

Ruminococcus\_lactaris

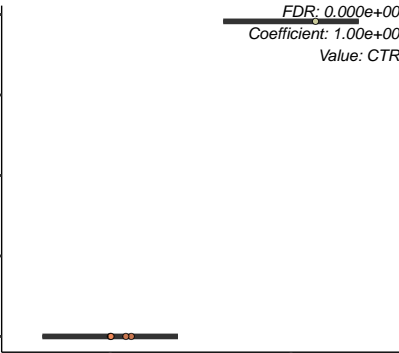
0.08  
0.06  
0.04  
0.02  
0.00

ADA (n=4)

CTR (n=1)

Group

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



Clostridium\_sp\_AF27\_2AA

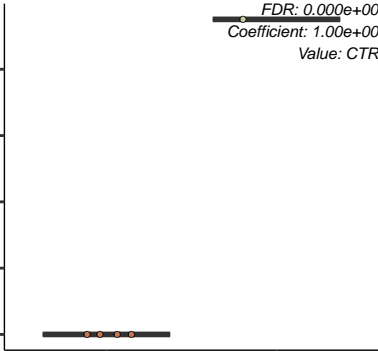
0.0020  
0.0015  
0.0010  
0.0005  
0.0000

ADA (n=4)

CTR (n=1)

Group

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



Collinsella\_SGB14861

0.09

0.06

0.03

0.00

ADA (n=4)

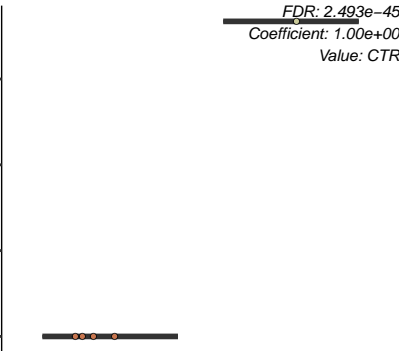
CTR (n=1)

Group

*FDR: 2.493e-45*

*Coefficient: 1.00e+00*

*Value: CTR*



Klebsiella\_pneumoniae

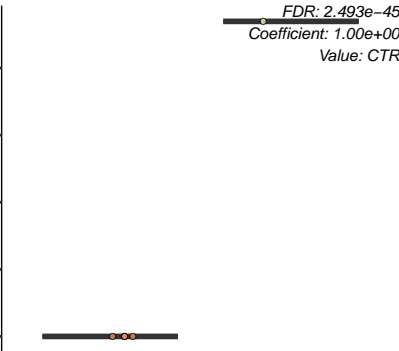
0.20  
0.15  
0.10  
0.05  
0.00

ADA (n=4)

CTR (n=1)

Group

*FDR: 2.493e-45*  
*Coefficient: 1.00e+00*  
*Value: CTR*



Faecalibacillus\_intestinalis

0.04

0.02

0.00

ADA (n=4)

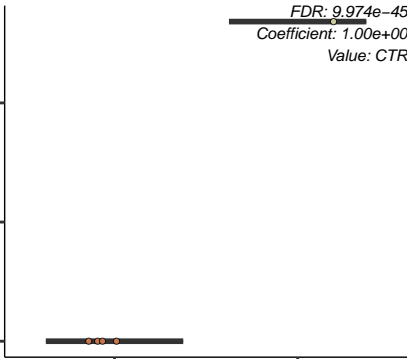
CTR (n=1)

Group

FDR:  $9.974 \times 10^{-45}$

Coefficient:  $1.00 \times 10^0$

Value: CTR



Staphylococcus\_SGB6340

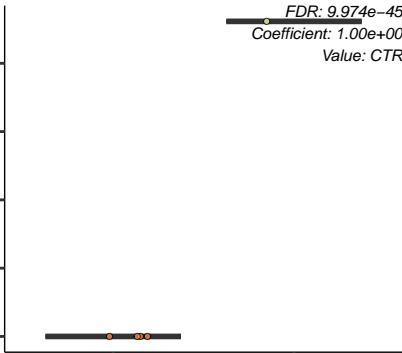
0.04  
0.03  
0.02  
0.01  
0.00

ADA (n=4)

CTR (n=1)

Group

*FDR: 9.974e-45*  
*Coefficient: 1.00e+00*  
*Value: CTR*



Blautia\_glucerasea

0.006

0.004

0.002

0.000

ADA (n=4)

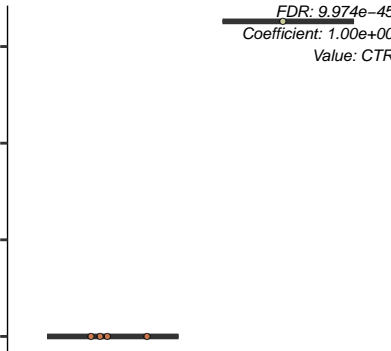
CTR (n=1)

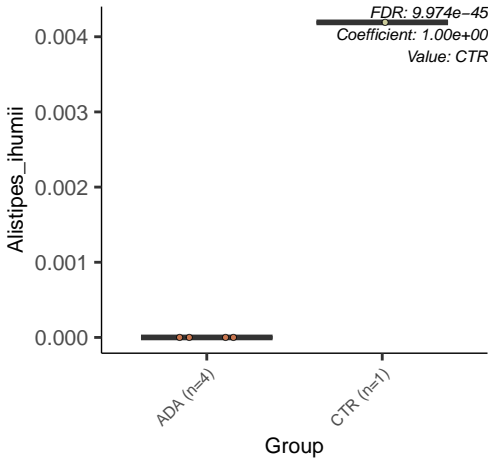
Group

FDR:  $9.974e-45$

Coefficient:  $1.00e+00$

Value: CTR







GGB3612\_SGB4882

0.04

0.02

0.00

ADA (n=4)

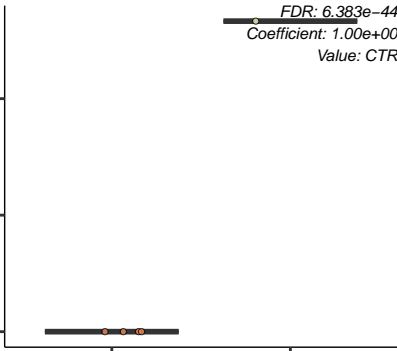
CTR (n=1)

Group

*FDR: 6.383e-44*

*Coefficient: 1.00e+00*

*Value: CTR*



Clostridium\_sp\_AF20\_17LB

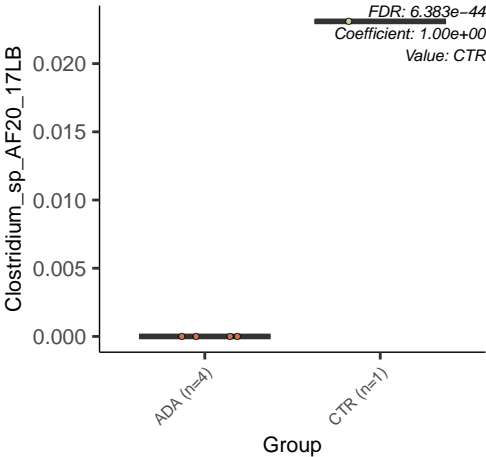
0.020  
0.015  
0.010  
0.005  
0.000

ADA (n=4)

CTR (n=1)

Group

*FDR: 6.383e-44*  
*Coefficient: 1.00e+00*  
*Value: CTR*



GGB9453\_SGB14845

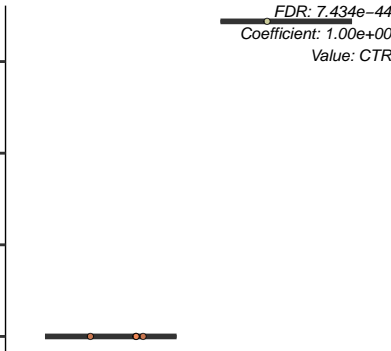
0.075  
0.050  
0.025  
0.000

ADA (n=4)

CTR (n=1)

Group

*FDR: 7.434e-44*  
*Coefficient: 1.00e+00*  
*Value: CTR*



Vescimonas\_SGB15087

0.000

0.025

0.050

0.075

*FDR: 7.434e-44*

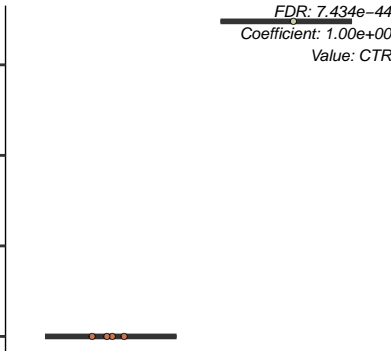
*Coefficient: 1.00e+00*

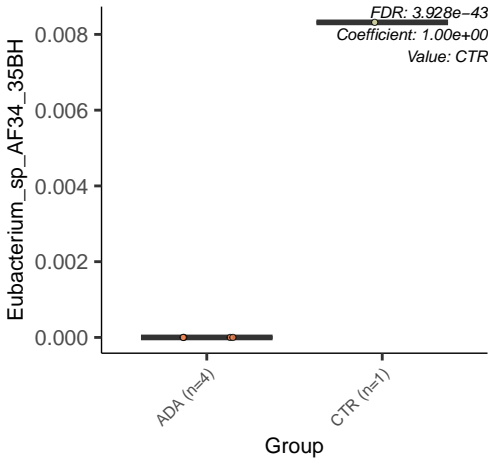
*Value: CTR*

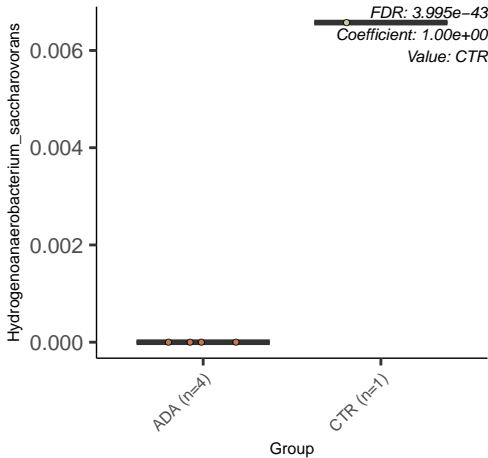
ADA (n=4)

CTR (n=1)

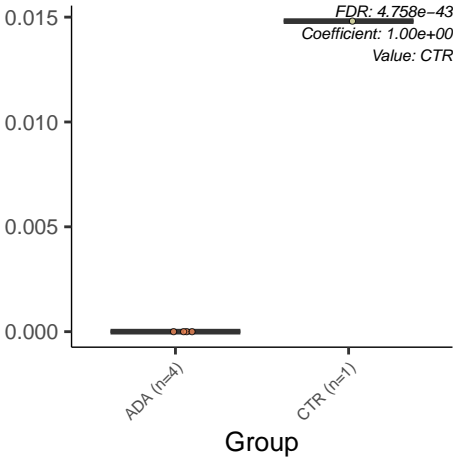
Group







GGB2653\_SGB3574



Eubacterium\_ramulus

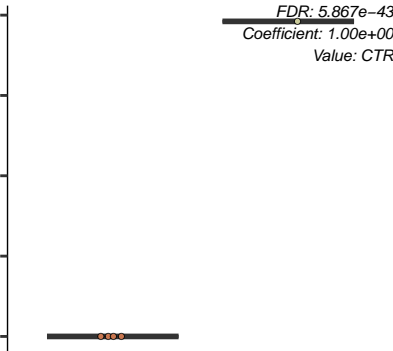
0.012  
0.009  
0.006  
0.003  
0.000

ADA (n=4)

CTR (n=1)

Group

*FDR: 5.867e-43*  
*Coefficient: 1.00e+00*  
*Value: CTR*





GGB9453\_SGB14844

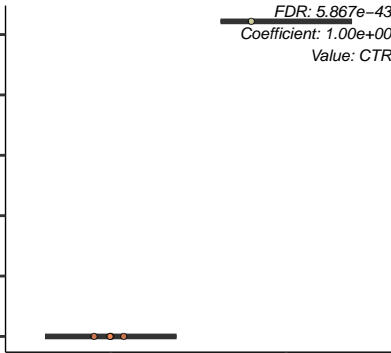
0.005  
0.004  
0.003  
0.002  
0.001  
0.000

ADA (n=4)

CTR (n=1)

Group

*FDR: 5.867e-43*  
*Coefficient: 1.00e+00*  
*Value: CTR*



GGB9619\_SGB15067

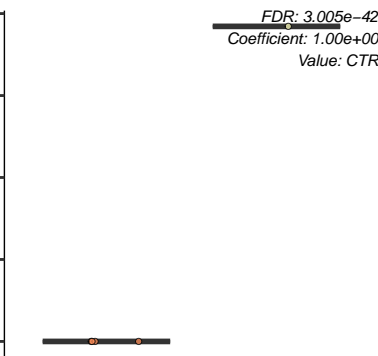
0.0020  
0.0015  
0.0010  
0.0005  
0.0000

ADA (n=4)

CTR (n=1)

Group

*FDR: 3.005e-42*  
*Coefficient: 1.00e+00*  
*Value: CTR*



GGB4566\_SGB6305

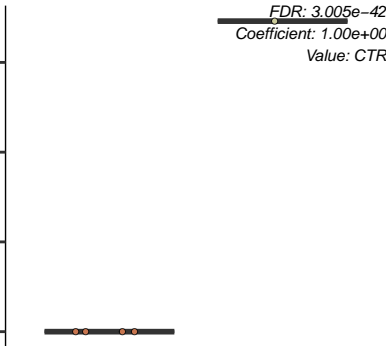
0.003  
0.002  
0.001  
0.000

ADA (n=4)

CTR (n=1)

Group

*FDR: 3.005e-42*  
*Coefficient: 1.00e+00*  
*Value: CTR*



GGB9708\_SGB15234

0.04

0.02

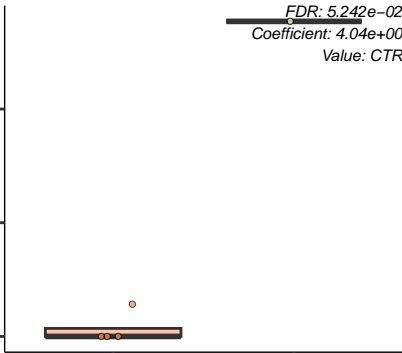
0.00

ADA (n=4)

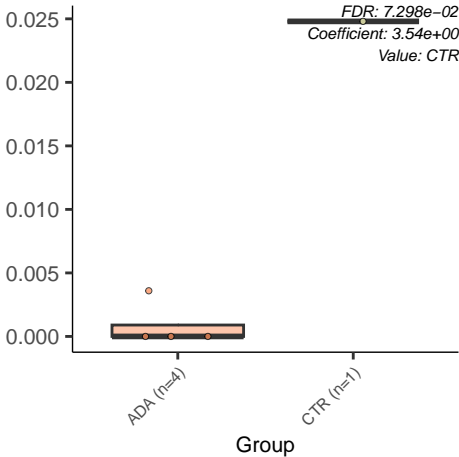
CTR (n=1)

Group

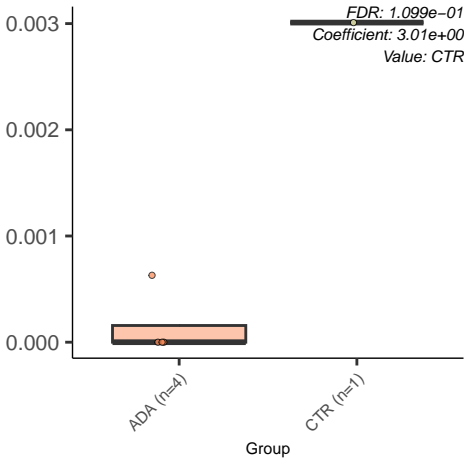
*FDR: 5.242e-02*  
*Coefficient: 4.04e+00*  
*Value: CTR*



Odoribacter\_splanchnicus



Adlercreutzia\_equolifaciens



Bacteroides\_cellulosilyticus

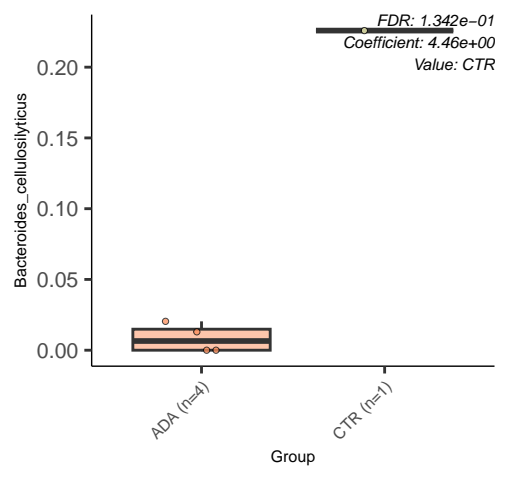
0.20  
0.15  
0.10  
0.05  
0.00

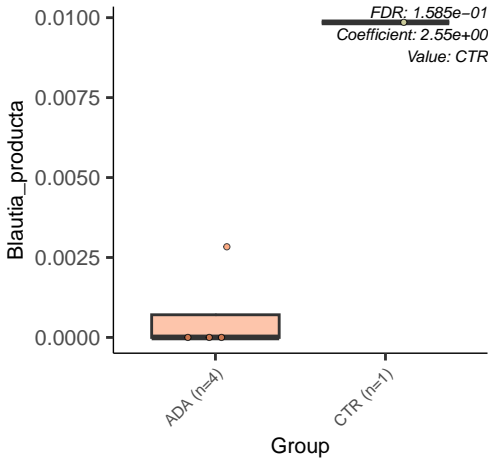
ADA (n=4)

CTR (n=1)

Group

*FDR: 1.342e-01*  
*Coefficient: 4.46e+00*  
*Value: CTR*







Firmicutes\_bacterium\_AF16\_15

0.09  
0.06  
0.03  
0.00

ADA (n=4)

CTR (n=1)

Group

*FDR: 2.028e-01*  
*Coefficient: 5.72e+00*  
*Value: CTR*

