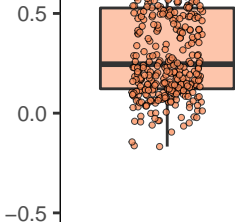


PWY.5659

FDR: 3.834e-04

Coefficient: -3.48e-02

Value: GenotypeTCR_KO.SiteCecum



0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteCecum

PWY.7228

FDR: 8.510e-04

Coefficient: -7.78e-02

Value: GenotypeTCR_KO.SiteCecum

0

-1

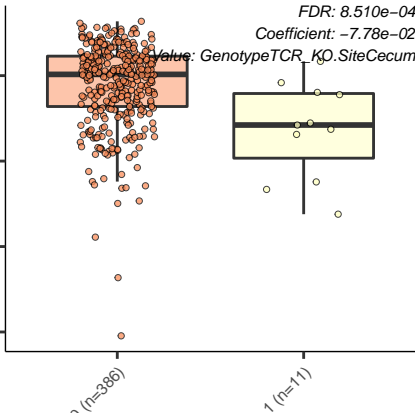
-2

-3

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteCecum



FASYN.ELONG.PWY

FDR: 1.752e-03

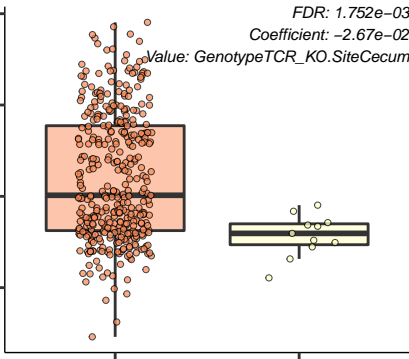
Coefficient: -2.67e-02

Value: GenotypeTCR_KO.SiteCecum

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteCecum



PWY.5913

FDR: 3.349e-03

Coefficient: -1.76e-01

Value: GenotypeTCR_KO.SiteCecum

0

-1

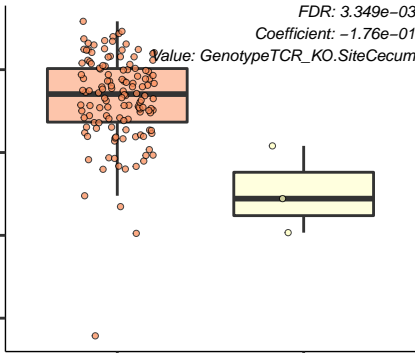
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-3

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteCecum

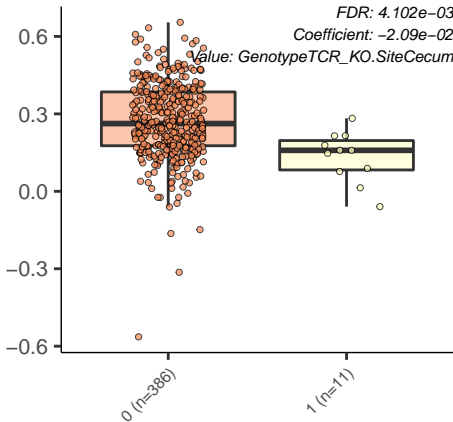


PWY.7220

FDR: 4.102e-03

Coefficient: -2.09e-02

Value: GenotypeTCR_KO.SiteCecum



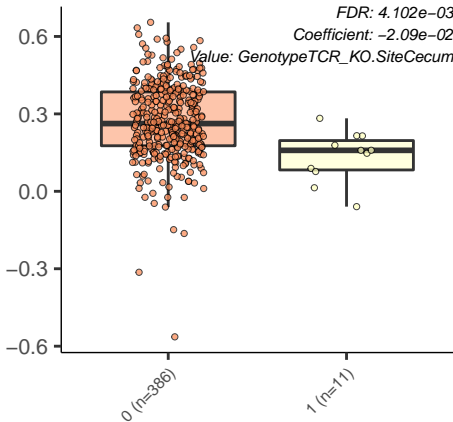
GenotypeTCR_KO.SiteCecum

PWY.7222

FDR: 4.102e-03

Coefficient: -2.09e-02

Value: GenotypeTCR_KO.SiteCecum



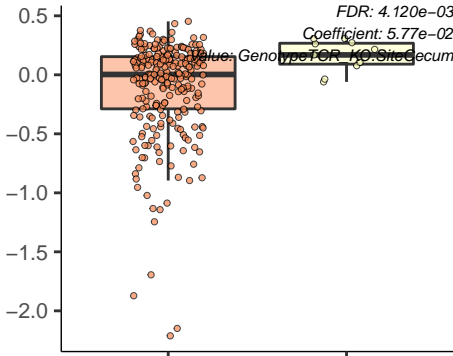
GenotypeTCR_KO.SiteCecum

COBALSYN.PWY

FDR: 4.120e-03

Coefficient: 5.77e-02

P-Value: GenotypeTCR_KO.SiteCecum



0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteCecum

PWY.6269

FDR: 6.246e-03

Coefficient: 5.63e-02

Value: GenotypeTCR_KO.SiteCecum

0

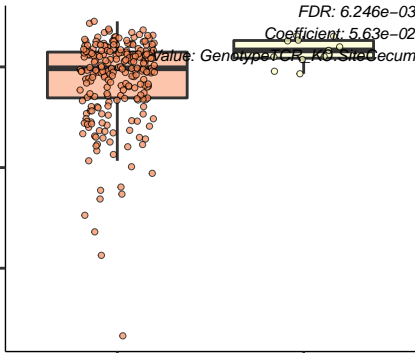
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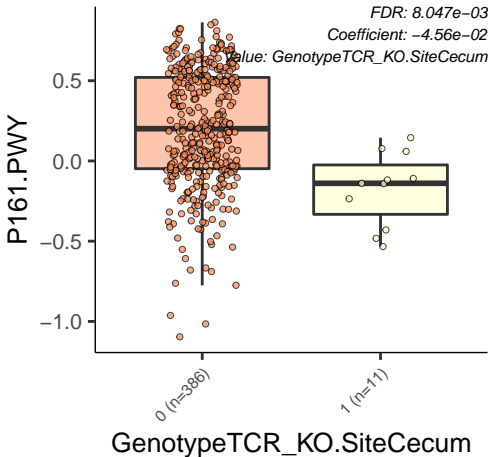
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0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteCecum





POLYISOPRENSYN.PWY

FDR: 1.250e-02

Coefficient: -2.27e-02

Value: GenotypeTCR_KO.SiteCecum

0.4

0.0

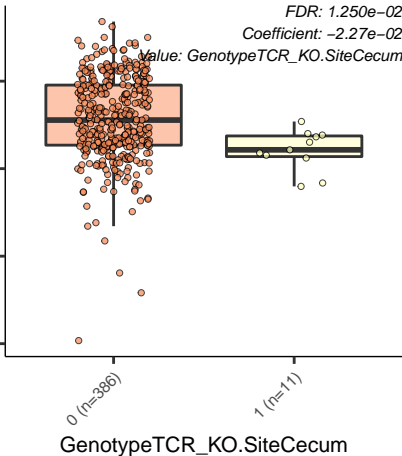
-0.4

-0.8

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteCecum



PWY.5509

FDR: 1.770e-02

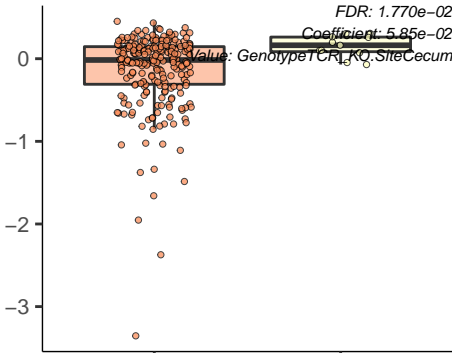
Coefficient: 5.85e-02

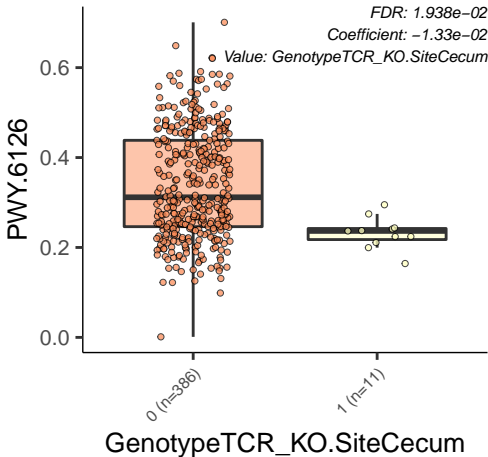
Value: GenotypeTCR_KO.SiteCecum

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteCecum



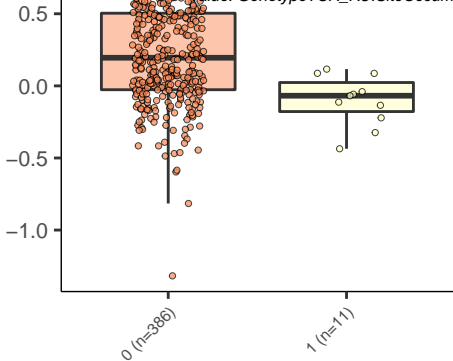


PWY0.1297

FDR: 2.007e-02

Coefficient: -3.64e-02

Value: GenotypeTCR_KO.SiteCecum



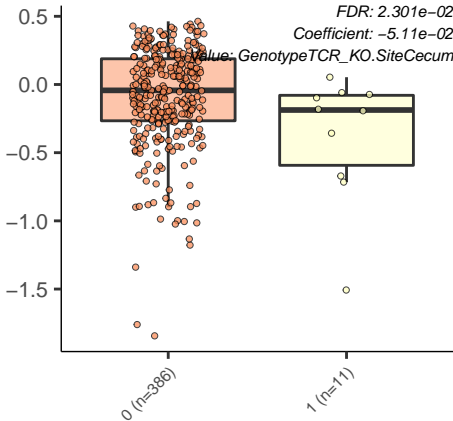
GenotypeTCR_KO.SiteCecum

PWY.6467

FDR: 2.301e-02

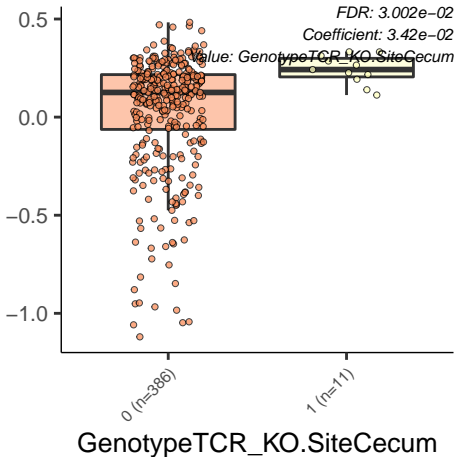
Coefficient: -5.11e-02

Value: GenotypeTCR_KO.SiteCecum

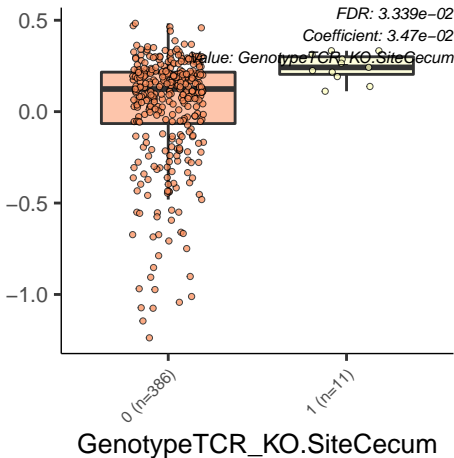


GenotypeTCR_KO.SiteCecum

PWY.7400



ARGSYN.PWY



PYRIDNUCSAL.PWY

FDR: 3.348e-02

Coefficient: 4.56e-02

Value: GenotypeTCR_KO.SiteCecum

0

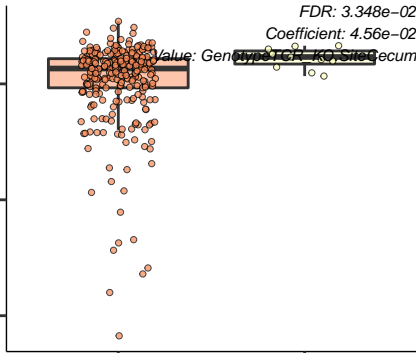
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-2

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteCecum

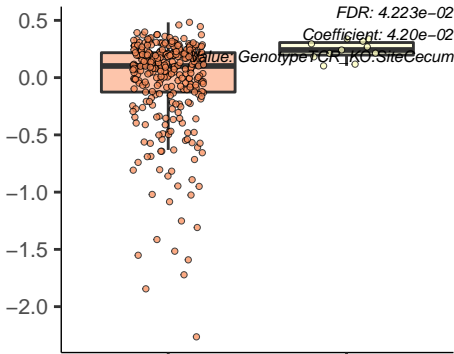


ARGSYNBSUB.PWY

FDR: 4.223e-02

Coefficient: 4.20e-02

Value: GenotypeTCR_KO.SiteCecum



0 (n=386)

1 (n=11)

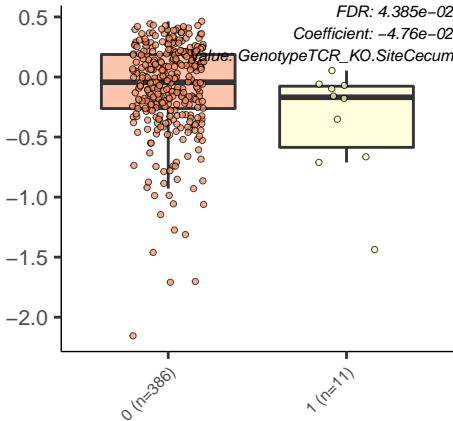
GenotypeTCR_KO.SiteCecum

PWY.1269

FDR: 4.385e-02

Coefficient: -4.76e-02

Value: GenotypeTCR_KO.SiteCecum



GenotypeTCR_KO.SiteCecum

PWY.5505

0

-1

-2

0 (n=386)

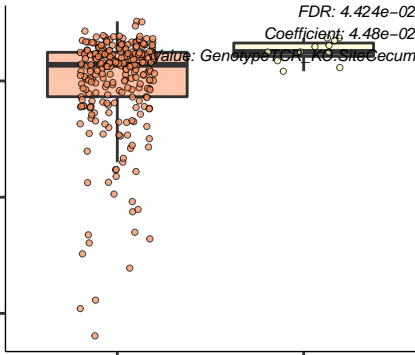
1 (n=11)

GenotypeTCR_KO.SiteCecum

FDR: 4.424e-02

Coefficient: 4.48e-02

Value: GenotypeTCR_KO.SiteCecum



PWY.7200

FDR: 5.666e-02

Coefficient: -4.89e-02

Value: GenotypeTCR_KO.SiteCecum

0

-1

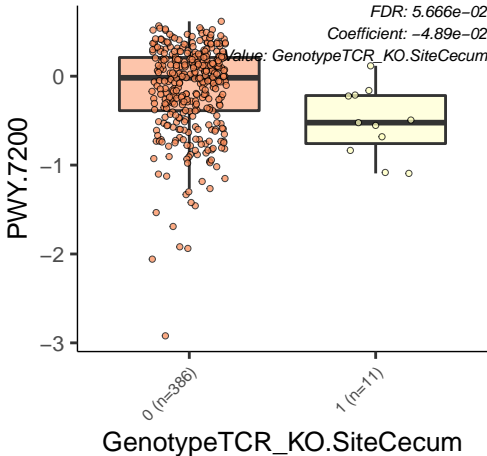
-2

-3

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteCecum

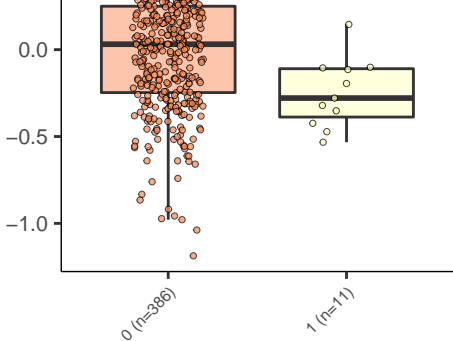


PWY.7187

FDR: 6.020e-02

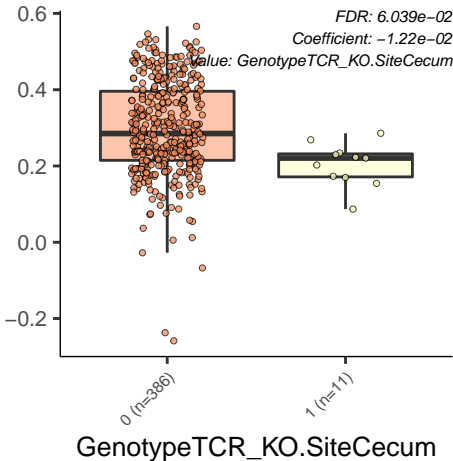
Coefficient: -3.26e-02

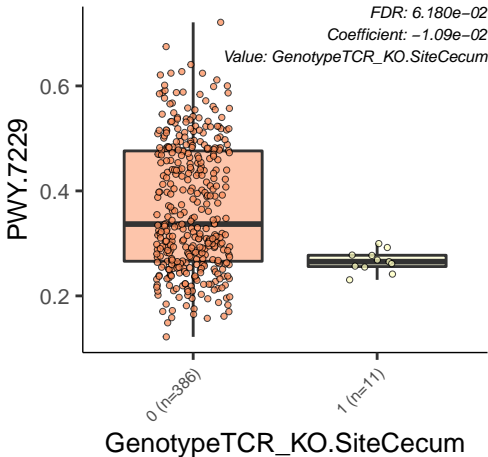
Value: GenotypeTCR_KO.SiteCecum



GenotypeTCR_KO.SiteCecum

PWY.5484





GLCMANNANAUT.PWY

FDR: 6.296e-02

Coefficient: -2.68e-02

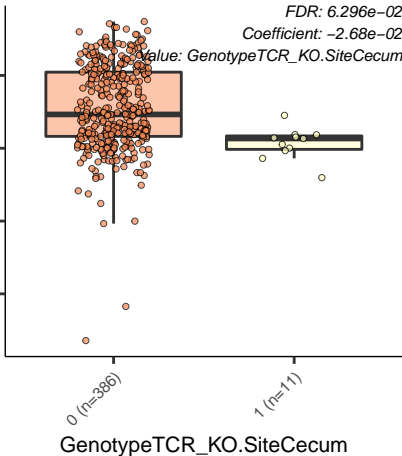
Value: GenotypeTCR_KO.SiteCecum

0.5
0.0
-0.5
-1.0

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteCecum



PWY.6125

FDR: 7.287e-02

Coefficient: -4.10e-02

Value: GenotypeTCR_KO.SiteCecum

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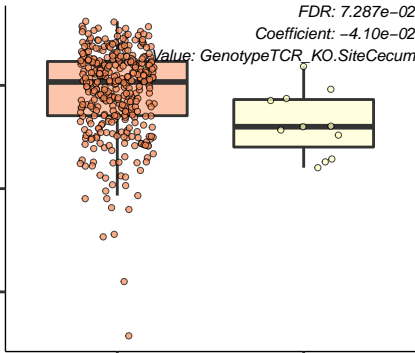
-1

-2

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteCecum

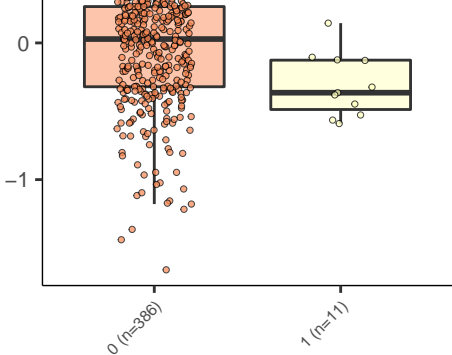


PWY0.166

FDR: 7.314e-02

Coefficient: -3.73e-02

Value: GenotypeTCR_KO.SiteCecum



GenotypeTCR_KO.SiteCecum

PWY.621

0
-1

0 (n=386)

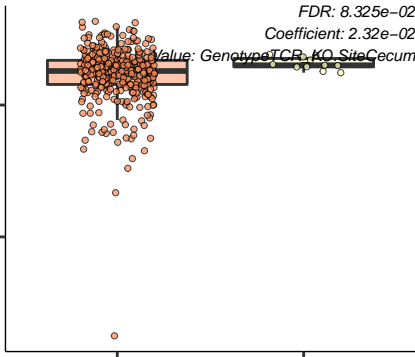
1 (n=11)

GenotypeTCR_KO.SiteCecum

FDR: 8.325e-02

Coefficient: 2.32e-02

Value: GenotypeTCR_KO.SiteCecum

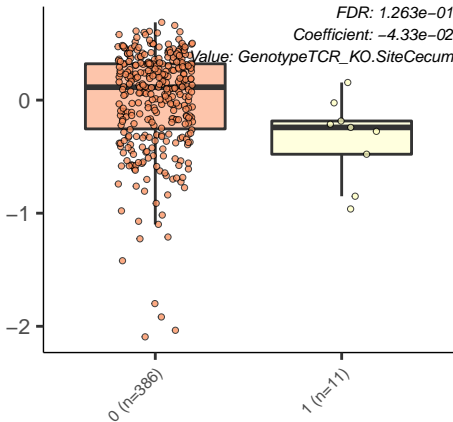


PWY.7392

FDR: 1.263e-01

Coefficient: -4.33e-02

Value: GenotypeTCR_KO.SiteCecum



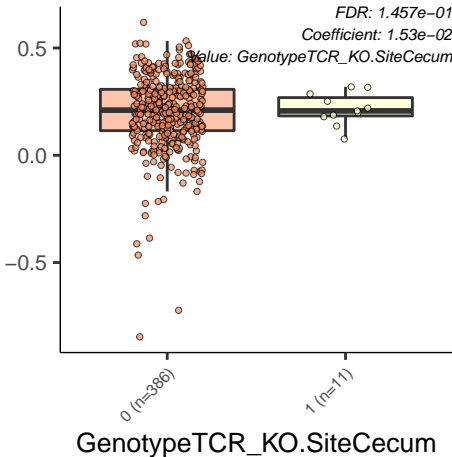
GenotypeTCR_KO.SiteCecum

OANTIGEN.PWY

FDR: 1.457e-01

Coefficient: 1.53e-02

Value: GenotypeTCR_KO.SiteCecum

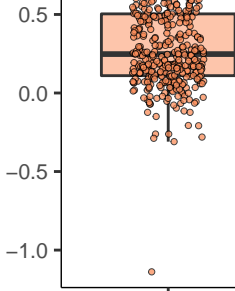


PWY0.1296

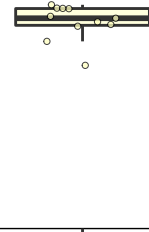
FDR: 1.558e-01

Coefficient: -1.99e-02

Value: GenotypeTCR_KO.SiteCecum



0 (n=386)



1 (n=11)

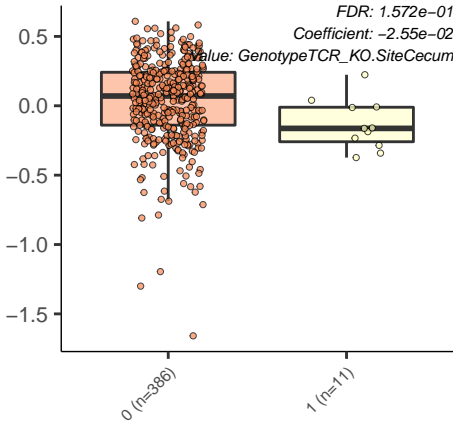
GenotypeTCR_KO.SiteCecum

PWY0.162

FDR: 1.572e-01

Coefficient: -2.55e-02

Value: GenotypeTCR_KO.SiteCecum



GenotypeTCR_KO.SiteCecum

PWY.7196

FDR: 1.622e-01

Coefficient: -3.30e-02

Value: GenotypeTCR_KO.SiteCecum

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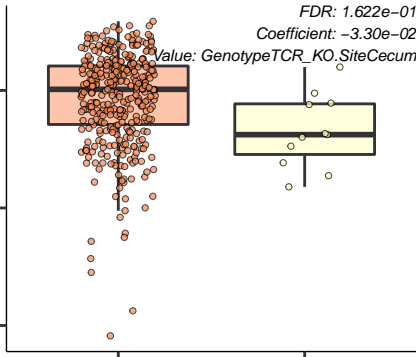
-1

-2

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteCecum

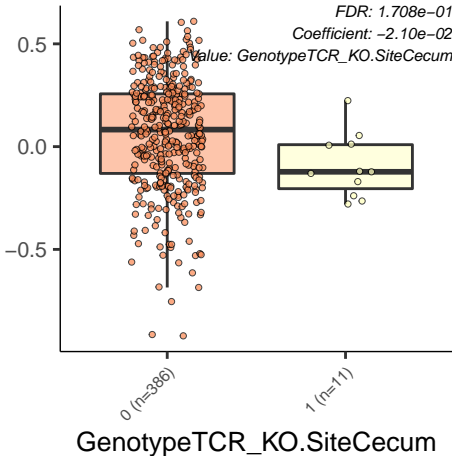


PWY.841

FDR: 1.708e-01

Coefficient: -2.10e-02

Value: GenotypeTCR_KO.SiteCecum



HOMOSER.METSYN.PWY

FDR: 1.887e-01

Coefficient: 9.39e-02

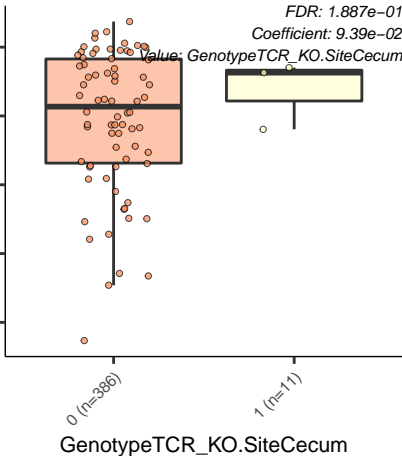
Value: GenotypeTCR_KO.SiteCecum

0.0
-0.5
-1.0
-1.5
-2.0

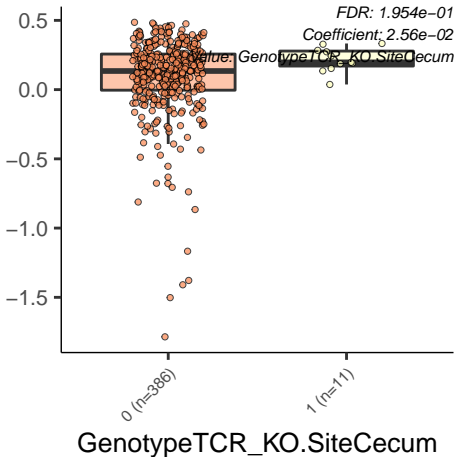
0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteCecum



PWY.6892

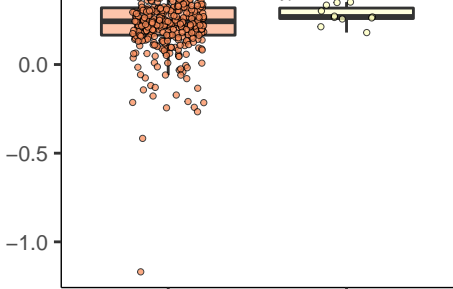


PWY.6317

FDR: 1.998e-01

Coefficient: 1.42e-02

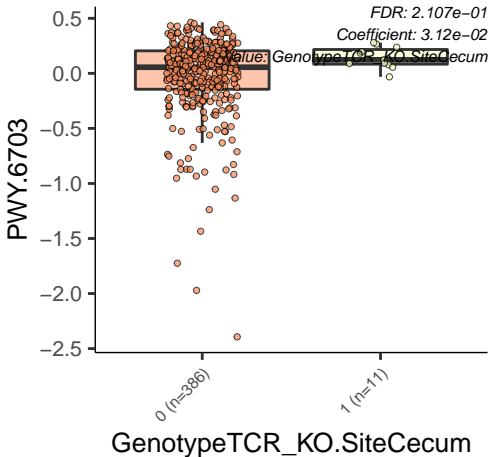
Value: GenotypeTCR_KO.SiteCecum



0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteCecum



GLYCOGENSYNTH.PWY

FDR: 2.430e-01

Coefficient: 1.51e-02

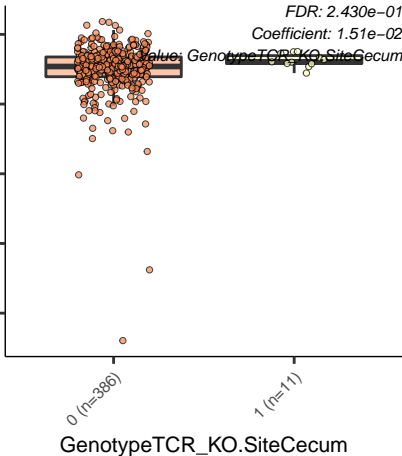
Value: GenotypeTCR_KO.SiteCecum

0.5
0.0
-0.5
-1.0
-1.5

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteCecum



GLUTORN.PWY

FDR: 2.435e-01

Coefficient: 2.87e-02

Value: GenotypeTCR_KO.SiteCecum

0

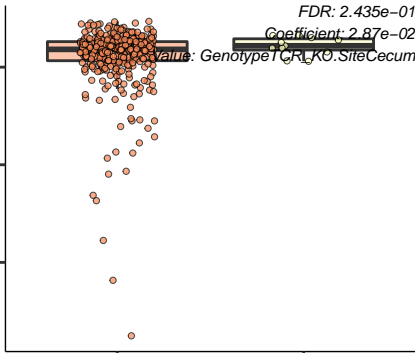
-1

-2

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteCecum

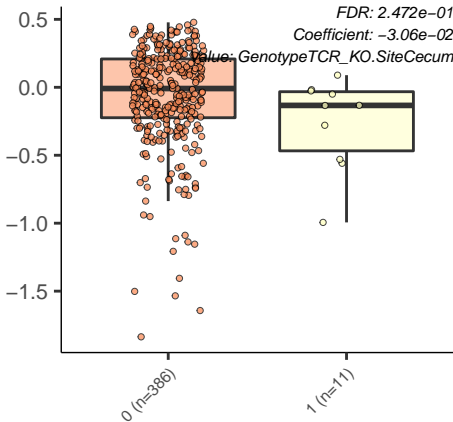


NAGLIPASYN.PWY

FDR: 2.472e-01

Coefficient: -3.06e-02

Value: GenotypeTCR_KO.SiteCecum



GenotypeTCR_KO.SiteCecum