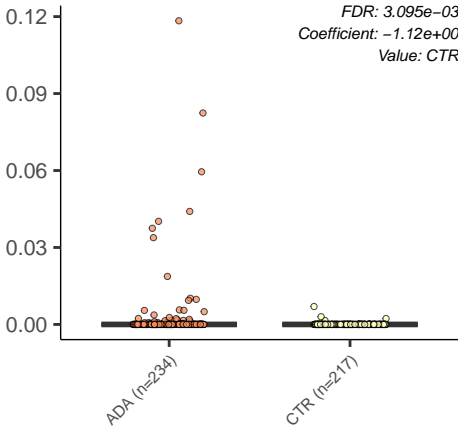




FDR: 3.095e-03

Coefficient: $-1.12e+00$

Value: CTR



Group

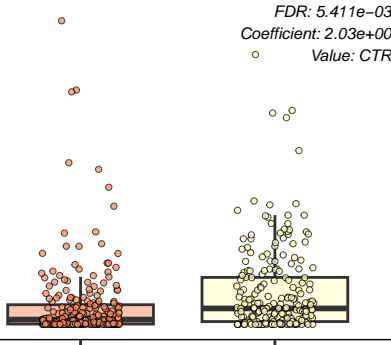
Phocaeicola_vulgatus

FDR: 5.411e-03
Coefficient: 2.03e+00
Value: CTR

ADA (n=234)

CTR (n=217)

Group



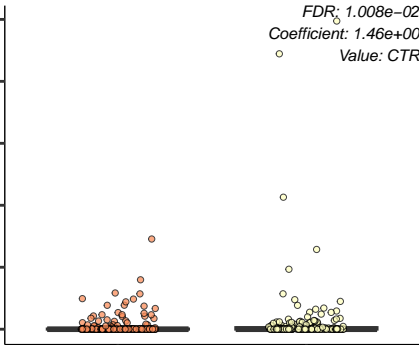
Streptococcus_thermophilus

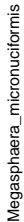
FDR: 1.008e-02
Coefficient: 1.46e+00
Value: CTR

ADA (n=234)

CTR (n=217)

Group





FDR: 1.008e-02

Coefficient: $-1.19\text{e}+00$

Value: CTR



Group

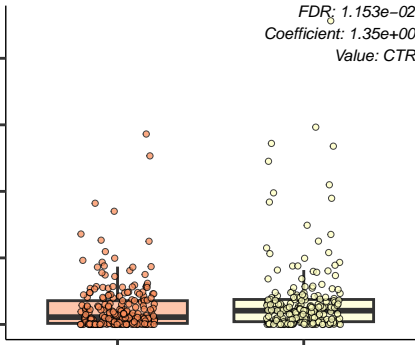
Clostridiaceae_bacterium

FDR: 1.153e-02
Coefficient: 1.35e+00
Value: CTR

ADA (n=234)

CTR (n=217)

Group



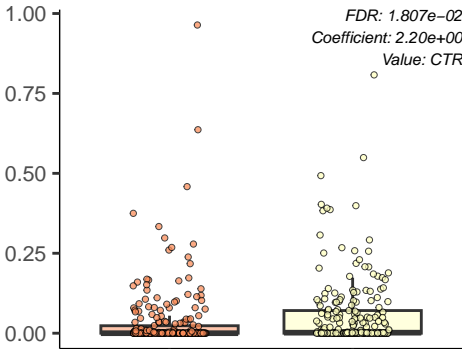
Phascolarctobacterium_faecium

FDR: 1.807e-02
Coefficient: 2.20e+00
Value: CTR

ADA (n=234)

CTR (n=217)

Group



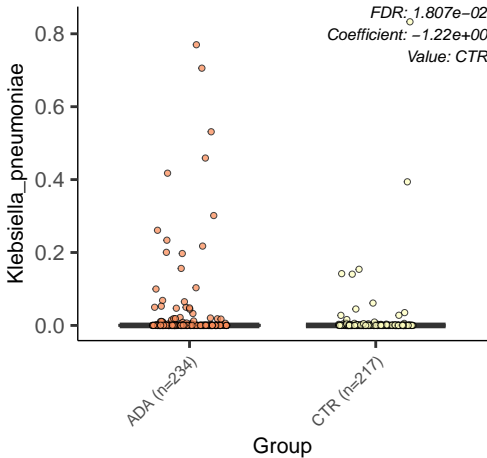
Klebsiella_pneumoniae

FDR: $1.807e-02$
Coefficient: $-1.22e+00$
Value: CTR

ADA (n=234)

CTR (n=217)

Group



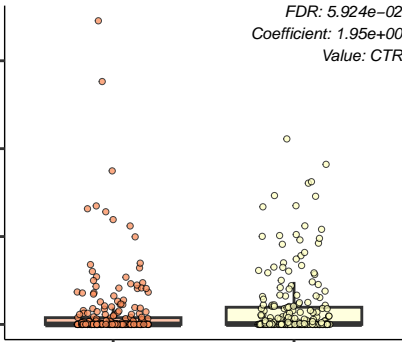
Ruminococcus_bicirculans

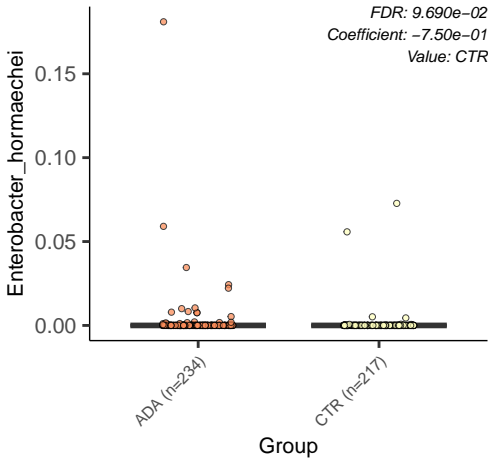
FDR: 5.924e-02
Coefficient: 1.95e+00
Value: CTR

ADA (n=234)

CTR (n=217)

Group





Parasutterella_excrementihominis

FDR: 1.142e-01
Coefficient: 1.55e+00
Value: CTR

0.9

0.6

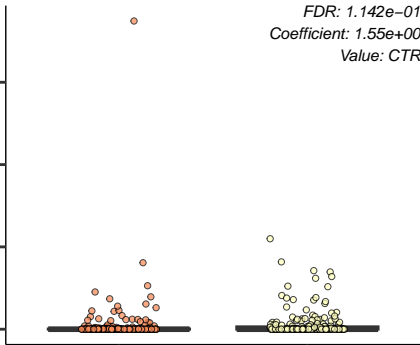
0.3

0.0

ADA (n=234)

CTR (n=217)

Group



Streptococcus_mitis

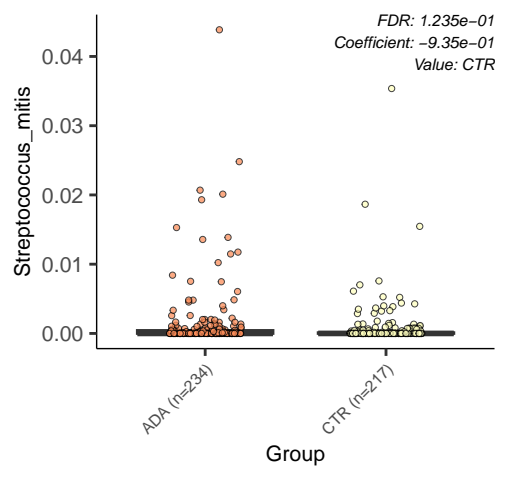
0.04
0.03
0.02
0.01
0.00

ADA (n=234)

CTR (n=217)

Group

FDR: 1.235e-01
Coefficient: -9.35e-01
Value: CTR



Blautia_faecis

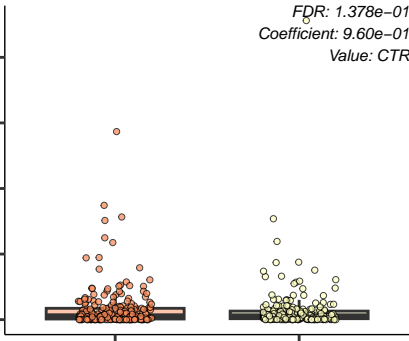
FDR: 1.378e-01
Coefficient: 9.60e-01
Value: CTR

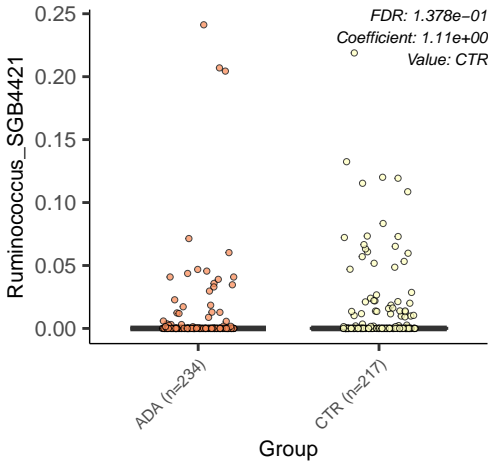
ADA (n=234)

CTR (n=217)

Group

0.4
0.3
0.2
0.1
0.0





Holdemanella_porci

0.4

0.2

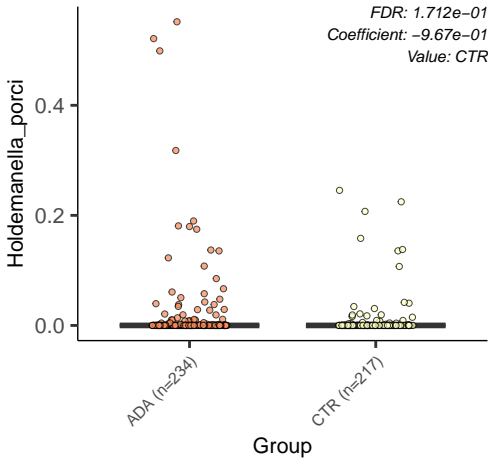
0.0

ADA (n=234)

CTR (n=217)

Group

FDR: 1.712e-01
Coefficient: -9.67e-01
Value: CTR



GGB9534_SGB14937

FDR: 1.747e-01
Coefficient: 1.17e+00
Value: CTR

0.3

0.2

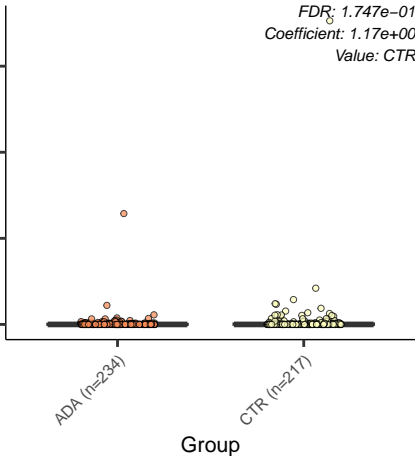
0.1

0.0

ADA (n=234)

CTR (n=217)

Group



GGB42994_SGB4746

0.06

0.04

0.02

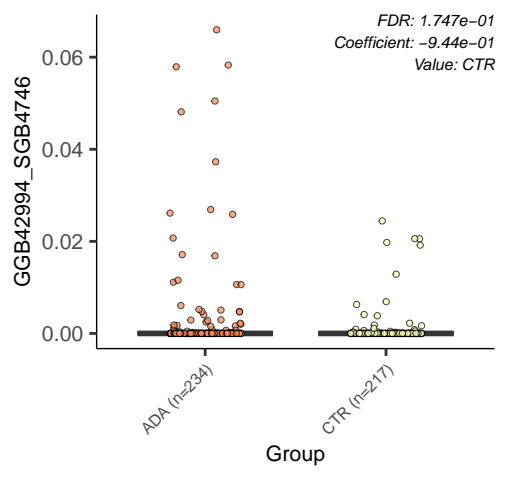
0.00

ADA (n=234)

CTR (n=217)

Group

FDR: 1.747e-01
Coefficient: -9.44e-01
Value: CTR



Clostridiales_bacterium_Choco116

FDR: 1.955e-01
Coefficient: -7.93e-01
Value: CTR

0.000

0.025

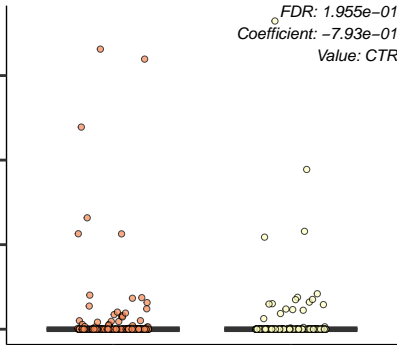
0.050

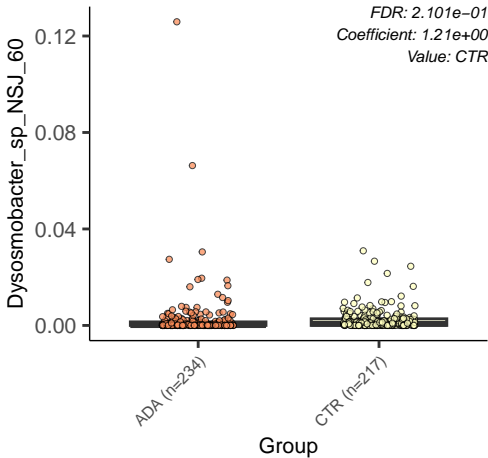
0.075

ADA (n=234)

CTR (n=217)

Group





GGB9614_SGB15049

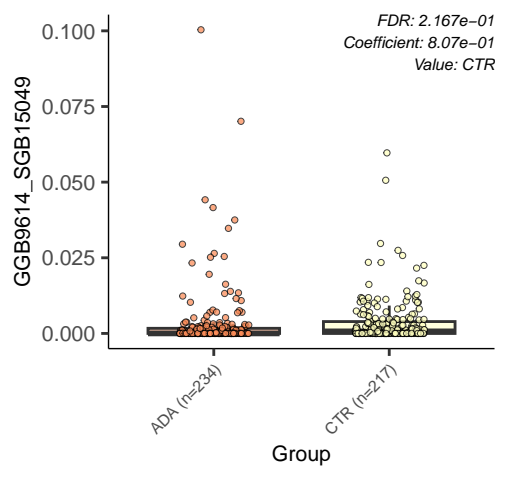
0.100
0.075
0.050
0.025
0.000

FDR: 2.167e-01
Coefficient: 8.07e-01
Value: CTR

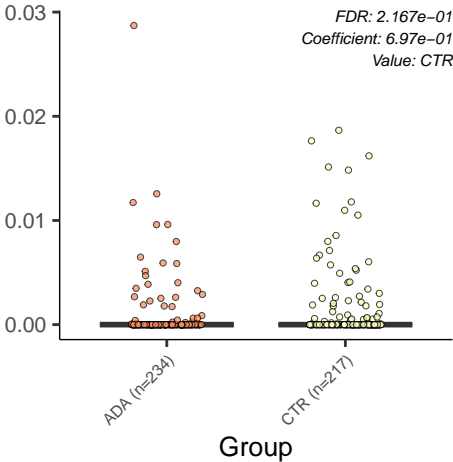
ADA (n=234)

CTR (n=217)

Group



GGB3109_SGB4119



Clostridium_sp_AM42_4

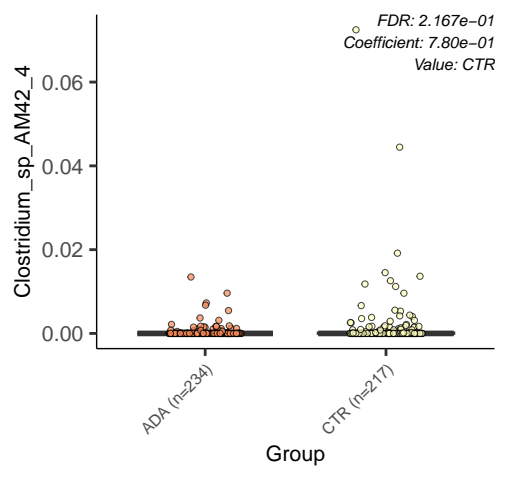
0.06
0.04
0.02
0.00

ADA (n=234)

CTR (n=217)

Group

FDR: 2.167e-01
Coefficient: 7.80e-01
Value: CTR



Veillonella_atypica

FDR: 2.283e-01

Coefficient: -1.17e+00

Value: CTR

0.20

0.15

0.10

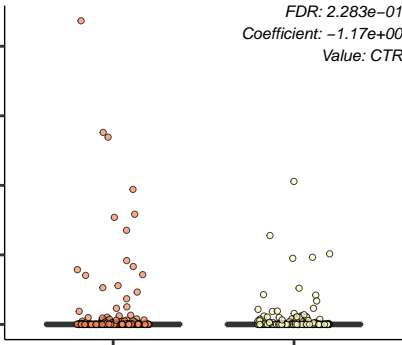
0.05

0.00

ADA (n=234)

CTR (n=217)

Group



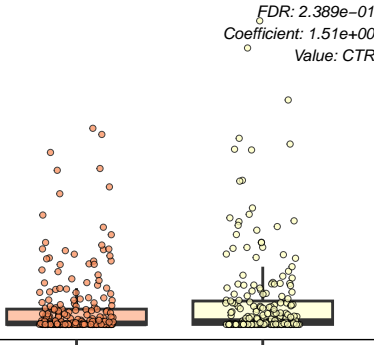
Lachnospira_eligens

FDR: 2.389e-01
Coefficient: 1.51e+00
Value: CTR

ADA (n=234)

CTR (n=217)

Group



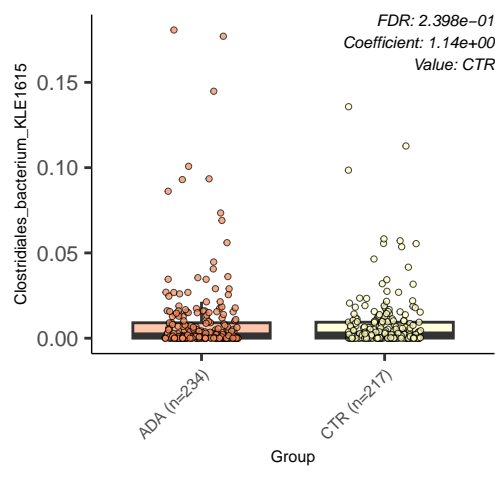
Clostridiales_bacterium_KLE1615

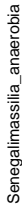
FDR: 2.398e-01
Coefficient: 1.14e+00
Value: CTR

ADA (n=234)

CTR (n=217)

Group





0.06
0.04
0.02
0.00

ADA (n=234)

CTR (n=217)

Group

FDR: 2.415e-01
Coefficient: -6.37e-01
Value: CTR

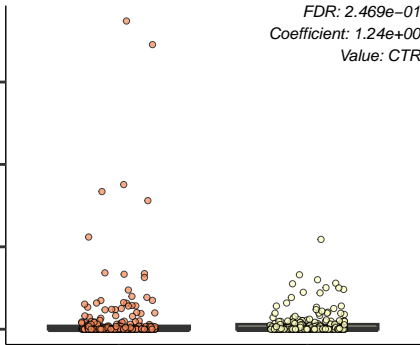
Faecalibacillus_intestinalis

FDR: 2.469e-01
Coefficient: 1.24e+00
Value: CTR

ADA (n=234)

CTR (n=217)

Group



Veillonella_tobetsuensis

