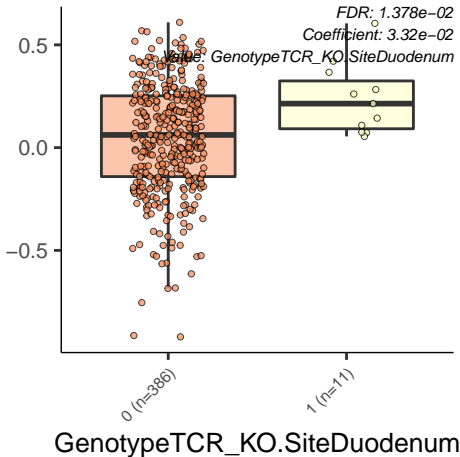
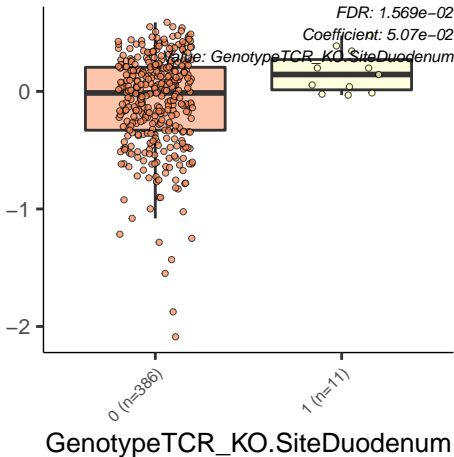


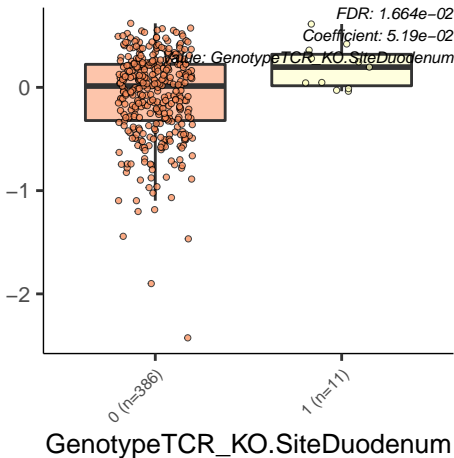
PWY.841



PWY.7196



PWY.6125



COMPLETE.ARO.PWY

FDR: 2.240e-02

Coefficient: -1.53e-02

Value: GenotypeTCR_KO.SiteDuodenum

0.50

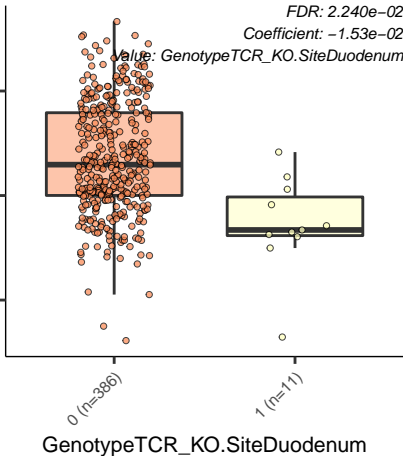
0.25

0.00

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteDuodenum



P42.PWY

FDR: $2.504e-02$

Coefficient: $-3.34e-02$

Value: GenotypeTCR_KO.SiteDuodenum

0

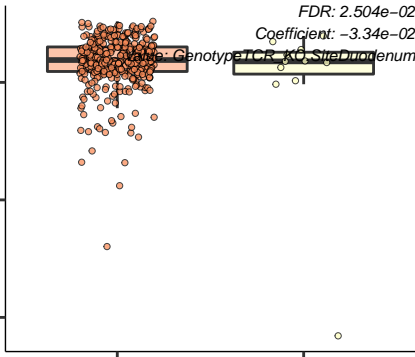
-1

-2

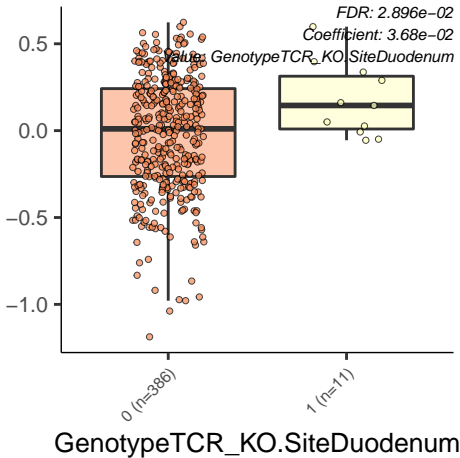
0 (n=386)

1 (n=11)

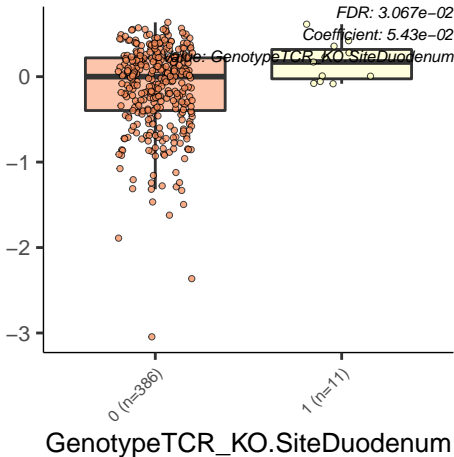
GenotypeTCR_KO.SiteDuodenum

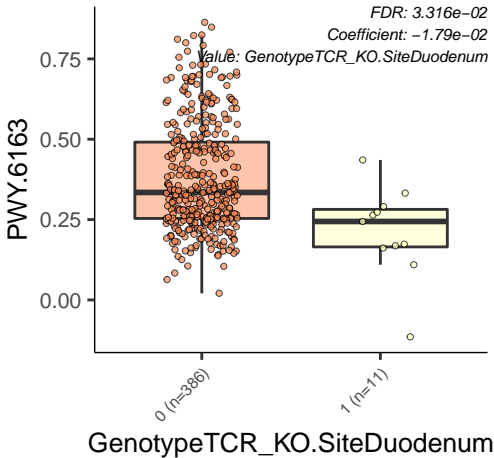


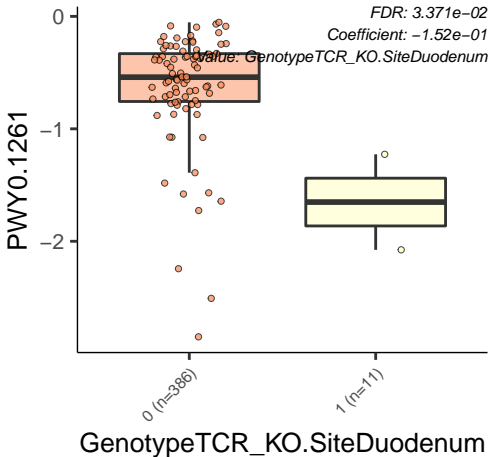
PWY.7187

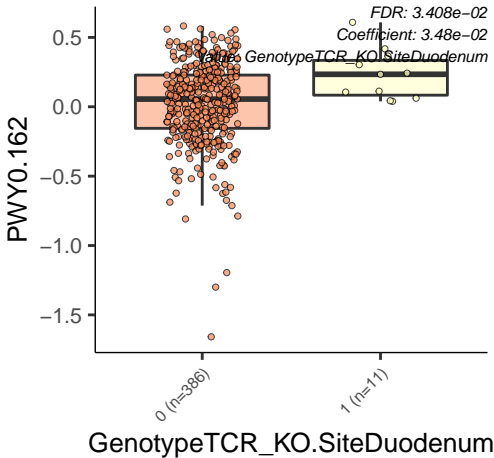


PWY.7228

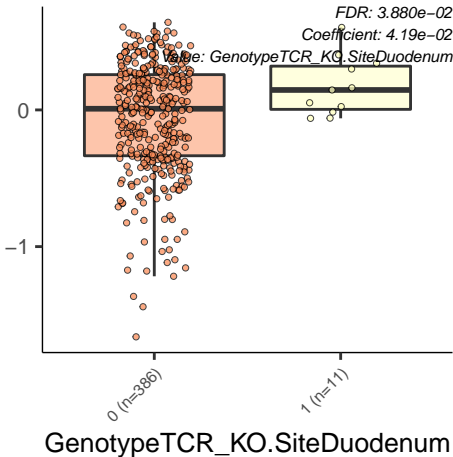


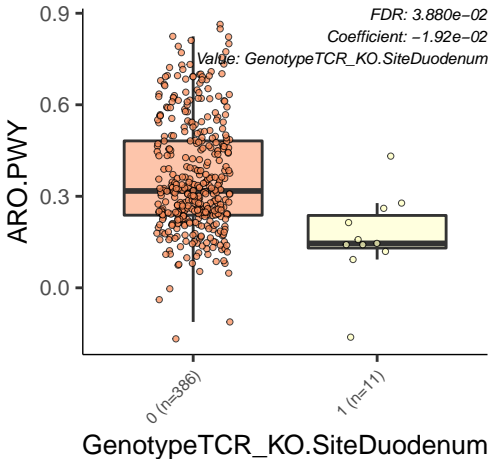






PWY0.166





DENOVOPURINE2.PWY

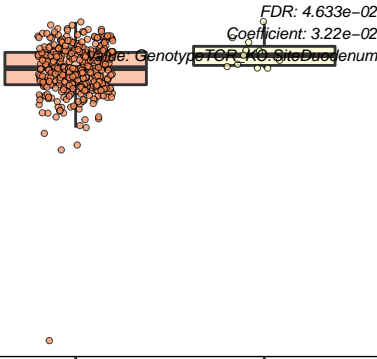
0
-1
-2
-3

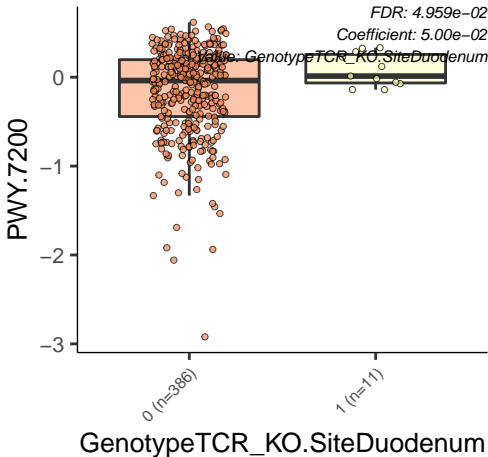
0 (n=386)

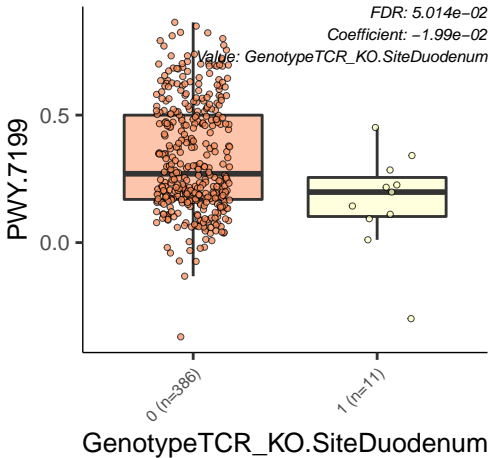
1 (n=11)

GenotypeTCR_KO.SiteDuodenum

FDR: 4.633e-02
Coefficient: 3.22e-02
Value: GenotypeTCR_KO.SiteDuodenum







NONMEVIP.PWY

FDR: 6.193e-02

Coefficient: -1.60e-02

Value: GenotypeTCR_KO.SiteDuodenum

0.75

0.50

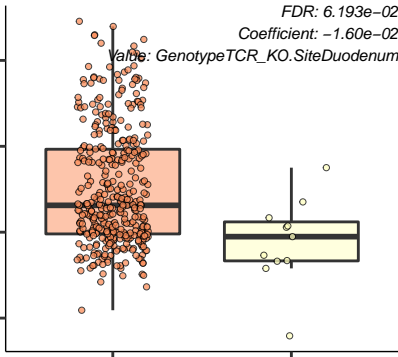
0.25

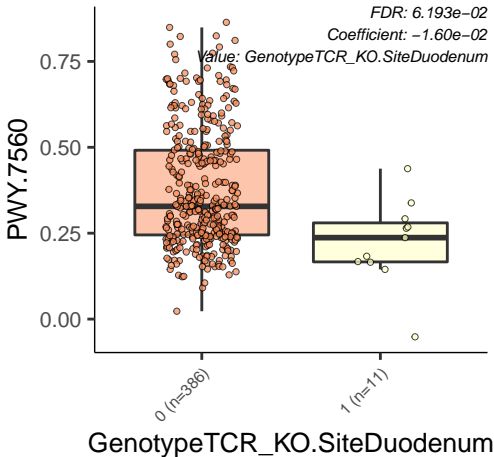
0.00

0 (n=386)

1 (n=11)

GenotypeTCR_KO.SiteDuodenum



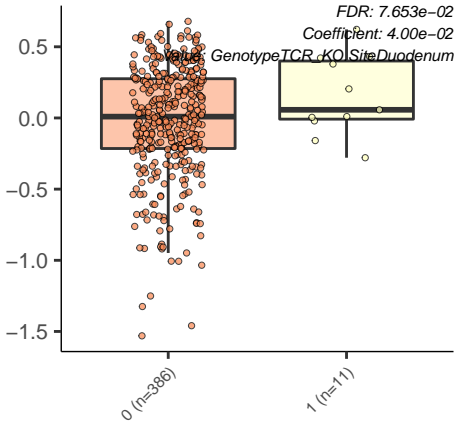


P124.PWY

FDR: $7.653e-02$

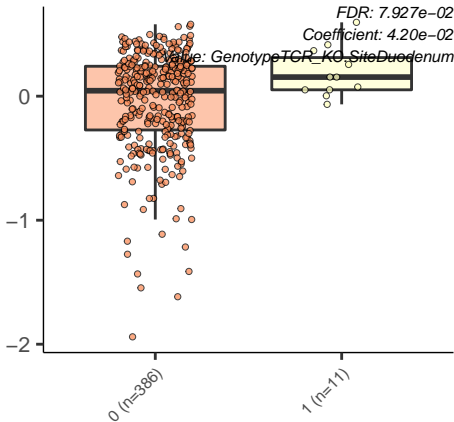
Coefficient: $4.00e-02$

Value: GenotypeTCR_KO.SiteDuodenum



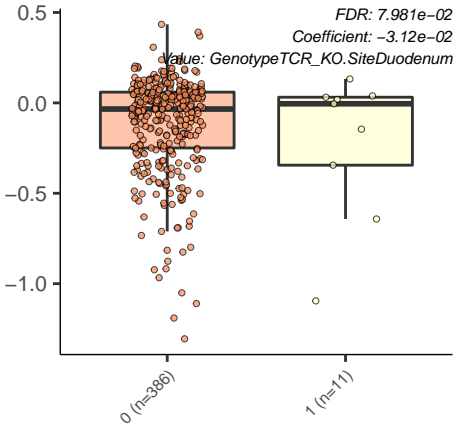
GenotypeTCR_KO.SiteDuodenum

PWY.7234



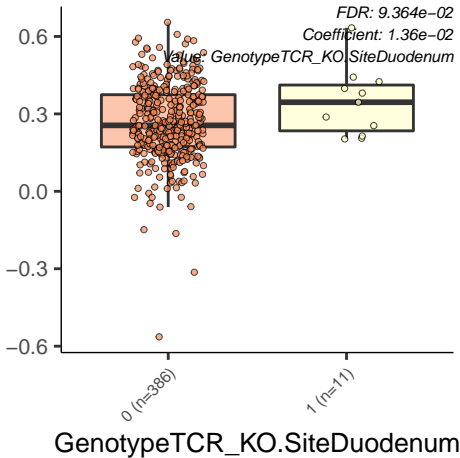
GenotypeTCR_KO.SiteDuodenum

PWY.5154

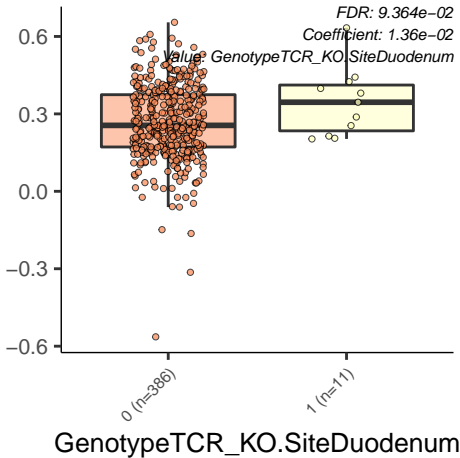


GenotypeTCR_KO.SiteDuodenum

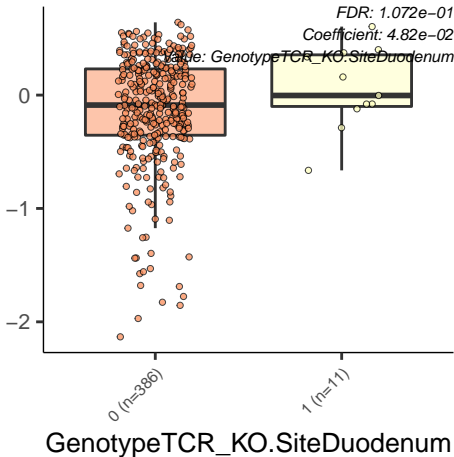
PWY.7220



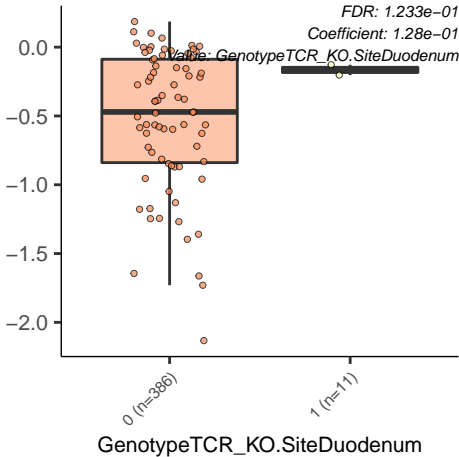
PWY.7222



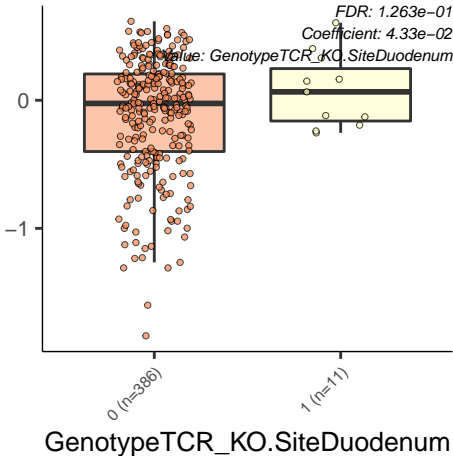
P122.PWY

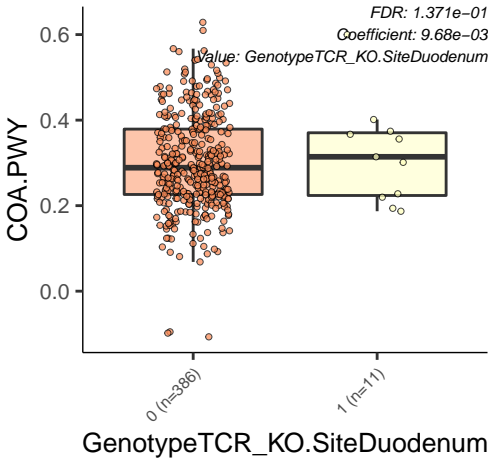


HOMOSER.METSYN.PWY

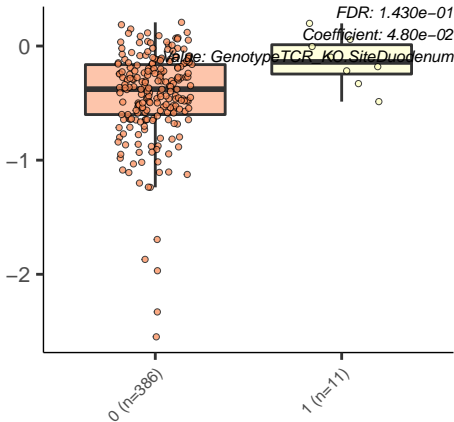


PWY.7184



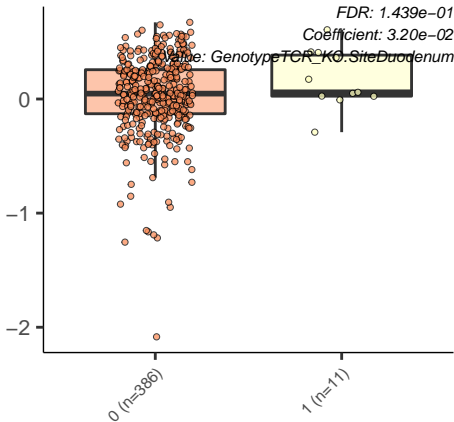


PRPP.PWY



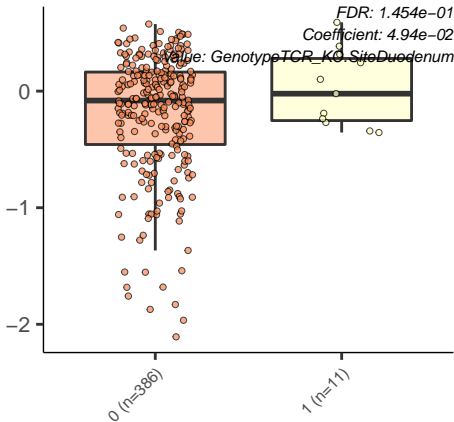
GenotypeTCR_KO.SiteDuodenum

PWY.6471



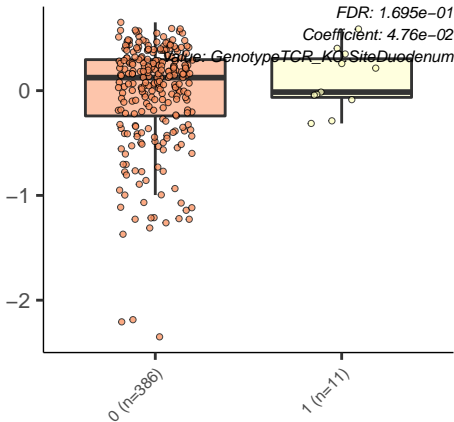
GenotypeTCR_KO.SiteDuodenum

PWY.7197

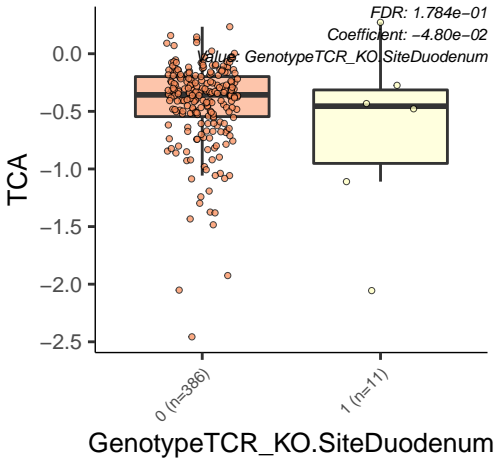


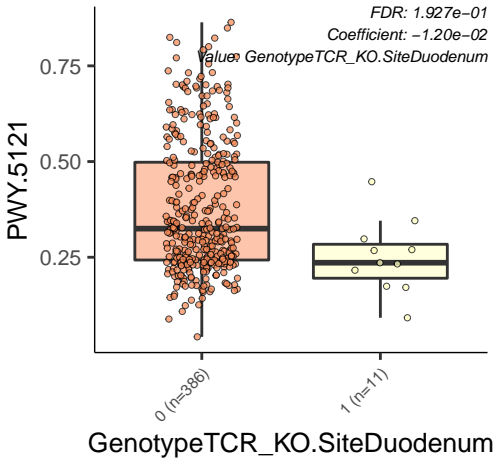
GenotypeTCR_KO.SiteDuodenum

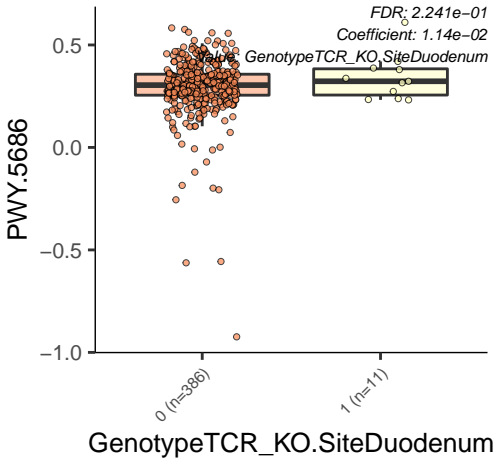
PWY.2941



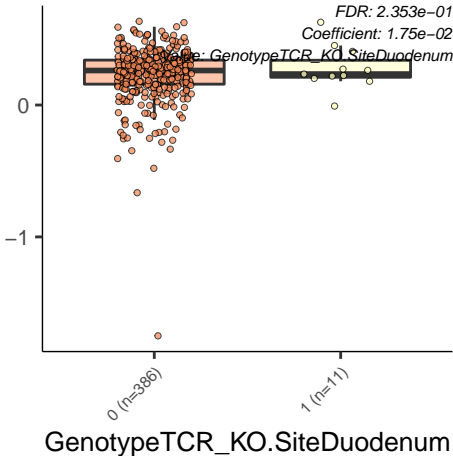
GenotypeTCR_KO.SiteDuodenum



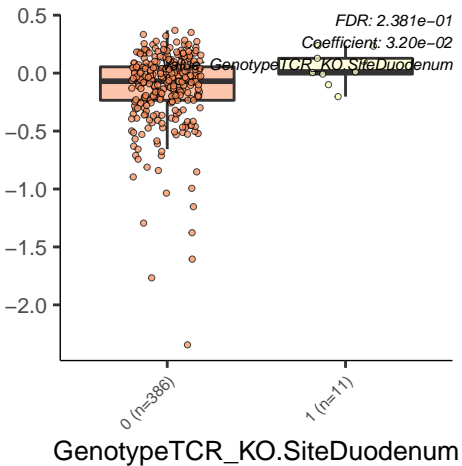




PWY.621



GLYCOLYSIS.E.D

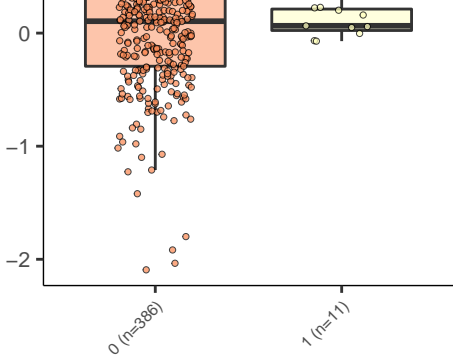


PWY.7392

FDR: 2.393e-01

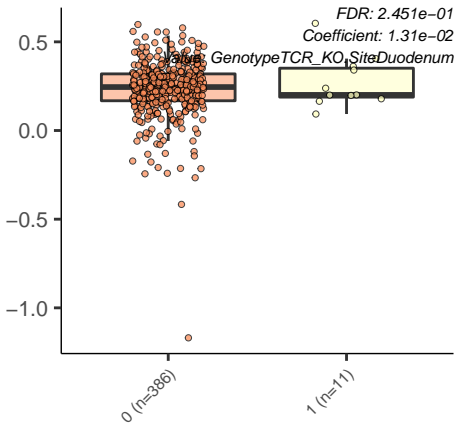
Coefficient: 3.27e-02

Value: GenotypeTCR_KO.SiteDuodenum



GenotypeTCR_KO.SiteDuodenum

PWY.6317



GenotypeTCR_KO.SiteDuodenum