

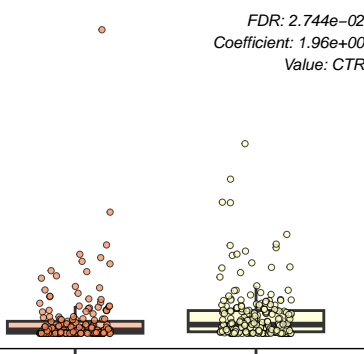
Phocaeicola\_vulgatus

*FDR: 2.744e-02*  
*Coefficient: 1.96e+00*  
*Value: CTR*

ADA (n=187)

CTR (n=214)

Group



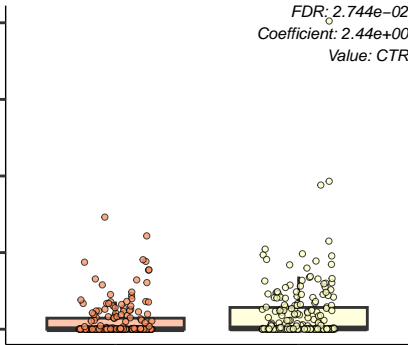
Phascolarctobacterium\_faecium

FDR:  $2.744e-02$   
Coefficient:  $2.44e+00$   
Value: CTR

ADA (n=187)

CTR (n=214)

Group



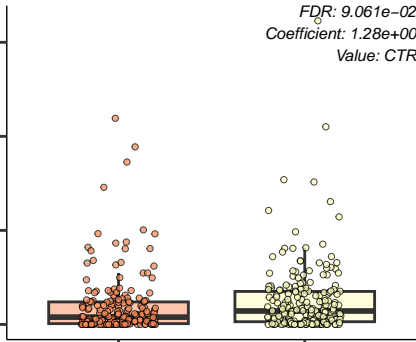
Clostridiaceae\_bacterium

*FDR: 9.061e-02*  
*Coefficient: 1.28e+00*  
*Value: CTR*

ADA (n=187)

CTR (n=214)

Group



Roseburia\_intestinalis

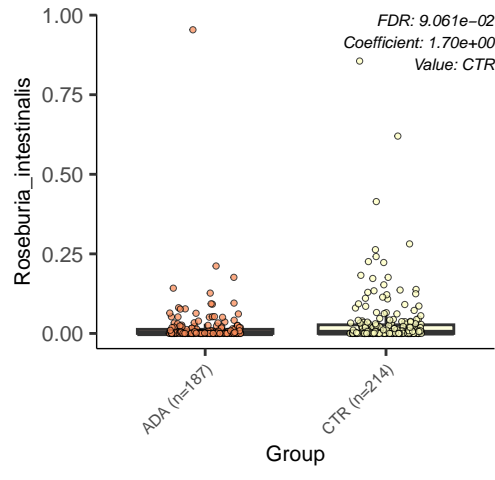
1.00  
0.75  
0.50  
0.25  
0.00

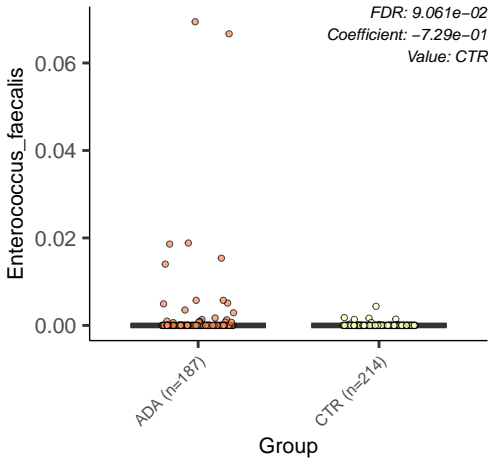
*FDR: 9.061e-02*  
*Coefficient: 1.70e+00*  
*Value: CTR*

ADA (n=187)

CTR (n=214)

Group





GGB9642\_SGB15119

*FDR: 1.488e-01*

*Coefficient: -6.86e-01*

*Value: CTR*

ADA (n=187)

CTR (n=214)

Group

0.00  
0.01  
0.02  
0.03  
0.04

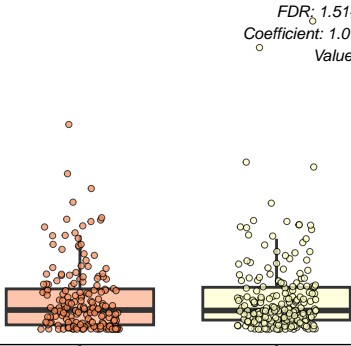
Faecalibacterium\_prausnitzii

FDR: 1.514e-01  
Coefficient: 1.07e+00  
Value: CTR

ADA (n=187)

CTR (n=214)

Group



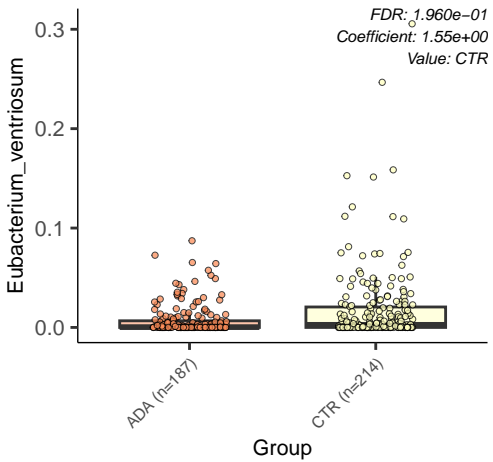
Eubacterium\_ventricosum

FDR: 1.960e-01  
Coefficient: 1.55e+00  
Value: CTR

ADA (n=187)

CTR (n=214)

Group





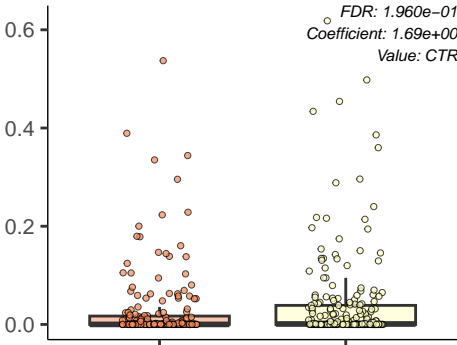
Ruminococcus\_bicirculans

FDR: 1.960e-01  
Coefficient: 1.69e+00  
Value: CTR

ADA (n=187)

CTR (n=214)

Group



Faecalicatena\_fissicatena

FDR: 1.960e-01  
Coefficient: 1.02e+00  
Value: CTR

ADA (n=187)

CTR (n=214)

Group



Streptococcus\_thermophilus

FDR: 1.960e-01  
Coefficient: 9.94e-01  
Value: CTR

0.00

0.05

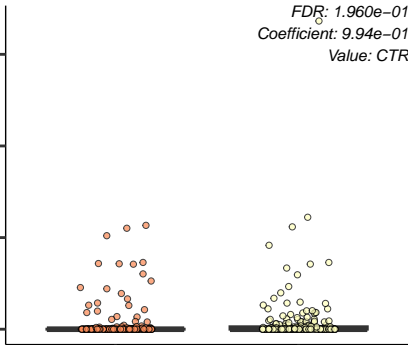
0.10

0.15

ADA (n=187)

CTR (n=214)

Group



Parasutterella\_excrementihominis

FDR: 1.960e-01  
Coefficient: 1.64e+00  
Value: CTR

0.4

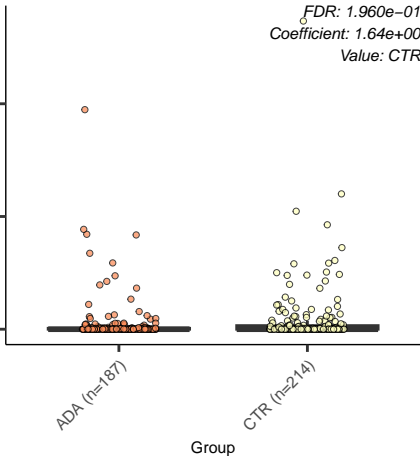
0.2

0.0

ADA (n=187)

CTR (n=214)

Group



Clostridium\_sp\_AM33\_3

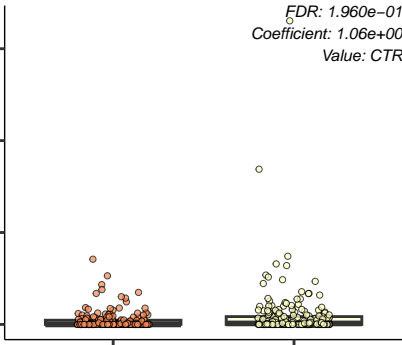
FDR: 1.960e-01  
Coefficient: 1.06e+00  
Value: CTR

0.15  
0.10  
0.05  
0.00

ADA (n=187)

CTR (n=214)

Group



Clostridium\_SGB4750

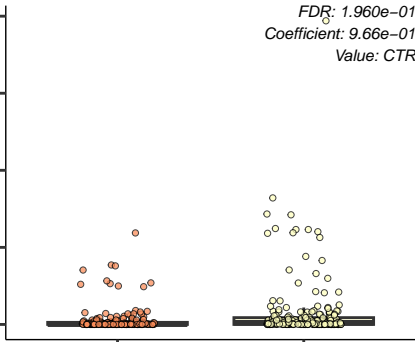
0.4  
0.3  
0.2  
0.1  
0.0

*FDR: 1.960e-01*  
*Coefficient: 9.66e-01*  
*Value: CTR*

ADA (n=187)

CTR (n=214)

Group



GGB9614\_SGB15049

*FDR: 1.960e-01*  
*Coefficient: 9.88e-01*  
*Value: CTR*

0.10

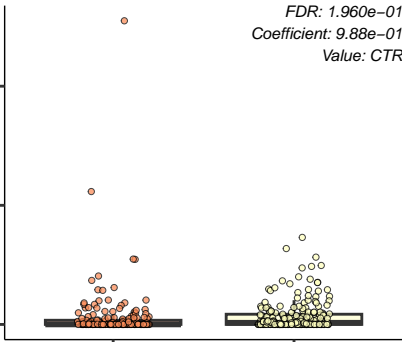
0.05

0.00

ADA (n=187)

CTR (n=214)

Group



GGB3109\_SGB4119

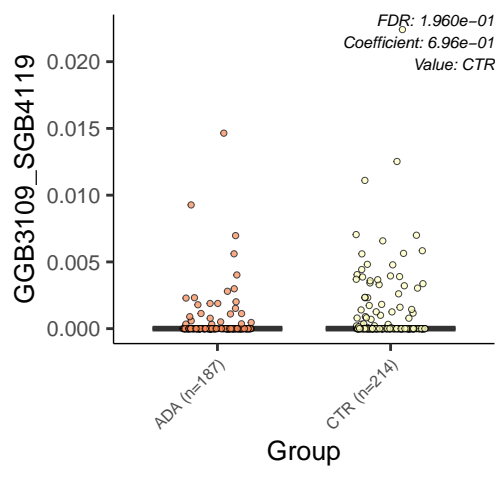
*FDR: 1.960e-01*  
*Coefficient: 6.96e-01*  
*Value: CTR*

0.020  
0.015  
0.010  
0.005  
0.000

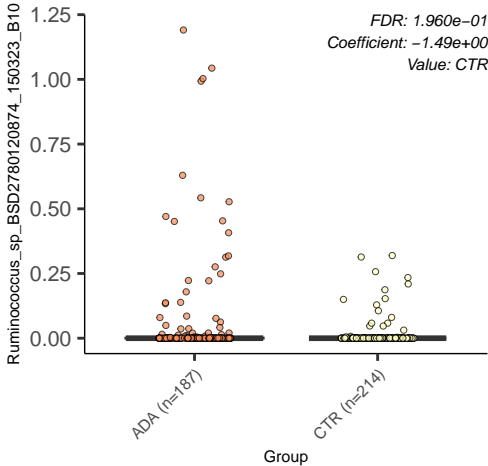
ADA (n=187)

CTR (n=214)

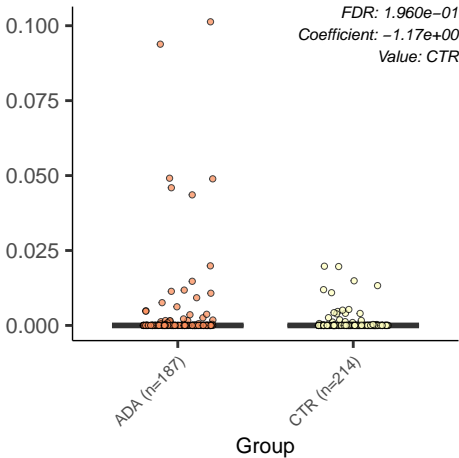
Group

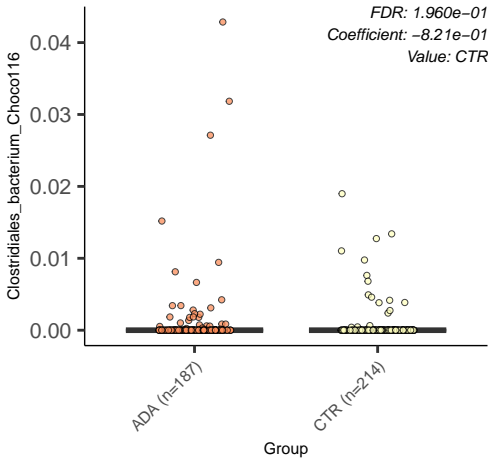






GGB42994\_SGB4746





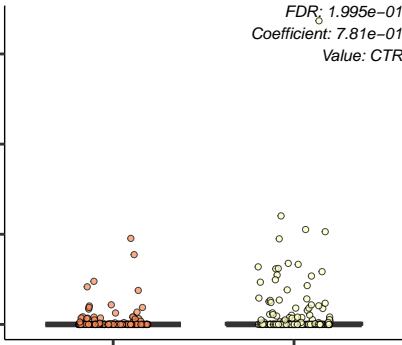
Clostridium\_sp\_AM42\_4

FDR: 1.995e-01  
Coefficient: 7.81e-01  
Value: CTR

ADA (n=187)

CTR (n=214)

Group



Klebsiella\_pneumoniae

FDR: 2.089e-01  
Coefficient: -9.38e-01  
Value: CTR

ADA (n=187)

CTR (n=214)

Group

