

CITRULBIO_PWY_L_citrulline_biosyn

0.020
0.015
0.010
0.005
0.000

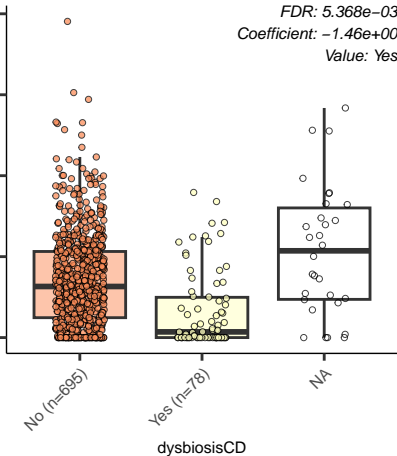
FDR: 5.368e-03
Coefficient: -1.46e+00
Value: Yes

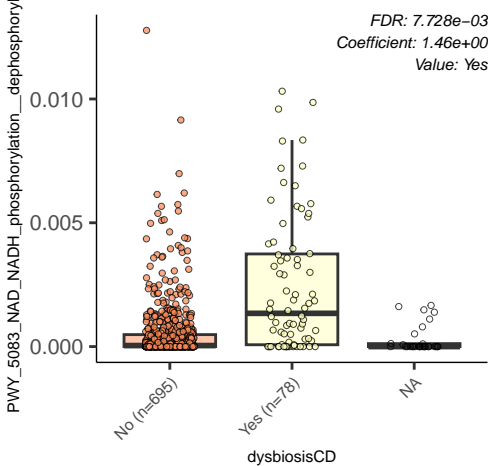
No (n=695)

Yes (n=78)

NA

dysbiosisCD





PWY66_400_glycolysis_VI

0.05
0.04
0.03
0.02
0.01
0.00

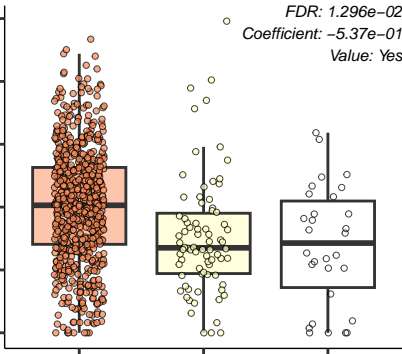
No (n=695)

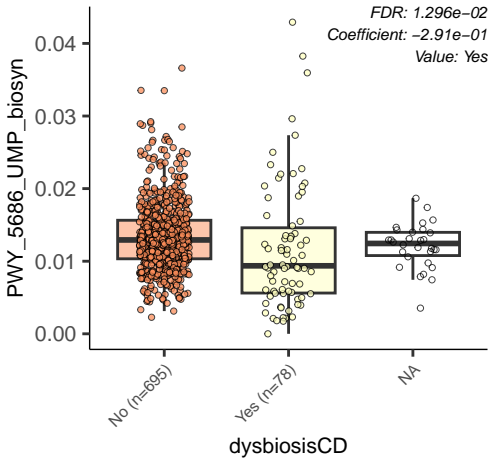
Yes (n=78)

NA

dysbiosisCD

FDR: 1.296e-02
Coefficient: -5.37e-01
Value: Yes





PWY_5723_Rubisco_shunt

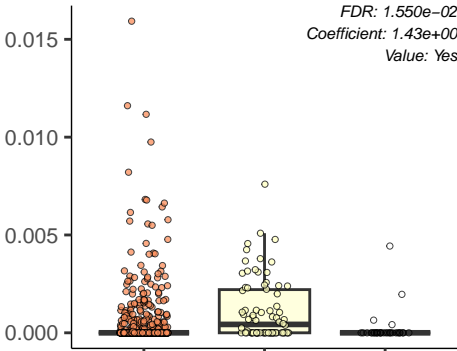
FDR: 1.550e-02
Coefficient: 1.43e+00
Value: Yes

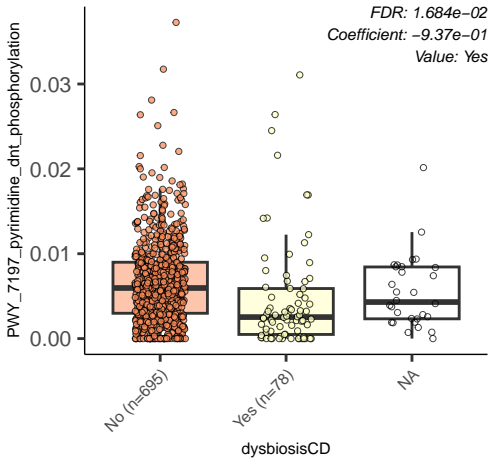
No (n=695)

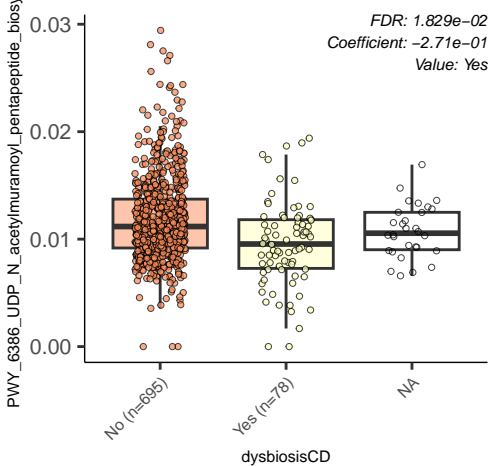
Yes (n=78)

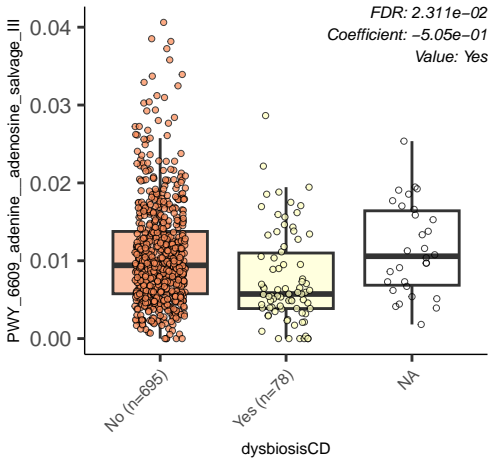
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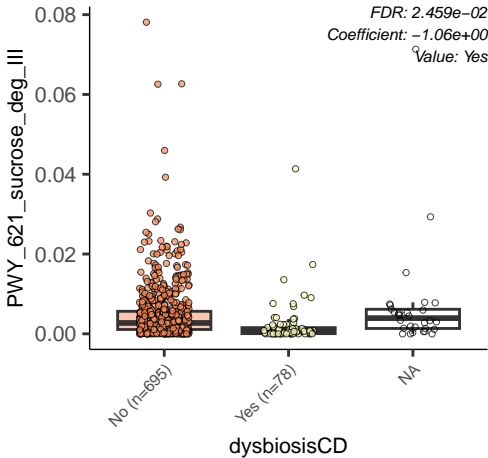
dysbiosisCD

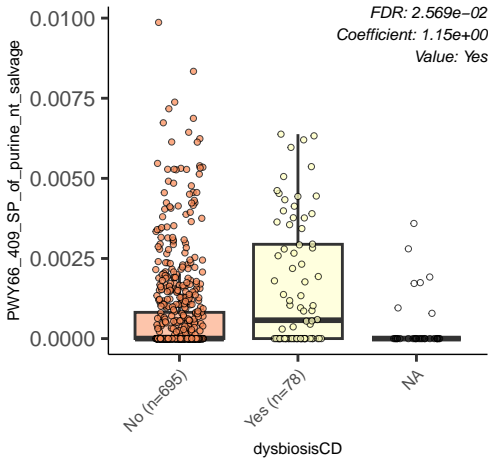


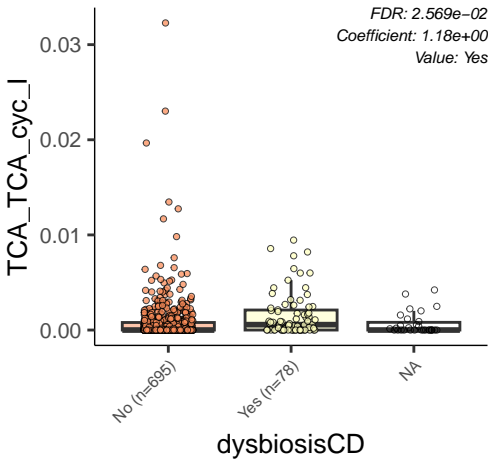


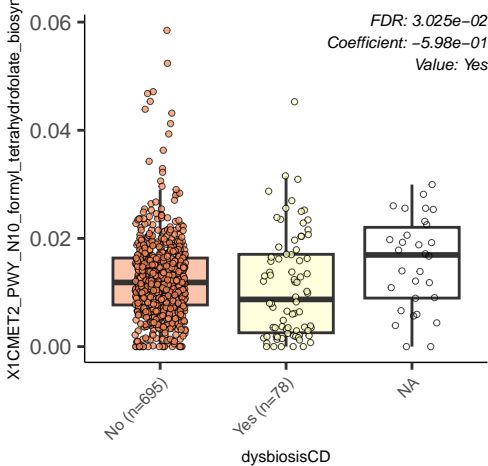


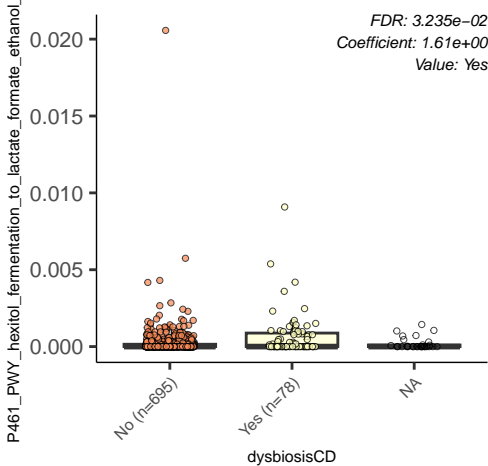


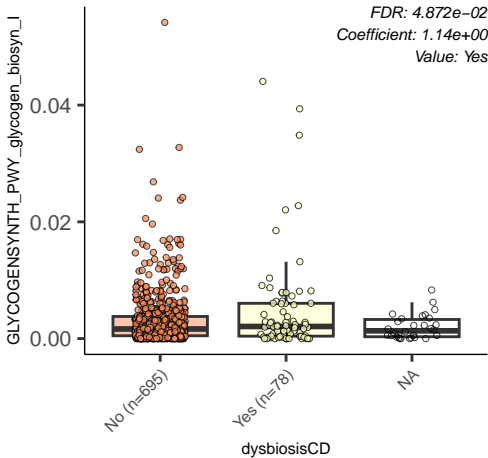


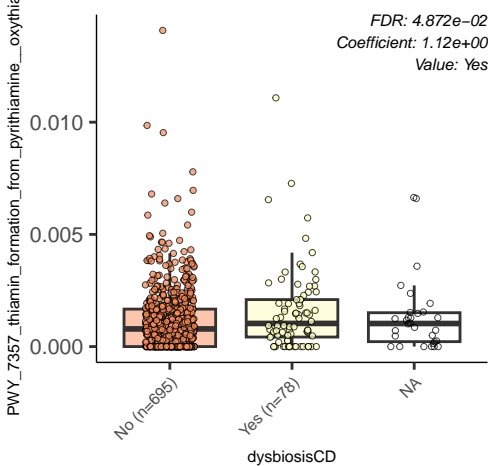


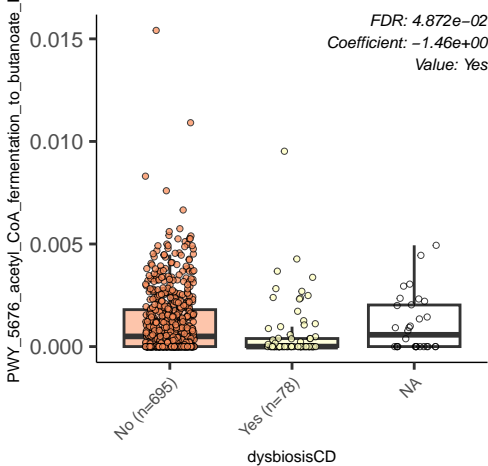


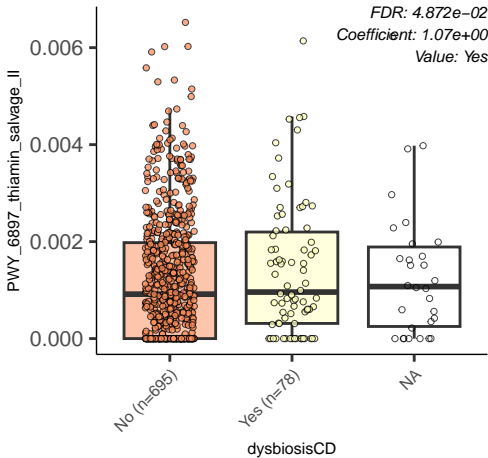


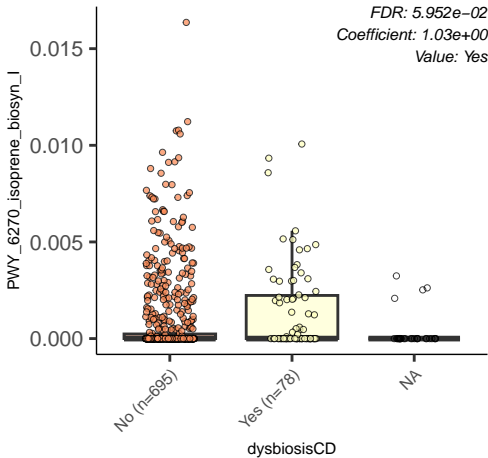


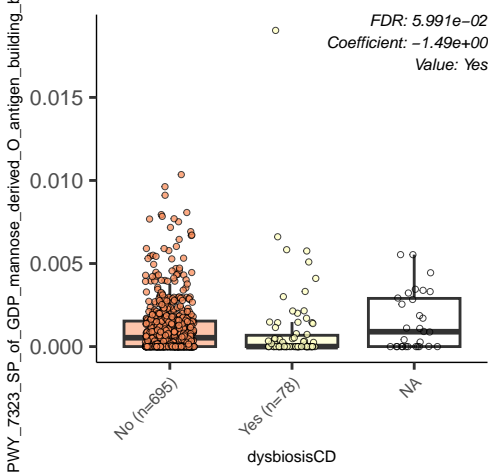


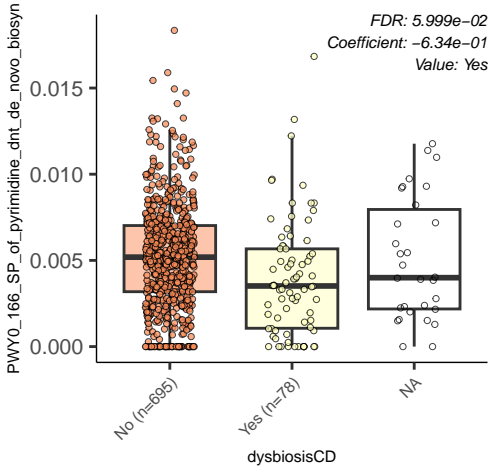


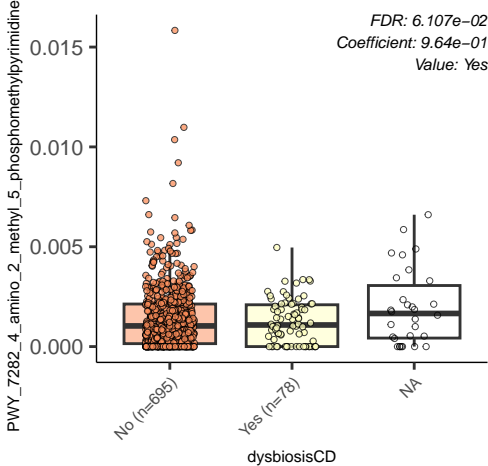


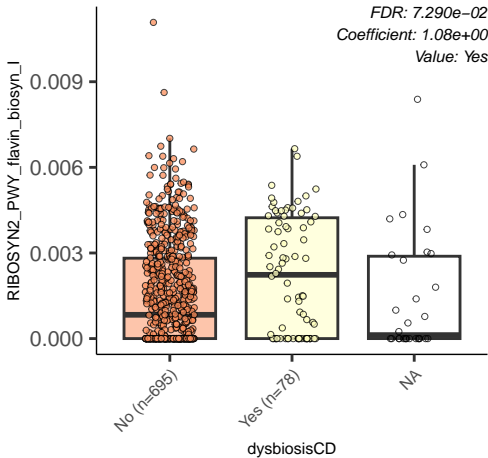


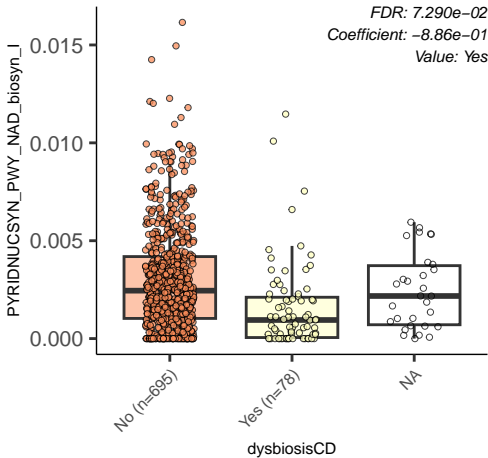


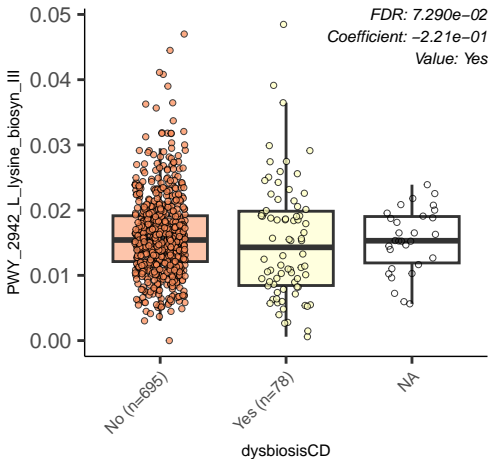


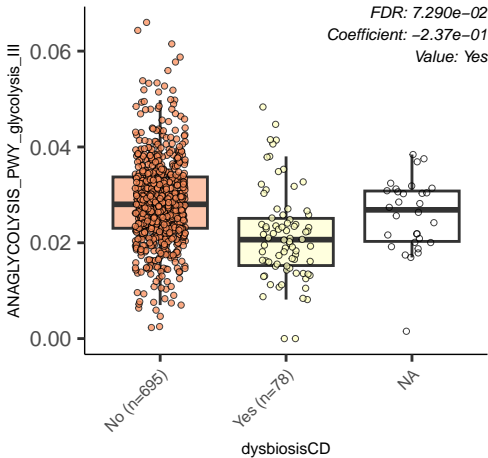


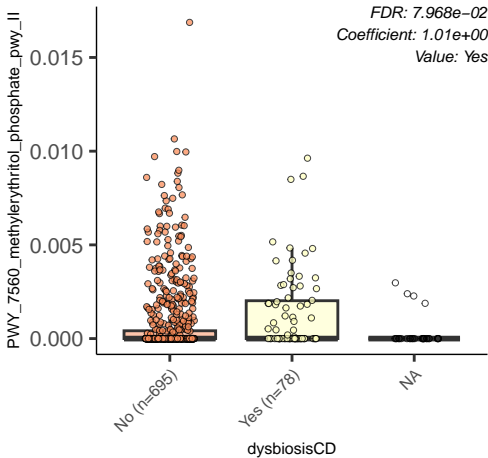












PWY_1042_glycolysis_IV

FDR: $8.525e-02$

Coefficient: $-2.37e-01$

Value: Yes

0.09

0.06

0.03

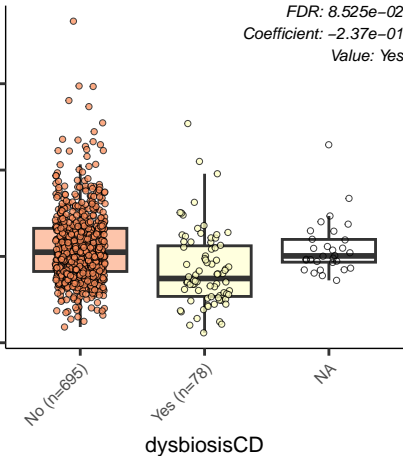
0.00

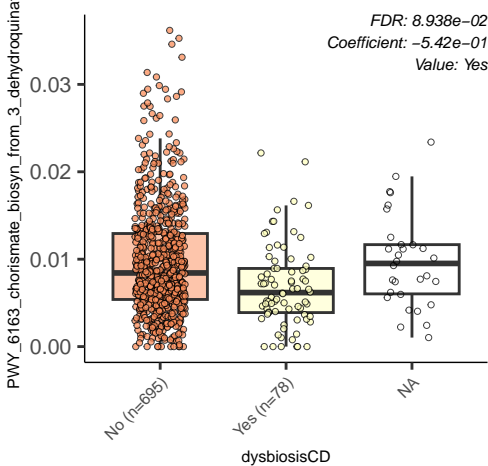
No (n=695)

Yes (n=78)

NA

dysbiosisCD





PWY_5690_TCA_cyc_II

FDR: 1.010e-01
Coefficient: 1.14e+00
Value: Yes

0.02

0.01

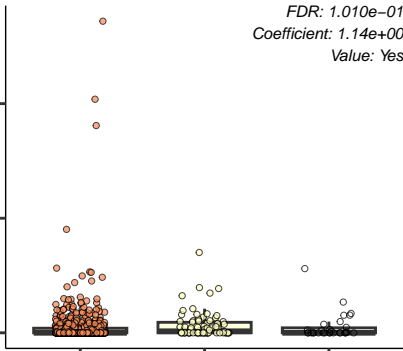
0.00

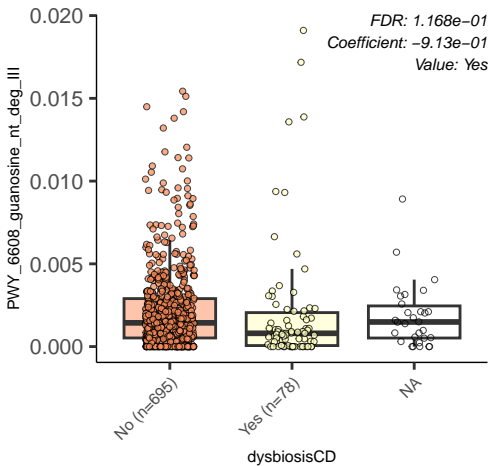
No (n=695)

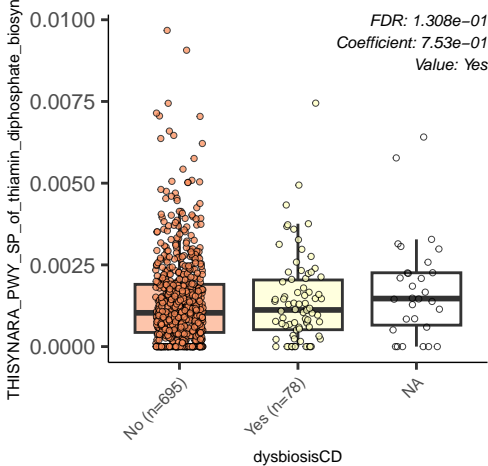
Yes (n=78)

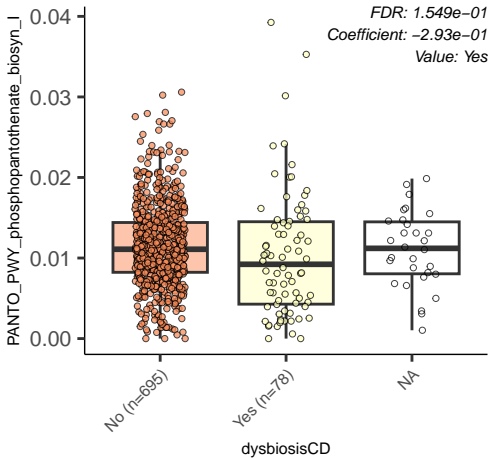
NA

dysbiosisCD









COA_PWY_1_coenzyme_A_biosyn_II

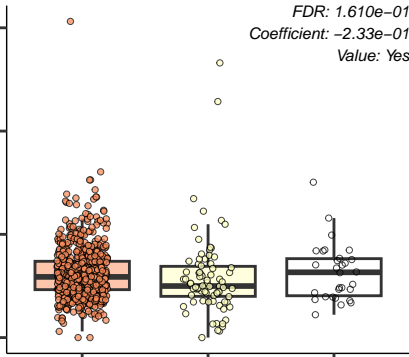
FDR: 1.610e-01
Coefficient: -2.33e-01
Value: Yes

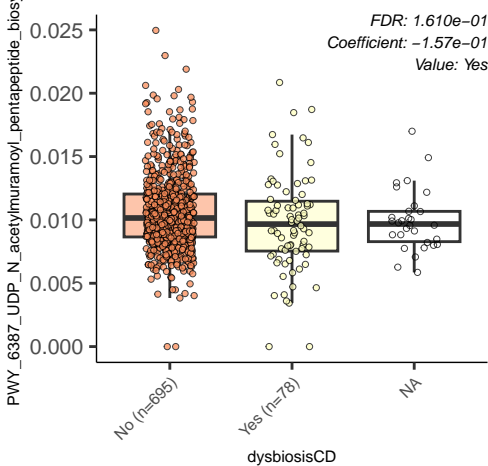
No (n=695)

Yes (n=78)

NA

dysbiosisCD





PWY_4984_urea_cyc

0.0100
0.0075
0.0050
0.0025
0.0000

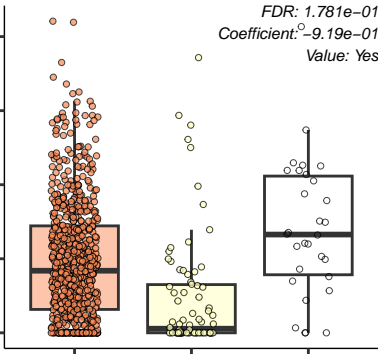
FDR: 1.781e-01
Coefficient: -9.19e-01
Value: Yes

No (n=695)

Yes (n=78)

NA

dysbiosisCD



PWY66_399_gluconeogenesis_III

FDR: $1.908e-01$
Coefficient: $-1.10e+00$
Value: Yes

0.04

0.02

0.00

No (n=695)

Yes (n=78)

NA

dysbiosisCD

