

PWY.7197

FDR: $7.353e-04$

Coefficient: $-1.54e-01$

Value: GenotypeTCR_KO.SiteProximal_Colon

0

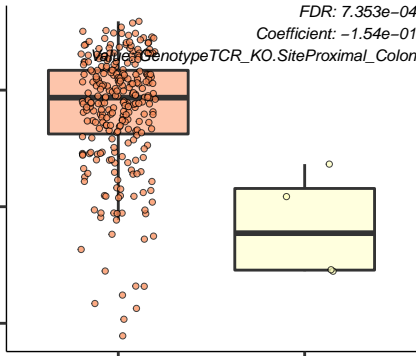
-1

-2

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteProximal_Colo



PWY.7234

FDR: 1.675e-03

Coefficient: -8.72e-02

Value: GenotypeTCR_KO.SiteProximal_Colon

0

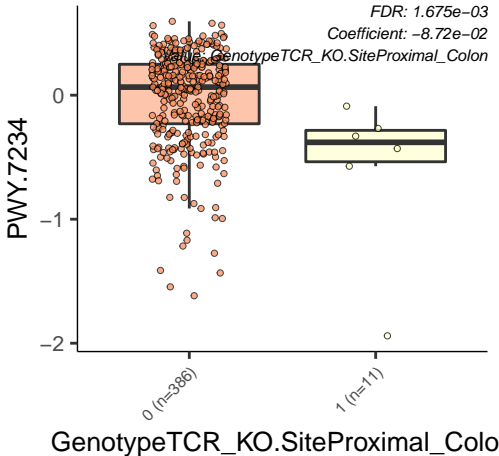
-1

-2

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteProximal_Colo

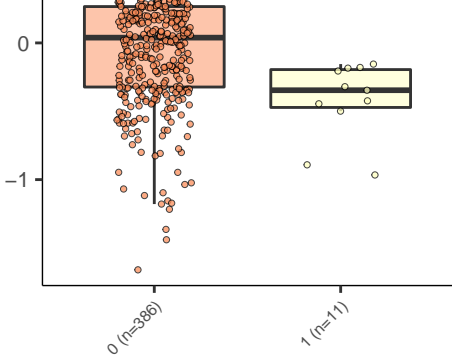


PWY0.166

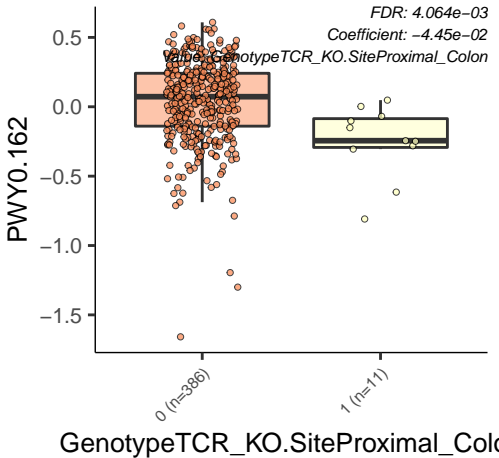
FDR: 3.565e-03

Coefficient: -5.54e-02

Value: GenotypeTCR_KO.SiteProximal_Colon



GenotypeTCR_KO.SiteProximal_Colo



PWY.7184

FDR: $9.273\text{e-}03$

Coefficient: $-1.03\text{e-}01$

Value: GenotypeTCR_KO.SiteProximal_Colon

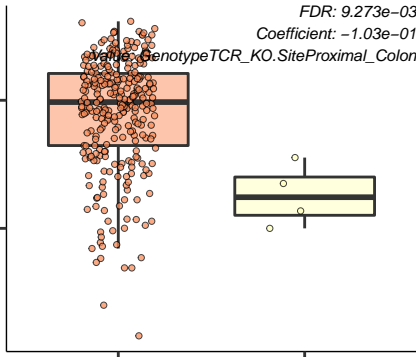
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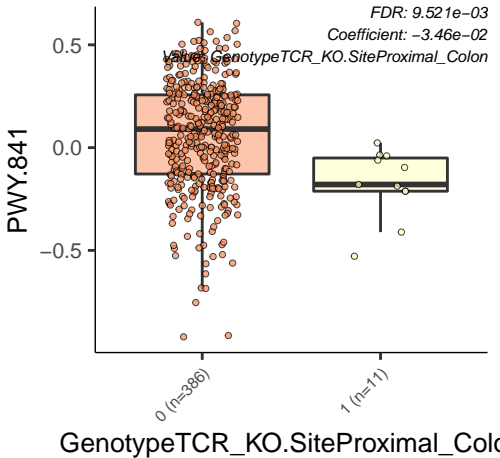
-1

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteProximal_Colo



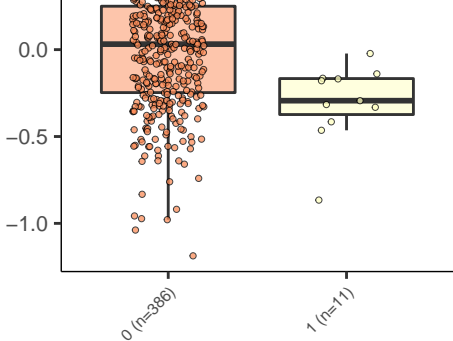


PWY.7187

FDR: 1.089e-02

Coefficient: -4.17e-02

Value: GenotypeTCR_KO.SiteProximal_Colon



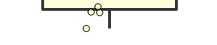
GenotypeTCR_KO.SiteProximal_Colon

COBALSYN.PWY

FDR: 1.463×10^{-2}

Coefficient: 5.07×10^{-2}

Value: GenotypeTCR_KO.SiteProximal_Colon



0 (n=386)

1 (n=11)

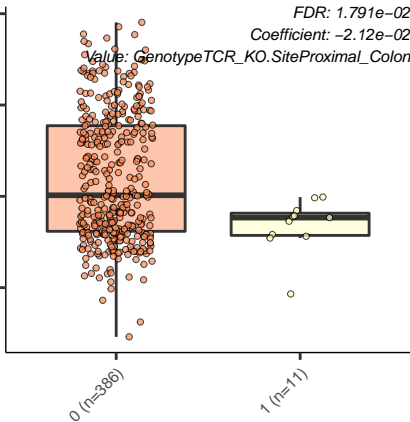
GenotypeTCR_KO.SiteProximal_Colon

FASYN.ELONG.PWY

FDR: 1.791e-02

Coefficient: -2.12e-02

Value: GenotypeTCR_KO.SiteProximal_Colon



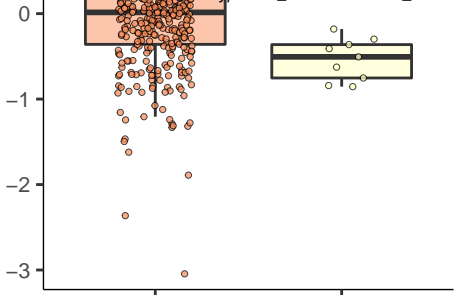
GenotypeTCR_KO.SiteProximal_Colo

PWY.7228

FDR: $1.993e-02$

Coefficient: $-6.27e-02$

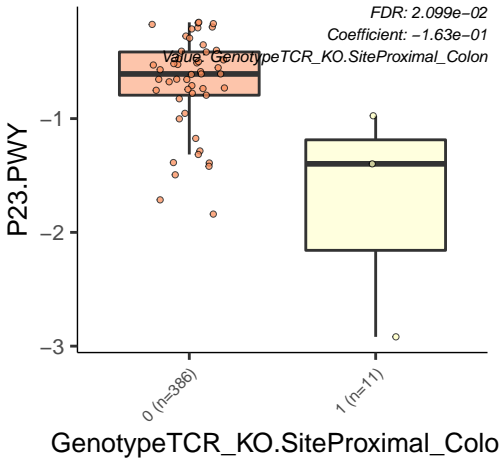
Value: GenotypeTCR_KO.SiteProximal_Colon



0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteProximal_Colo



HOMOSER.METSYN.PWY

FDR: 2.730e-02

Coefficient: 1.00e-01

Valpha_GenotypeTCR_KO.SiteProximal_Colon

0.0

-0.5

-1.0

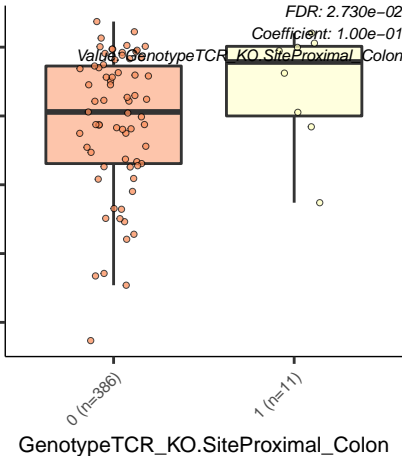
-1.5

-2.0

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteProximal_Colon



PWY.6269

FDR: 2.790×10^{-2}

Coefficient: 4.72×10^{-2}

GenotypeTCR_KO.SiteProximal_Colon

0

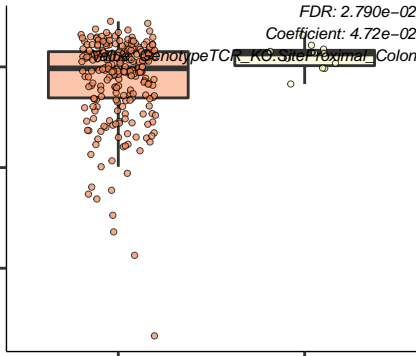
-1

-2

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteProximal_Colo

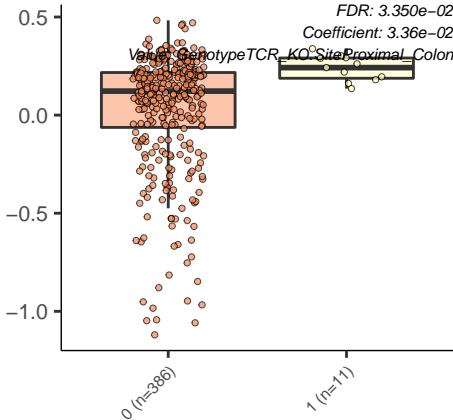


PWY.7400

FDR: 3.350e-02

Coefficient: 3.36e-02

Value: GenotypeTCR_KO.SiteProximal_Colon



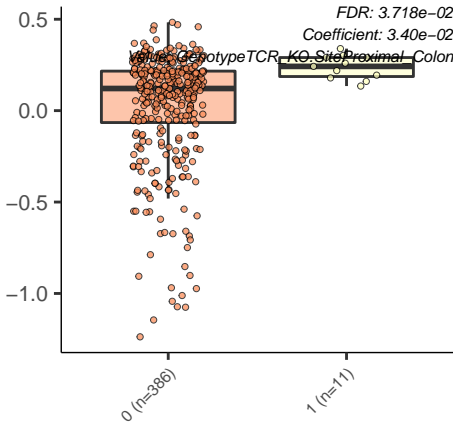
GenotypeTCR_KO.SiteProximal_Colon

ARGSYN.PWY

FDR: 3.718×10^{-2}

Coefficient: 3.40×10^{-2}

Value: GenotypeTCR_KO.SiteProximal_Colon



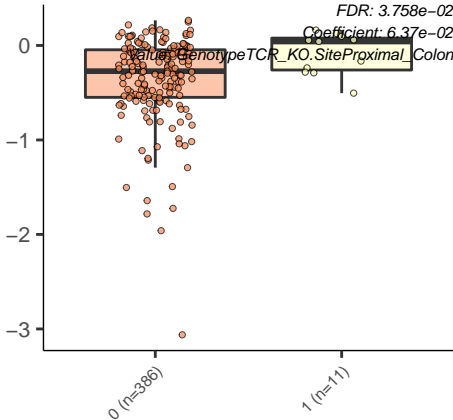
GenotypeTCR_KO.SiteProximal_Colon

PWY.5189

FDR: 3.758e-02

Coefficient: 6.37e-02

Value: GenotypeTCR_KO.SiteProximal_Colon



GenotypeTCR_KO.SiteProximal_Colo

PYRIDNUCSAL.PWY

FDR: $4.091e-02$

Coefficient: $4.41e-02$

GenotypeTCR_KO.SiteProximal_Colon

0

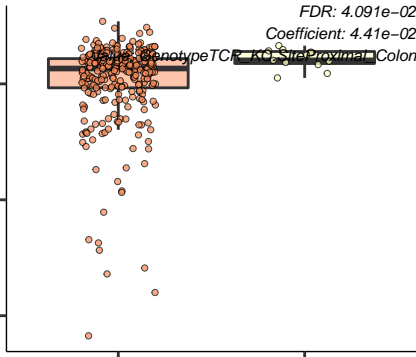
-1

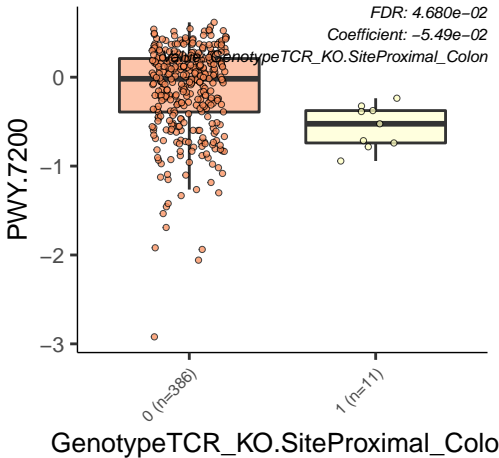
-2

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteProximal_Colo



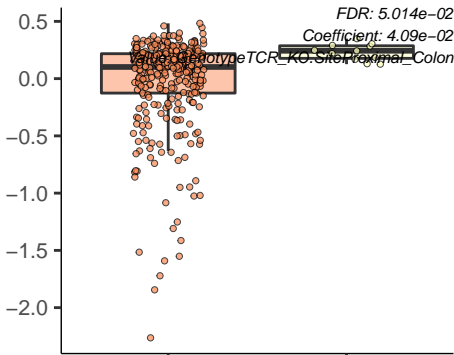


ARGSYNBSUB.PWY

FDR: 5.014e-02

Coefficient: 4.09e-02

Value of GenotypeTCR_KO.SiteProximal_Colon



0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteProximal_Colon

DENOVOPURINE2.PWY

FDR: $5.146e-02$

Coefficient: $-3.16e-02$

Value: GenotypeTCR_KO.SiteProximal_Colon

0

-1

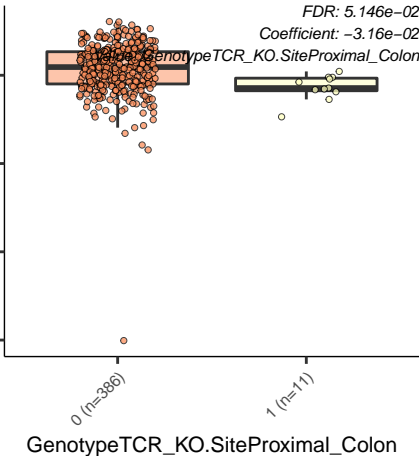
-2

-3

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteProximal_Colon



PWY.5509

FDR: 5.339e-02

Coefficient: 4.94e-02

Value-GenotypeTCR_KO.SiteProximal_Colon

0

-1

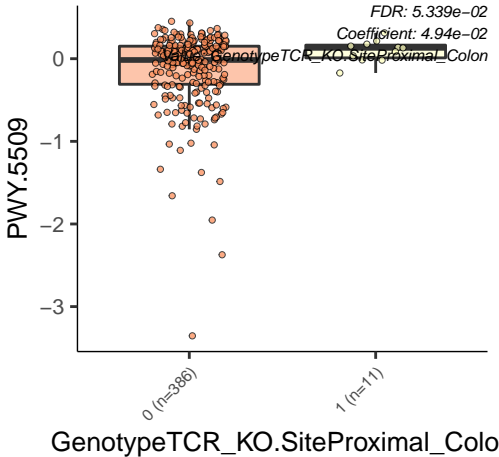
-2

-3

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteProximal_Colo

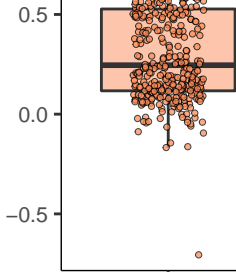


PWY.5659

FDR: 6.039×10^{-2}

Coefficient: -2.05×10^{-2}

Value: GenotypeTCR_KO.SiteProximal_Colon



0 (n=386)

1 (n=11)

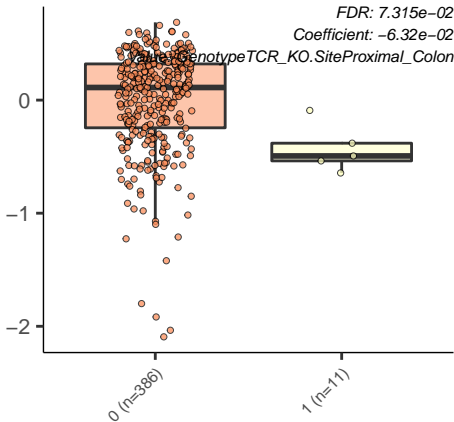
GenotypeTCR_KO.SiteProximal_Colon

PWY.7392

FDR: 7.315e-02

Coefficient: -6.32e-02

Value: GenotypeTCR_KO.SiteProximal_Colon



GenotypeTCR_KO.SiteProximal_Colo

PWY.5505

FDR: 7.653e-02

Coefficient: 4.05e-02

Value: GenotypeTCR_KO.SiteProximal_Colon

0

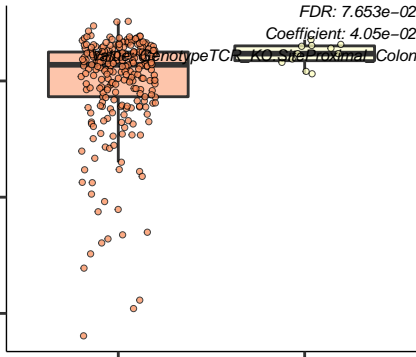
-1

-2

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteProximal_Colo

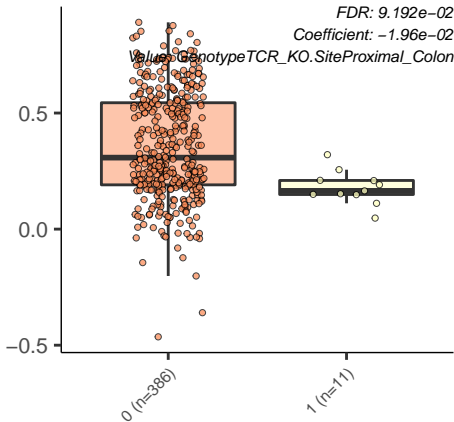


PWY0.1586

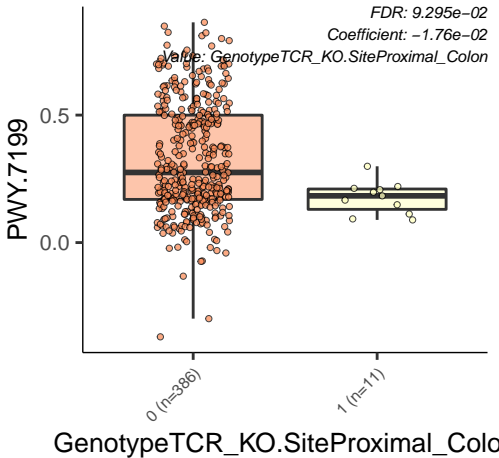
FDR: 9.192×10^{-2}

Coefficient: -1.96×10^{-2}

Value GenotypeTCR_KO.SiteProximal_Colon



GenotypeTCR_KO.SiteProximal_Colon



PWY.6125

FDR: 1.172e-01

Coefficient: -4.03e-02

Value: GenotypeTCR_KO.SiteProximal_Colon

0

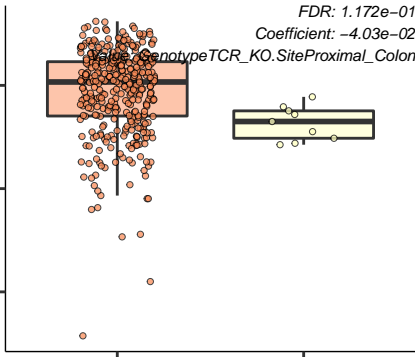
-1

-2

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteProximal_Colo



PWY.5347

FDR: 1.233e-01

Coefficient: 5.02e-02

Value GenotypeTCR_KO.SiteProximal_Colon

0.0

-0.5

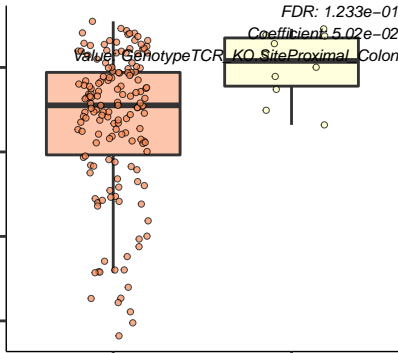
-1.0

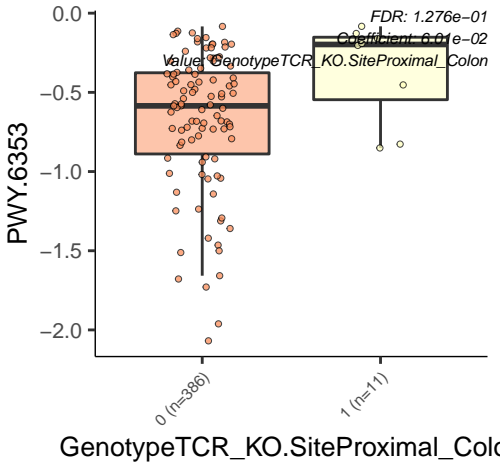
-1.5

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteProximal_Colon





PWY.7196

FDR: 1.359e-01

Coefficient: -3.75e-02

Value: GenotypeTCR_KO.SiteProximal_Colon

0

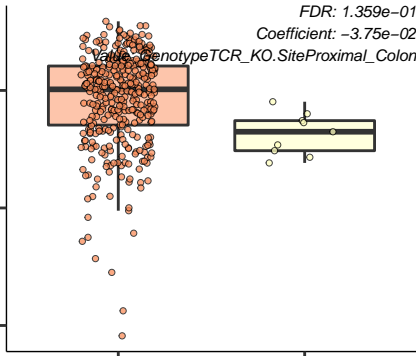
-1

-2

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteProximal_Colo



GALACTUROCAT.PWY

FDR: 1.572e-01

Coefficient: 2.73e-02

Value GenotypeTCR_KO.SiteProximal_Colon

0.5

0.0

-0.5

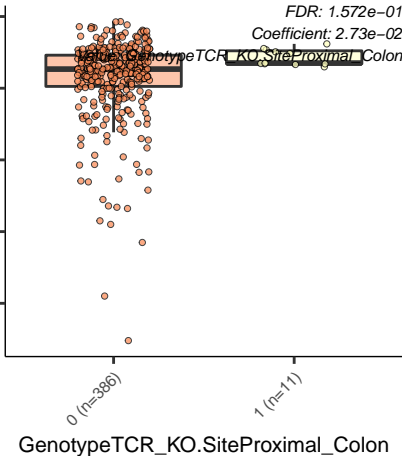
-1.0

-1.5

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteProximal_Colon

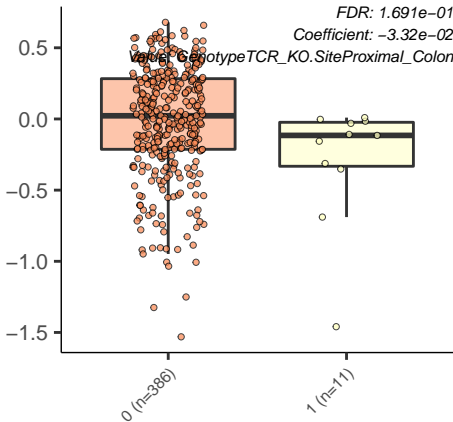


P124.PWY

FDR: 1.691e-01

Coefficient: -3.32e-02

Value GenotypeTCR_KO.SiteProximal_Colon



GenotypeTCR_KO.SiteProximal_Colon

PWY.5188

FDR: 1.759e-01

Coefficient: 3.87e-02

Value: GenotypeTCR_KO.SiteProximal_Colon

0.0

-0.5

-1.0

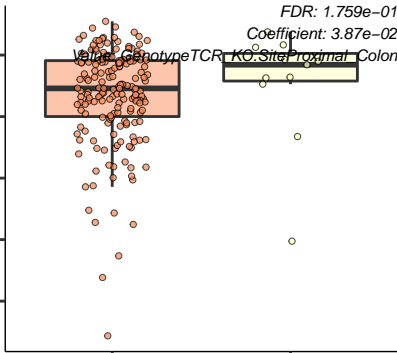
-1.5

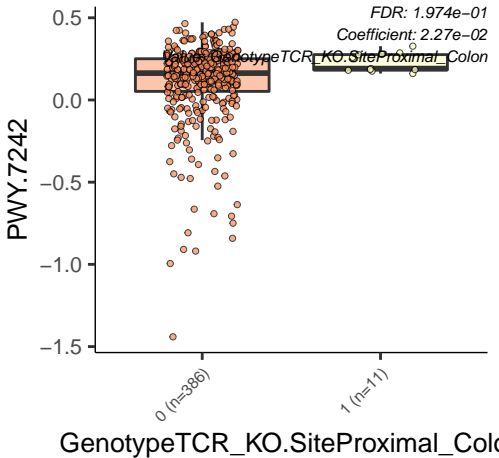
-2.0

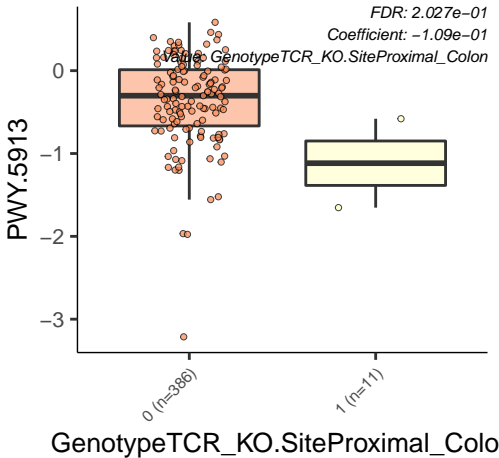
0 (n=386)

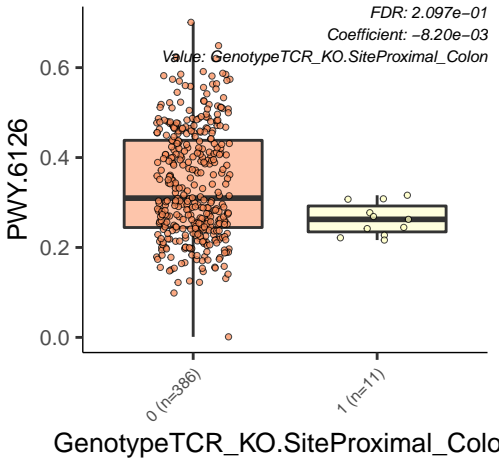
1 (n=11)

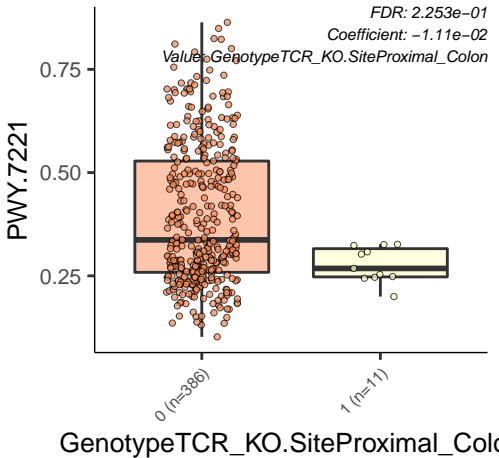
GenotypeTCR_KO.SiteProximal_Colon











PWY.621

FDR: 2.289e-01

Coefficient: 1.77e-02

Valve GenotypeTCR_KO.SiteProximal_Colon

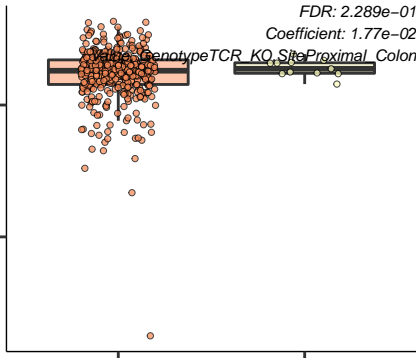
0

-1

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteProximal_Colo



GLUTORN.PWY

