

Faecalibacillus_intestinalis

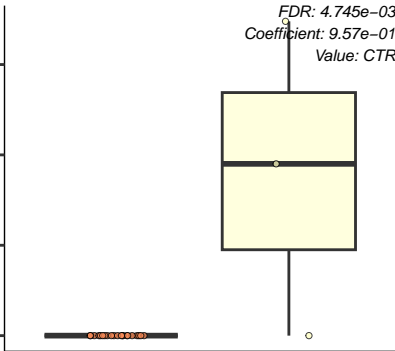
0.015
0.010
0.005
0.000

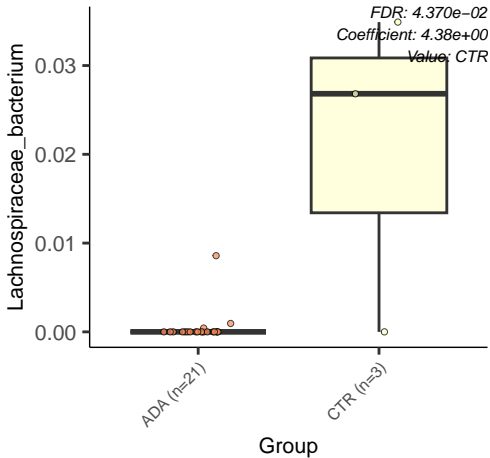
ADA (n=21)

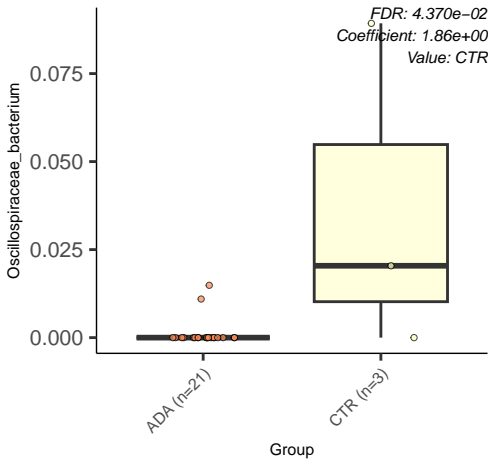
CTR (n=3)

Group

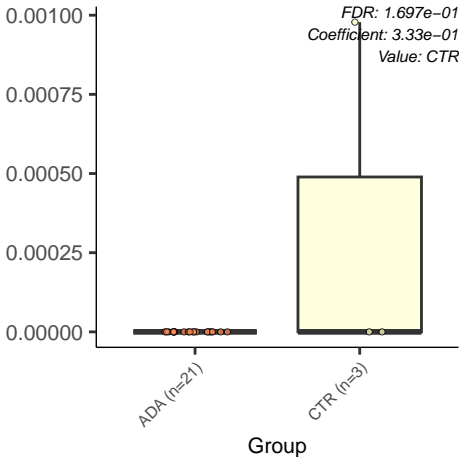
FDR: 4.745e-03
Coefficient: 9.57e-01
Value: CTR







Clostridium_SGB4909



Clostridium_sp_AF27_2AA

0.010

0.005

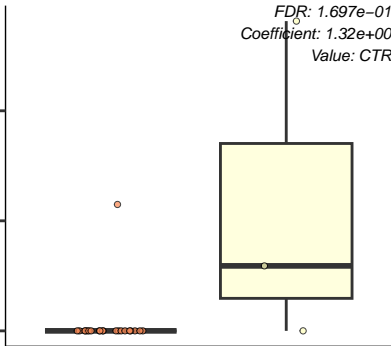
0.000

ADA (n=21)

CTR (n=3)

Group

FDR: 1.697e-01
Coefficient: 1.32e+00
Value: CTR



Clostridium_sp_NSJ_42

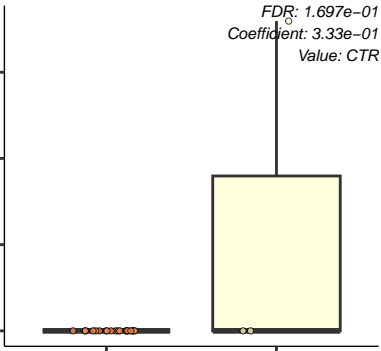
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0.0050
0.0025
0.0000

ADA (n=21)

CTR (n=3)

Group

FDR: 1.697e-01
Coefficient: 3.33e-01
Value: CTR



Vescimonas_SGB15087

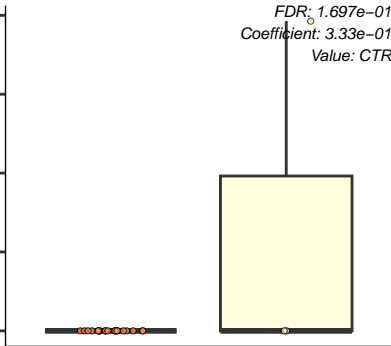
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0.003
0.002
0.001
0.000

FDR: 1.697e-01
Coefficient: 3.33e-01
Value: CTR

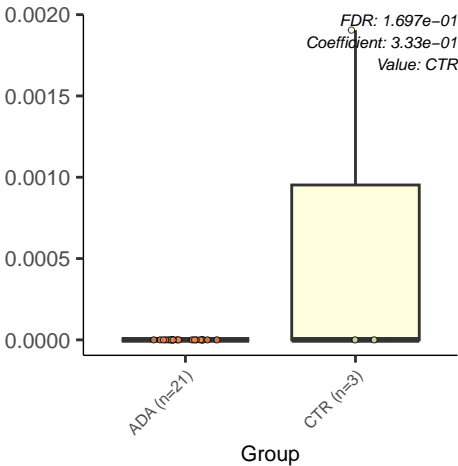
ADA (n=21)

CTR (n=3)

Group



GGB9453_SGB14844



GGB33512_SGB15201

0.0075

0.0050

0.0025

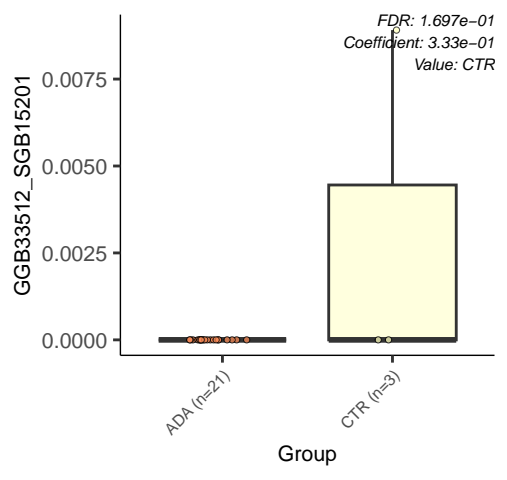
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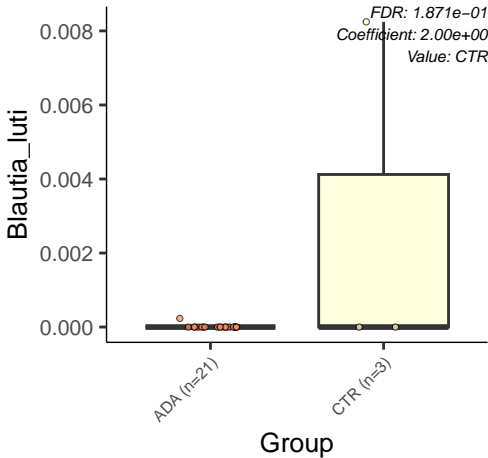
ADA (n=21)

CTR (n=3)

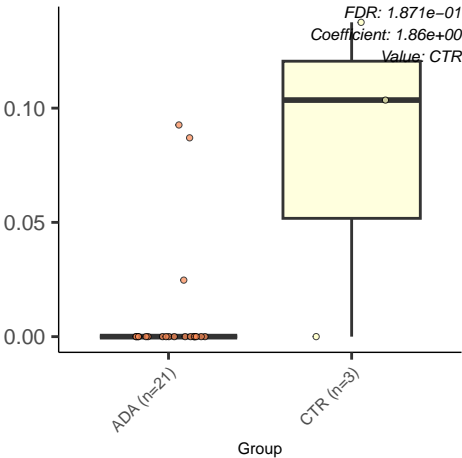
Group

FDR: 1.697e-01
Coefficient: 3.33e-01
Value: CTR

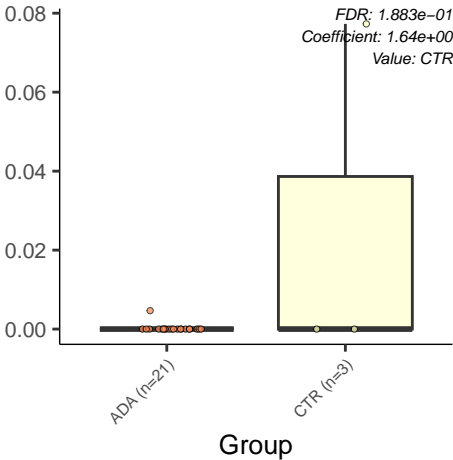




Bifidobacterium_pseudocatenulatum



GGB3614_SGB4886



Butyricimonas_virosa

0.020

0.015

0.010

0.005

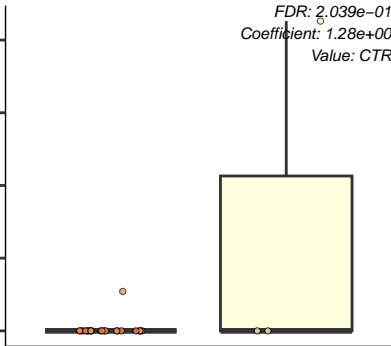
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ADA (n=21)

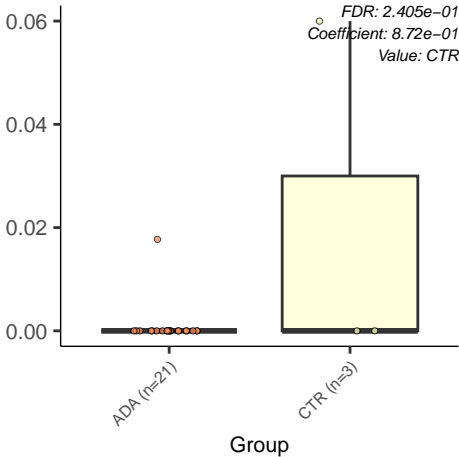
CTR (n=3)

Group

FDR: 2.039e-01
Coefficient: 1.28e+00
Value: CTR



GGB9345_SGB14311



Roseburia_intestinalis

FDR: 2.405e-01
Coefficient: 4.04e+00
Value: CTR

ADA (n=21)

CTR (n=3)

Group

0.3

0.2

0.1

0.0

Gemmiger_SGB15295

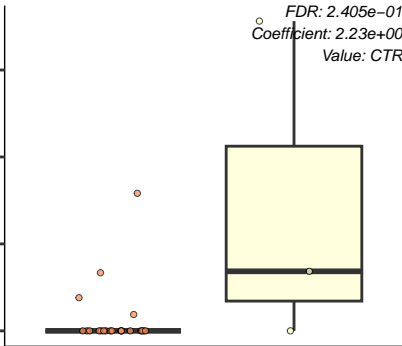
0.06
0.04
0.02
0.00

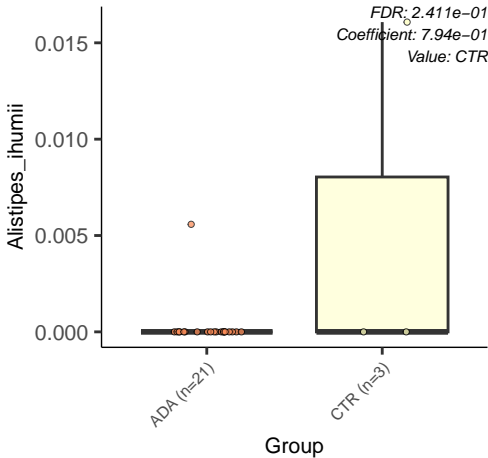
ADA (n=21)

CTR (n=3)

Group

FDR: 2.405e-01
Coefficient: 2.23e+00
Value: CTR





Mesosutterella_multiformis

0.06

0.04

0.02

0.00

ADA (n=21)

CTR (n=3)

Group

FDR: 2.411e-01
Coefficient: 2.01e+00
Value: CTR

