

PWY0.1241..ADP.L.glycero..beta..D.manno.heptose.biosynt

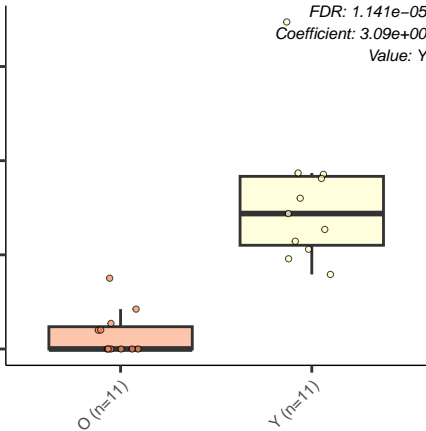
FDR: 1.141e-05
Coefficient: 3.09e+00
Value: Y

30
20
10
0

O (n=11)

Y (n=11)

Groups



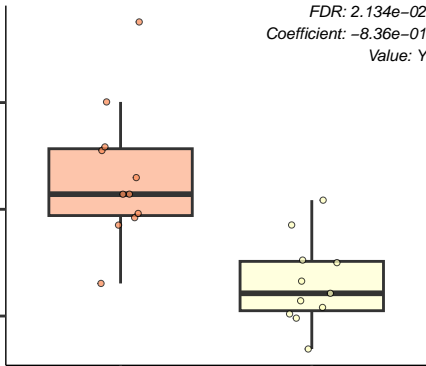
PWY.6606..guanosine.nucleotides.degradation.II

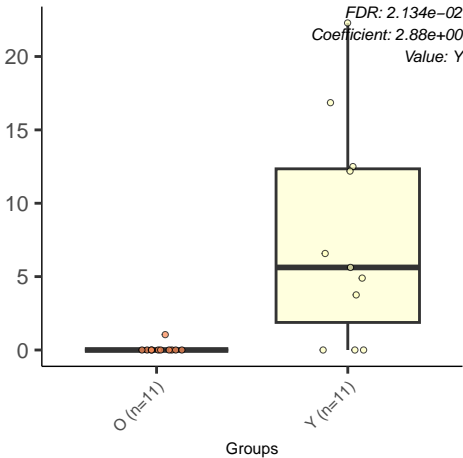
FDR: 2.134e-02
Coefficient: -8.36e-01
Value: Y

O (n=11)

Y (n=11)

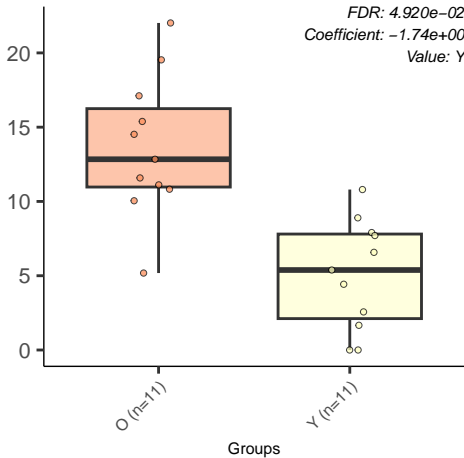
Groups





PWY.241..C4.photosynthetic.carbon.assimilation.cycle..NADP.

FDR: 4.920e-02
Coefficient: -1.74e+00
Value: Y



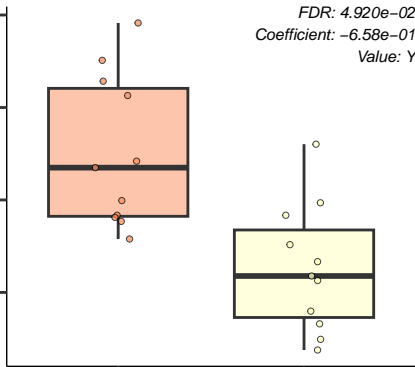
PWY.6527...stachyose.degradation

FDR: 4.920e-02
Coefficient: -6.58e-01
Value: Y

O (n=11)

Y (n=11)

Groups



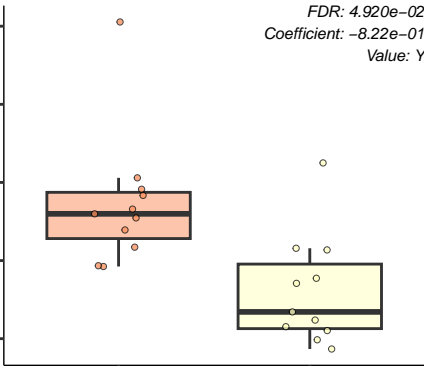
PWY.6608..guanosine.nucleotides.degradation.III

FDR: 4.920e-02
Coefficient: -8.22e-01
Value: Y

O (n=11)

Y (n=11)

Groups



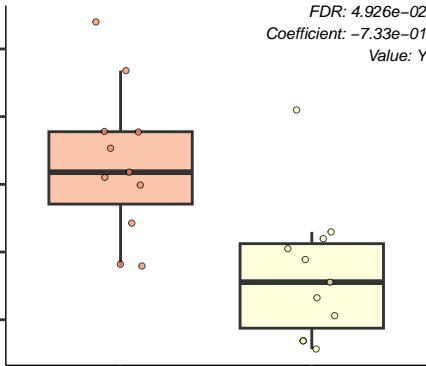
P164.PWY..purine.nucleobases.degradation.l..anaerobic

FDR: 4.926e-02
Coefficient: -7.33e-01
Value: Y

O (n=11)

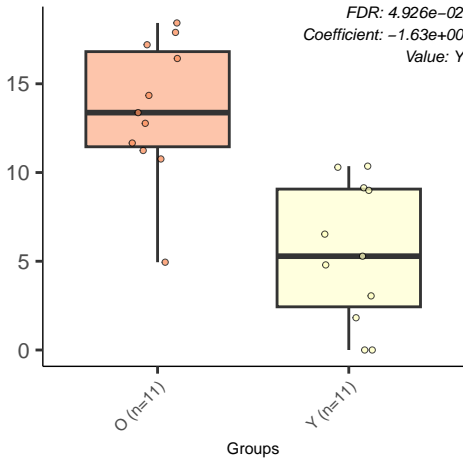
Y (n=11)

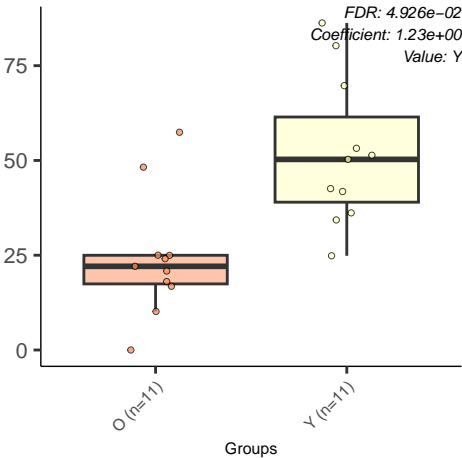
Groups



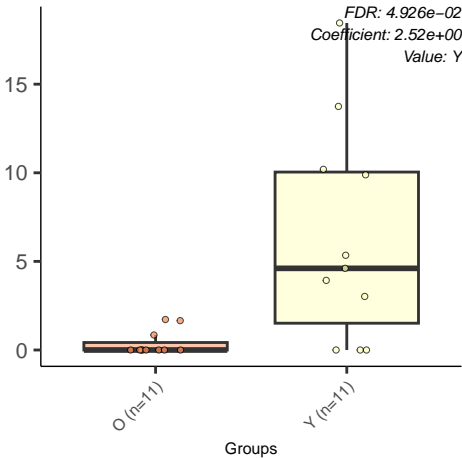
PWY.7117..C4.photosynthetic.carbon.assimilation.cycle..PEPC

FDR: 4.926e-02
Coefficient: -1.63e+00
Value: Y



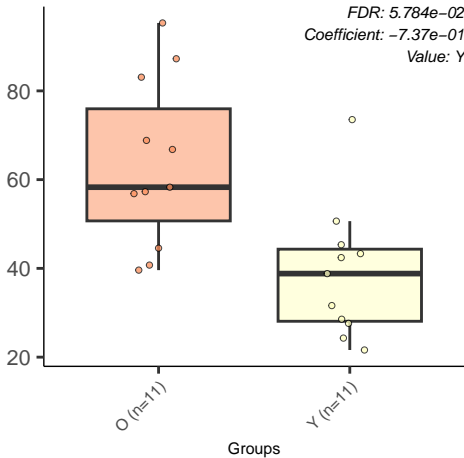


PYRIDOXSYN.PWY..pyridoxal.5..phosphate.biosynthesis



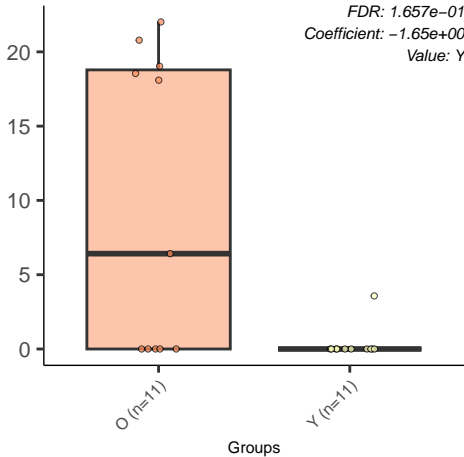
PWY.6545..pyrimidine.deoxyribonucleotides.de.novo.biosynth

FDR: 5.784e-02
Coefficient: -7.37e-01
Value: Y

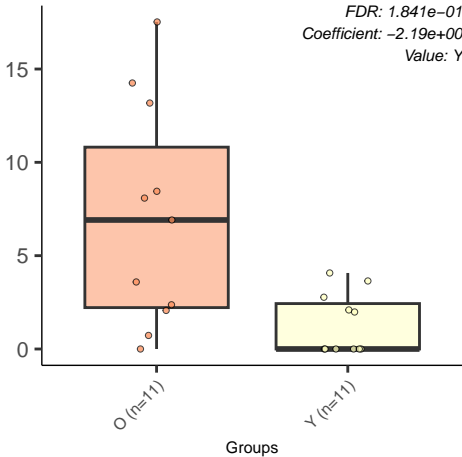


COBALSYN.PWY..adenosylcobalamin.salvage.from.cobinan

FDR: 1.657e-01
Coefficient: -1.65e+00
Value: Y

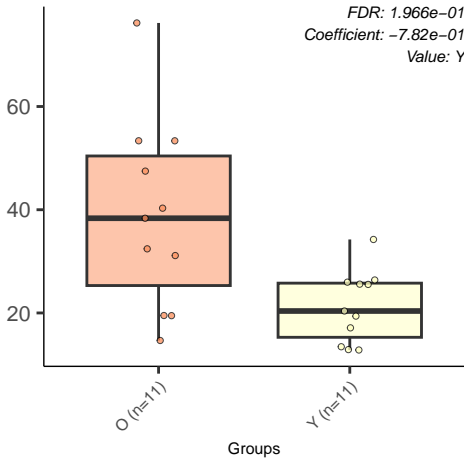


FDR: 1.841e-01
Coefficient: -2.19e+00
Value: Y



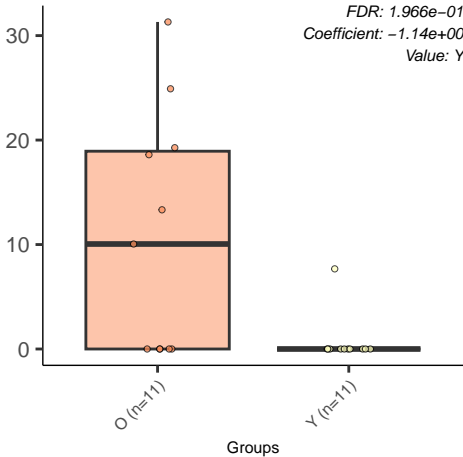
PHOSLIPSYN.PWY..superpathway.of.phospholipid.biosynthesis.

FDR: 1.966e-01
Coefficient: -7.82e-01
Value: Y



PWY.6969..TCA.cycle.V..2.oxoglutarate.ferredoxin.oxidoredu

FDR: 1.966e-01
Coefficient: -1.14e+00
Value: Y



FUCCAT.PWY..fucose.degradation

FDR: 2.293e-01
Coefficient: 1.13e+00
Value: Y

10

5

0

O (n=11)

Y (n=11)

Groups

