

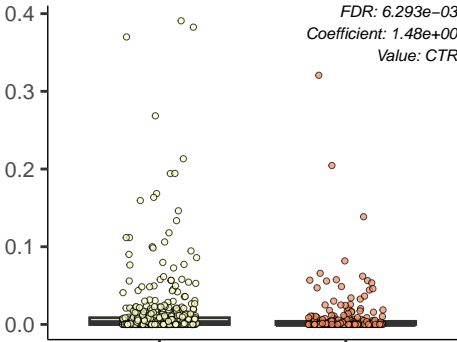
Clostridium_SGB4750

FDR: 6.293e-03
Coefficient: 1.48e+00
Value: CTR

CTR (n=512)

ADA (n=276)

Group



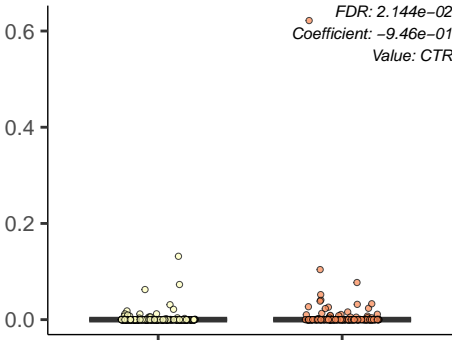
GGB3614_SGB4886

FDR: 2.144×10^{-2}
Coefficient: -9.46×10^{-1}
Value: CTR

CTR (n=512)

ADA (n=276)

Group



Streptococcus_thermophilus

FDR: 2.674e-02
Coefficient: 1.29e+00
Value: CTR

1.0

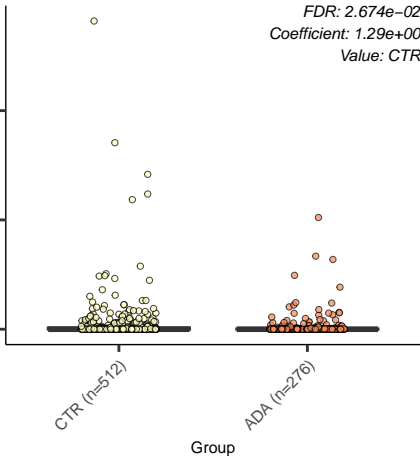
0.5

0.0

CTR (n=512)

ADA (n=276)

Group



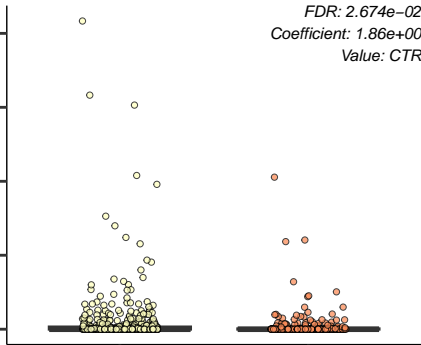
Bifidobacterium_pseudocatenulatum

FDR: 2.674e-02
Coefficient: 1.86e+00
Value: CTR

CTR (n=512)

ADA (n=276)

Group



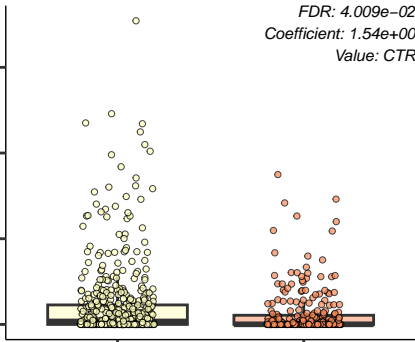
Eubacterium_ventricosum

FDR: 4.009e-02
Coefficient: 1.54e+00
Value: CTR

CTR (n=512)

ADA (n=276)

Group



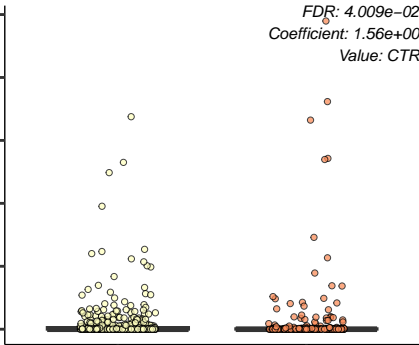
Erysipelatoclostridium_ramosum

FDR: 4.009e-02
Coefficient: 1.56e+00
Value: CTR

CTR (n=512)

ADA (n=276)

Group



Enterococcus_faecalis

0.2

0.1

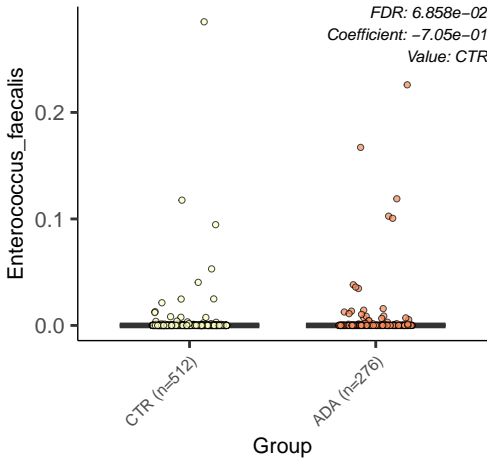
0.0

CTR (n=512)

ADA (n=276)

Group

FDR: $6.858e-02$
Coefficient: $-7.05e-01$
Value: CTR



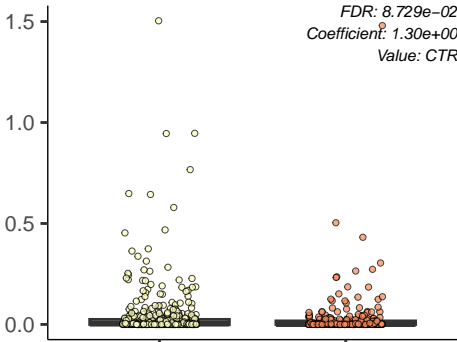
Bacteroides_xylanisolvens

FDR: 8.729×10^{-2}
Coefficient: 1.30×10^0
Value: CTR

CTR (n=512)

ADA (n=276)

Group



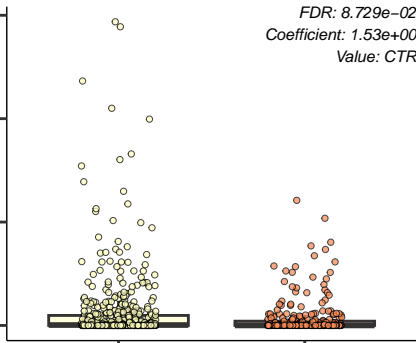
Ruminococcus_bircirculans

FDR: 8.729e-02
Coefficient: 1.53e+00
Value: CTR

CTR (n=512)

ADA (n=276)

Group



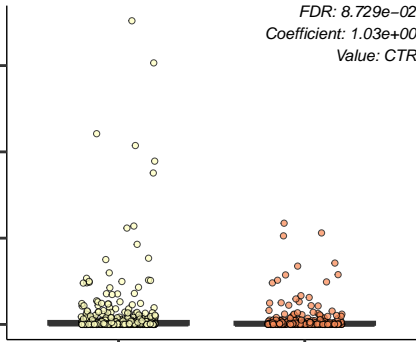
Streptococcus_salivarius

FDR: 8.729e-02
Coefficient: 1.03e+00
Value: CTR

CTR (n=512)

ADA (n=276)

Group



GGB9642_SGB15119

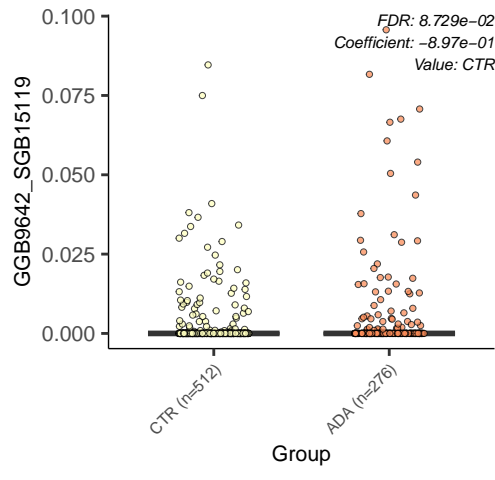
0.100
0.075
0.050
0.025
0.000

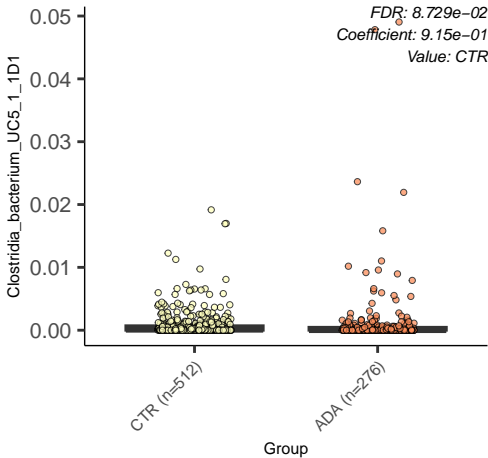
CTR (n=512)

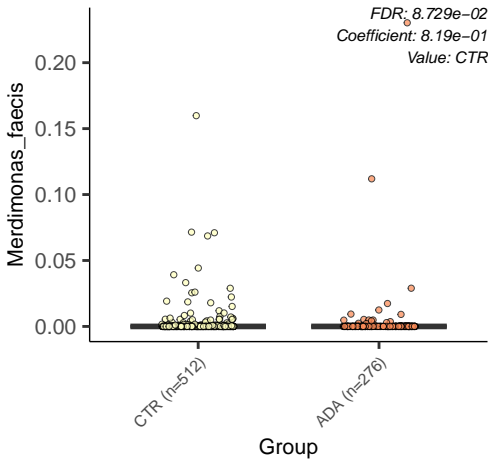
ADA (n=276)

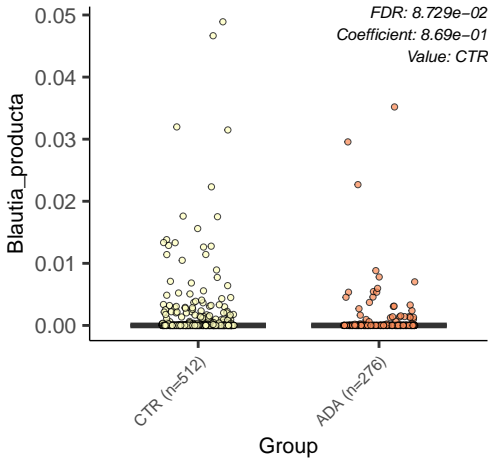
Group

FDR: $8.729e-02$
Coefficient: $-8.97e-01$
Value: CTR









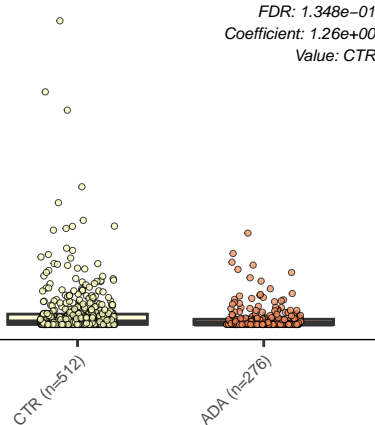
Lacrimispora_amygdalina

FDR: 1.348e-01
Coefficient: 1.26e+00
Value: CTR

CTR (n=512)

ADA (n=276)

Group



Collinsella_SGB4121

0.06

0.04

0.02

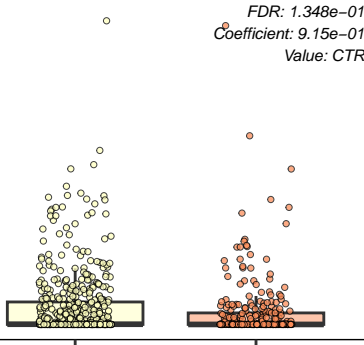
0.00

CTR (n=512)

ADA (n=276)

Group

FDR: 1.348e-01
Coefficient: 9.15e-01
Value: CTR



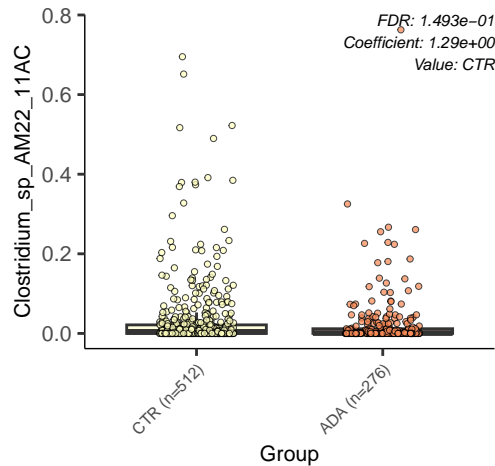
Clostridium_sp_AM22_11AC

FDR: 1.493e-01
Coefficient: 1.29e+00
Value: CTR

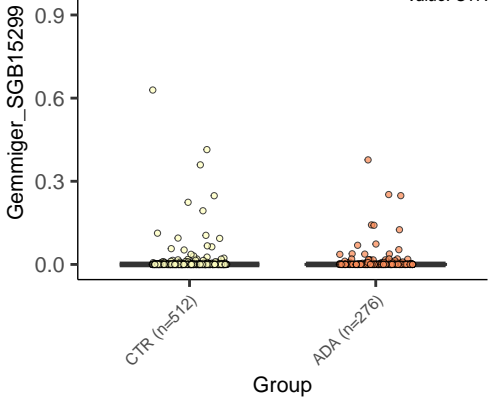
CTR (n=512)

ADA (n=276)

Group



Value: CTR



Eggerthella_lenta

0.3

0.2

0.1

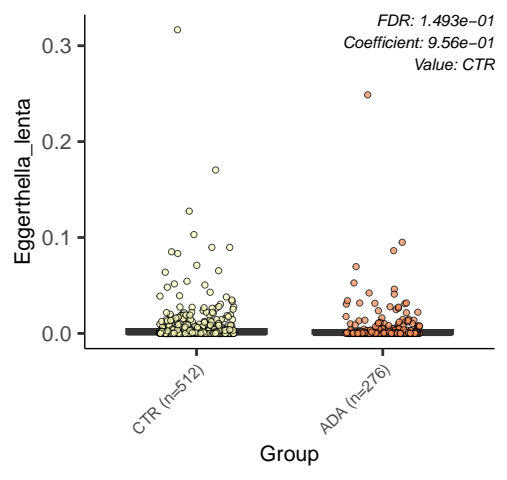
0.0

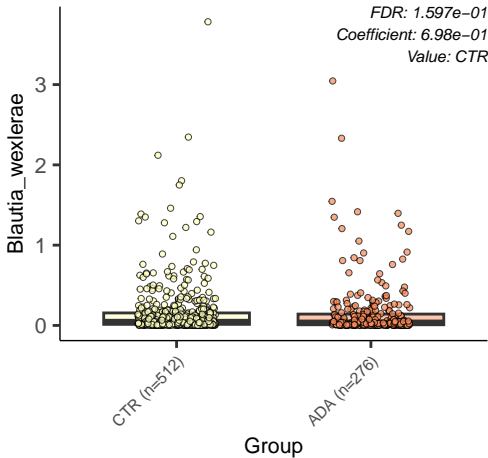
CTR (n=512)

ADA (n=276)

Group

FDR: 1.493e-01
Coefficient: 9.56e-01
Value: CTR





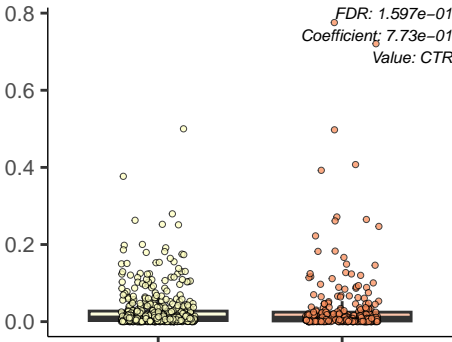
Blautia_faecis

FDR: 1.597e-01
Coefficient: 7.73e-01
Value: CTR

CTR (n=512)

ADA (n=276)

Group



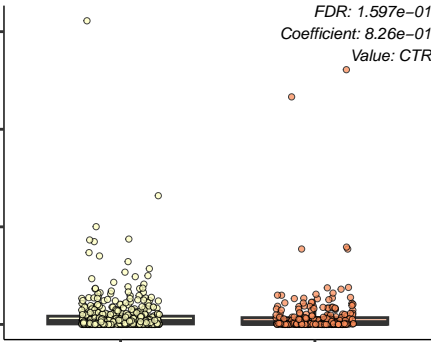
Anaerostipes_hadrus

FDR: 1.597e-01
Coefficient: 8.26e-01
Value: CTR

CTR (n=512)

ADA (n=276)

Group



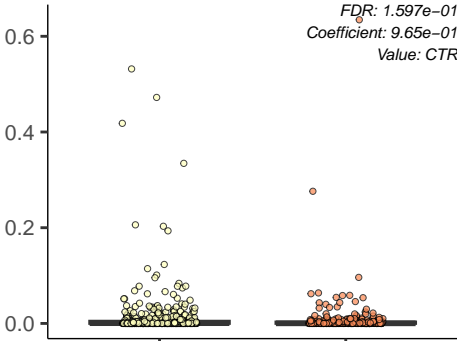
Enterocloster_bolteae

FDR: 1.597e-01
Coefficient: 9.65e-01
Value: CTR

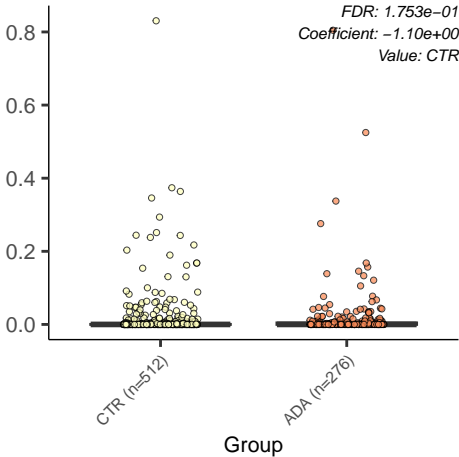
CTR (n=512)

ADA (n=276)

Group



Bacteroides_finegoldii



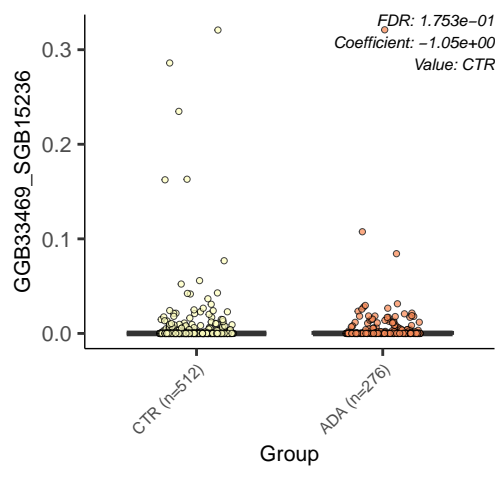
GGB33469_SGB15236

FDR: 1.753e-01
Coefficient: -1.05e+00
Value: CTR

CTR (n=512)

ADA (n=276)

Group



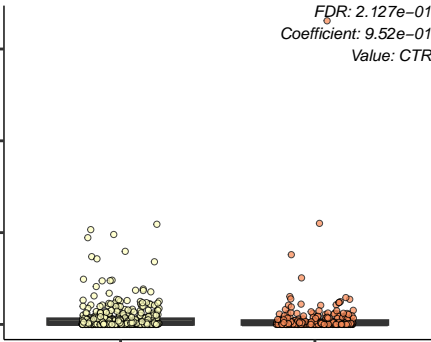
Roseburia_inulinivorans

FDR: 2.127e-01
Coefficient: 9.52e-01
Value: CTR

CTR (n=512)

ADA (n=276)

Group



GGB9635_SGB15106

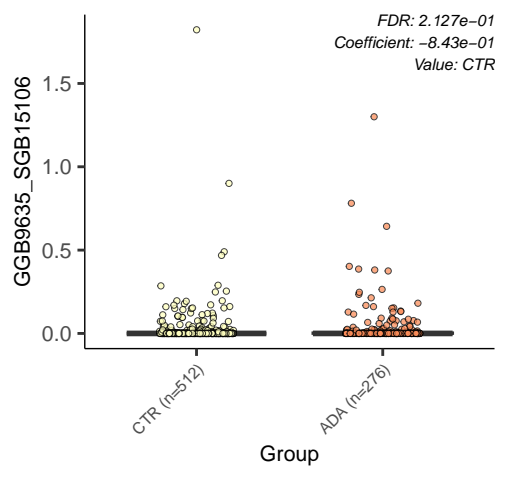
FDR: 2.127e-01
Coefficient: -8.43e-01
Value: CTR

CTR (n=512)

ADA (n=276)

Group

1.5
1.0
0.5
0.0



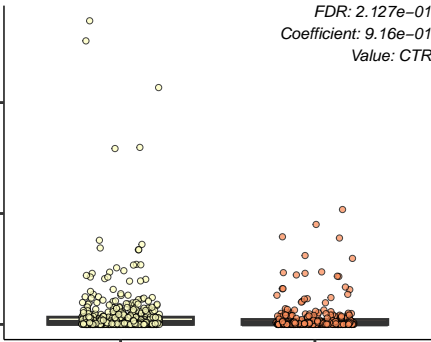
Bifidobacterium_longum

FDR: 2.127e-01
Coefficient: 9.16e-01
Value: CTR

CTR (n=512)

ADA (n=276)

Group



GGB47687_SGB2286

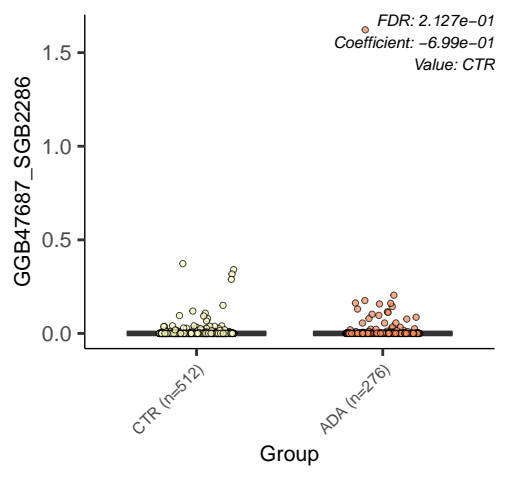
FDR: 2.127e-01
Coefficient: -6.99e-01
Value: CTR

1.5
1.0
0.5
0.0

CTR (n=512)

ADA (n=276)

Group



Ruminococcus_gnavus

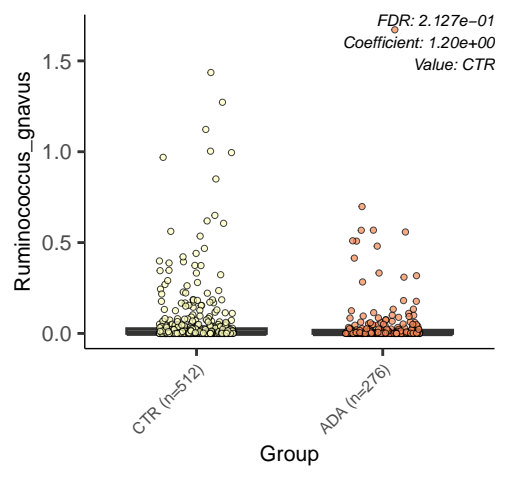
FDR: 2.127e-01
Coefficient: 1.20e+00
Value: CTR

CTR (n=512)

ADA (n=276)

Group

1.5
1.0
0.5
0.0



Butyricimonas_SGB15260

FDR: $2.127e-01$
Coefficient: $-7.82e-01$
Value: CTR

0.6

0.4

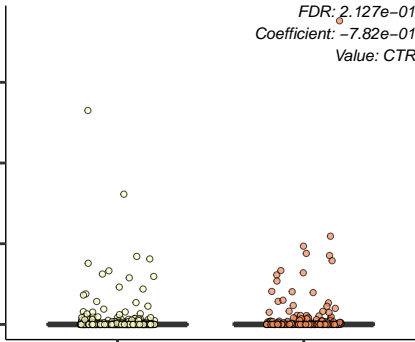
0.2

0.0

CTR (n=512)

ADA (n=276)

Group



Eubacterium_SGB6276

FDR: 2.159×10^{-1}
Coefficient: -7.55×10^{-1}
Value: CTR

0.3

0.2

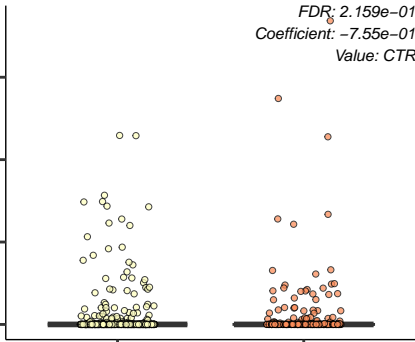
0.1

0.0

CTR (n=512)

ADA (n=276)

Group



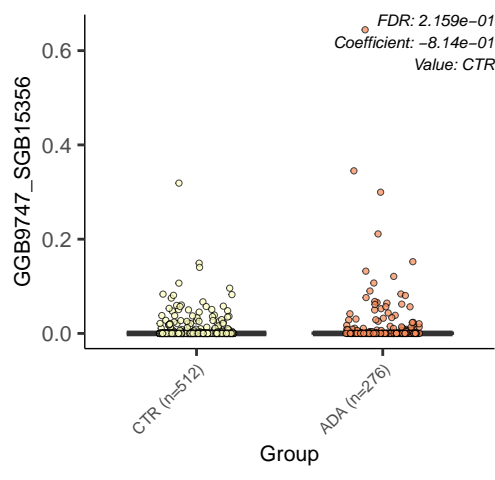
GGB9747_SGB15356

FDR: 2.159e-01
Coefficient: -8.14e-01
Value: CTR

CTR (n=512)

ADA (n=276)

Group



Candidatus_Cibionibacter_quicibialis

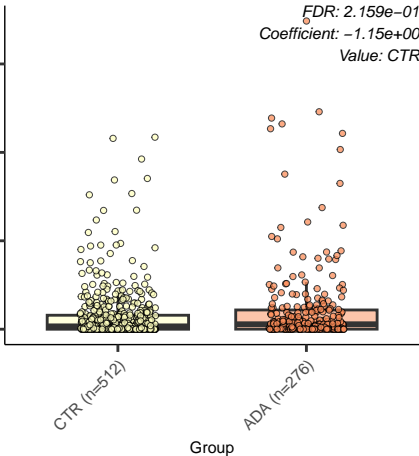
FDR: 2.159×10^{-1}
Coefficient: -1.15×10^0
Value: CTR

1.5
1.0
0.5
0.0

CTR (n=512)

ADA (n=276)

Group



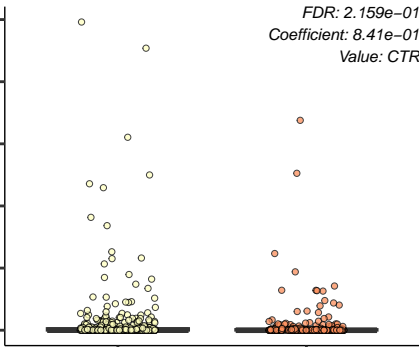
Haemophilus_parainfluenzae

FDR: 2.159e-01
Coefficient: 8.41e-01
Value: CTR

CTR (n=512)

ADA (n=276)

Group



Butyricoccus_sp_AM29_23AC

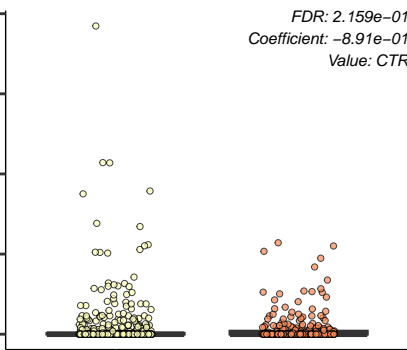
0.20
0.15
0.10
0.05
0.00

FDR: 2.159e-01
Coefficient: -8.91e-01
Value: CTR

CTR (n=512)

ADA (n=276)

Group



Desulfovibrio_piger

0.3

0.2

0.1

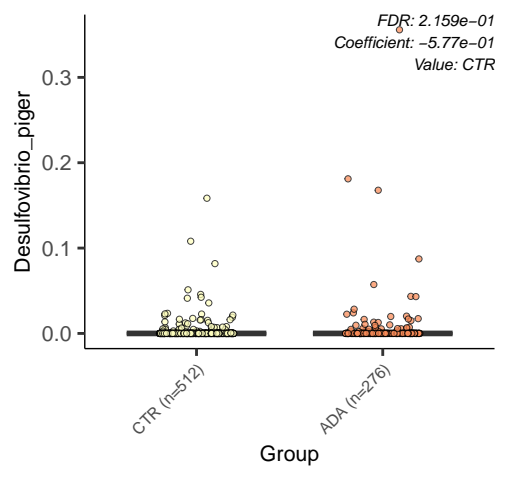
0.0

CTR (n=512)

ADA (n=276)

Group

FDR: 2.159e-01
Coefficient: -5.77e-01
Value: CTR



Ruminococcus_callidus

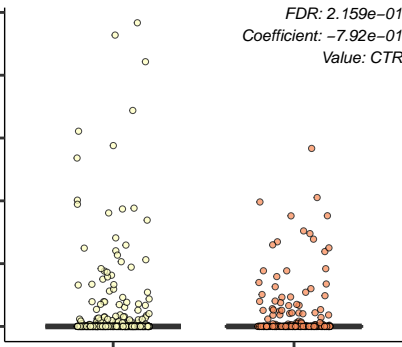
0.25
0.20
0.15
0.10
0.05
0.00

FDR: 2.159e-01
Coefficient: -7.92e-01
Value: CTR

CTR (n=512)

ADA (n=276)

Group



GGB3304_SGB4367

FDR: $2.159e-01$
Coefficient: $-8.33e-01$
Value: CTR

0.4

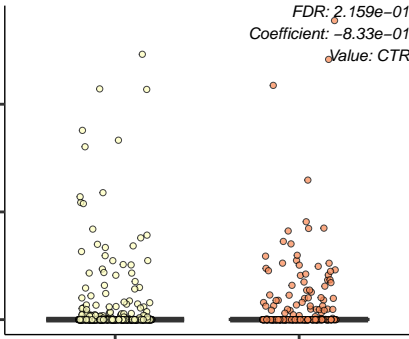
0.2

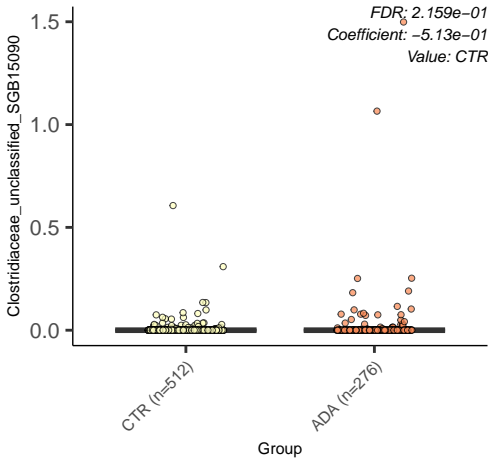
0.0

CTR (n=512)

ADA (n=276)

Group





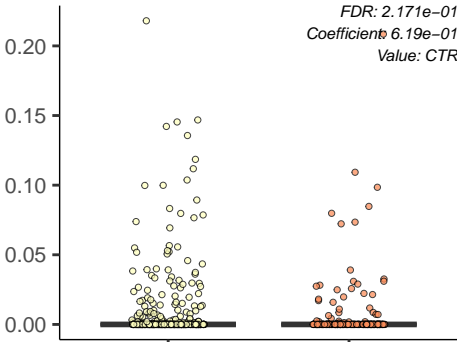
Ruminococcus_SGB4421

FDR: 2.171e-01
Coefficient: 6.19e-01
Value: CTR

CTR (n=512)

ADA (n=276)

Group



Streptococcus_parasanguinis

FDR: 2.270e-01
Coefficient: 8.24e-01
Value: CTR

0.00

0.25

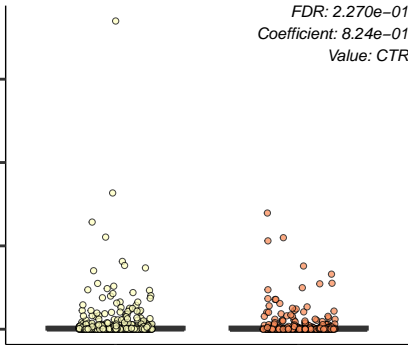
0.50

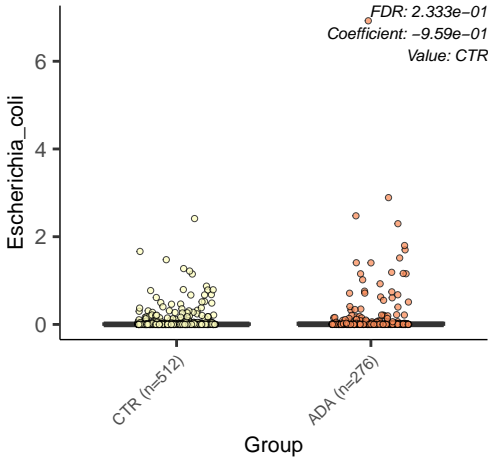
0.75

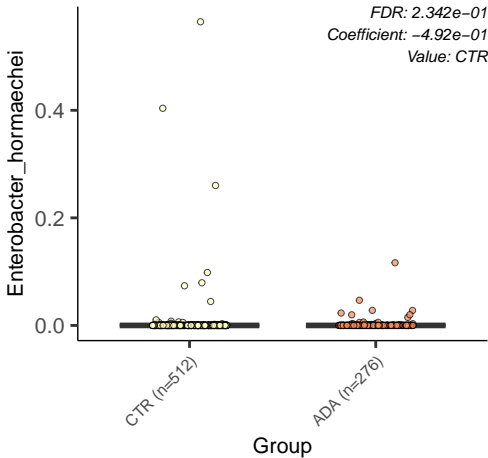
CTR (n=512)

ADA (n=276)

Group







Sellimonas_intestinalis

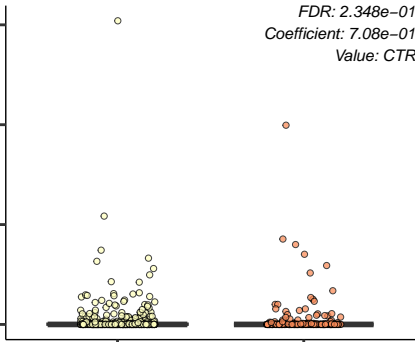
0.3
0.2
0.1
0.0

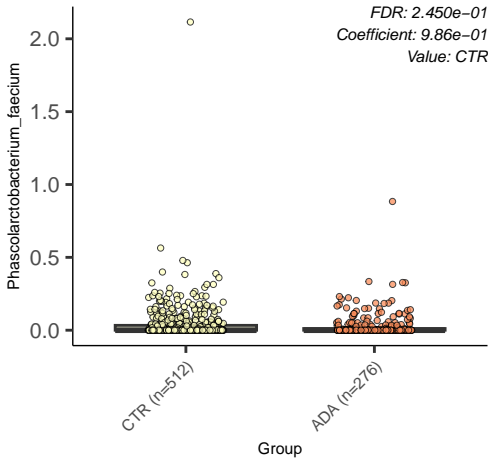
FDR: 2.348e-01
Coefficient: 7.08e-01
Value: CTR

CTR (n=512)

ADA (n=276)

Group





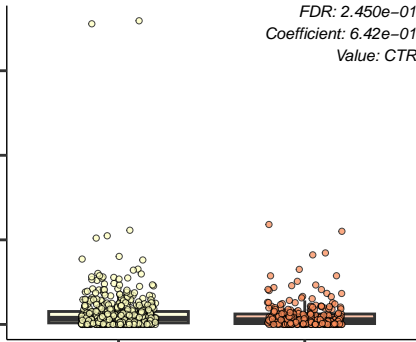
Clostridiaceae_bacterium

FDR: 2.450e-01
Coefficient: 6.42e-01
Value: CTR

CTR (n=512)

ADA (n=276)

Group



Enterocloster_citroniae

FDR: 2.450e-01
Coefficient: 7.22e-01
Value: CTR

0.03

0.02

0.01

0.00

CTR (n=512)

ADA (n=276)

Group

