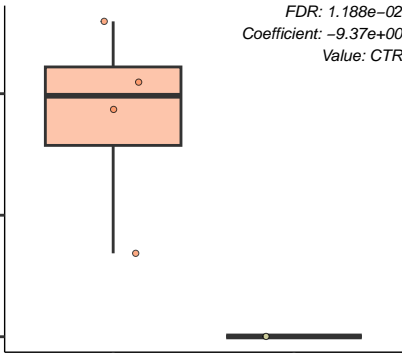
0.00

ADA (n=4)

CTR (n=1)

Group

FDR: $1.188e-02$
Coefficient: $-9.37e+00$
Value: CTR



Fusicatenibacter_saccharivorans

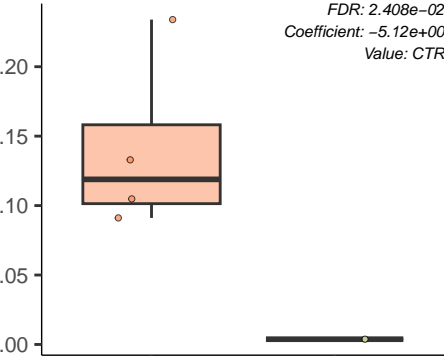
FDR: 2.408e-02
Coefficient: -5.12e+00
Value: CTR

ADA (n=4)

CTR (n=1)

Group

0.00
0.05
0.10
0.15
0.20



Bacteroides_ovatus

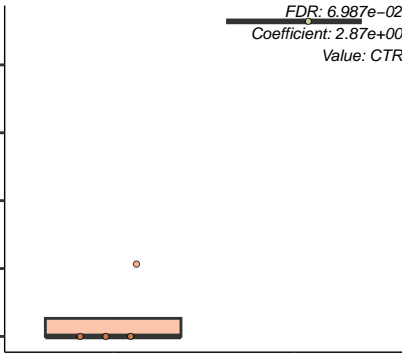
0.04
0.03
0.02
0.01
0.00

ADA (n=4)

CTR (n=1)

Group

FDR: 6.987e-02
Coefficient: 2.87e+00
Value: CTR



Clostridiaceae_bacterium_AF18_31LB

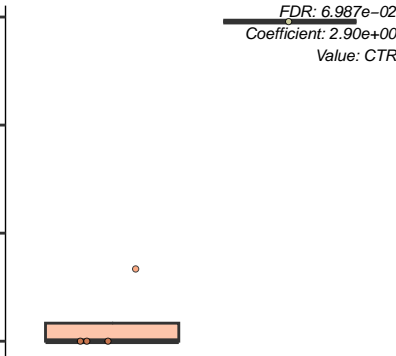
0.015
0.010
0.005
0.000

ADA (n=4)

CTR (n=1)

Group

FDR: 6.987e-02
Coefficient: 2.90e+00
Value: CTR



Clostridiaceae_bacterium

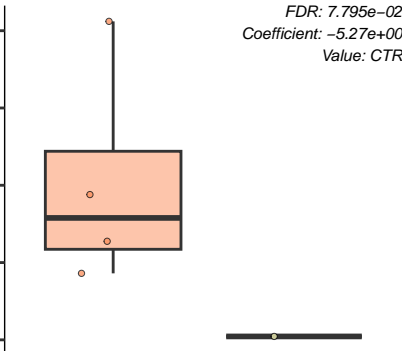
0.08
0.06
0.04
0.02
0.00

FDR: 7.795e-02
Coefficient: -5.27e+00
Value: CTR

ADA (n=4)

CTR (n=1)

Group



Phocaeicola_vulgatus

0.0

0.2

0.4

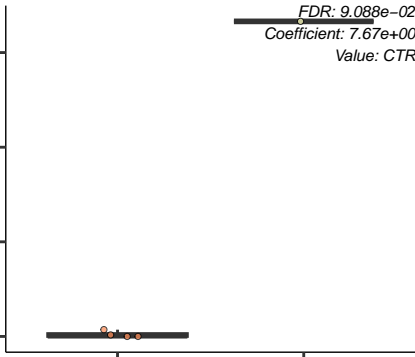
0.6

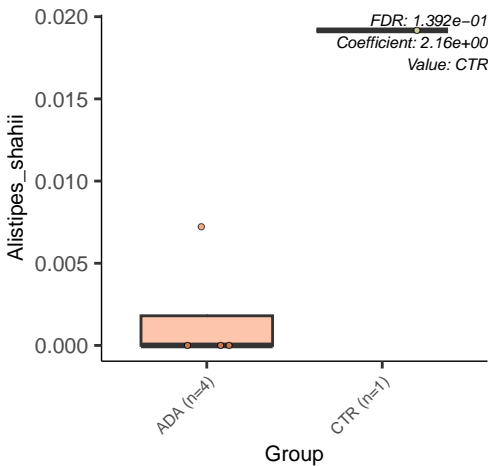
ADA (n=4)

CTR (n=1)

Group

FDR: 9.088e-02
Coefficient: 7.67e+00
Value: CTR





Eubacterium_rectale

0.5
0.4
0.3
0.2
0.1

FDR: 1.690e-01
Coefficient: -2.52e+00
Value: CTR

ADA (n=4)

CTR (n=1)

Group

