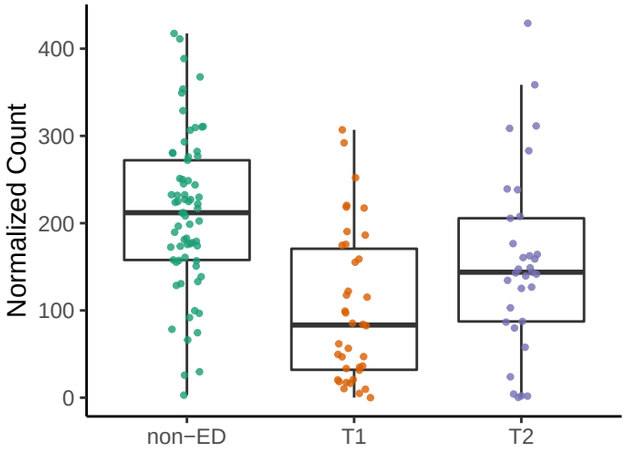
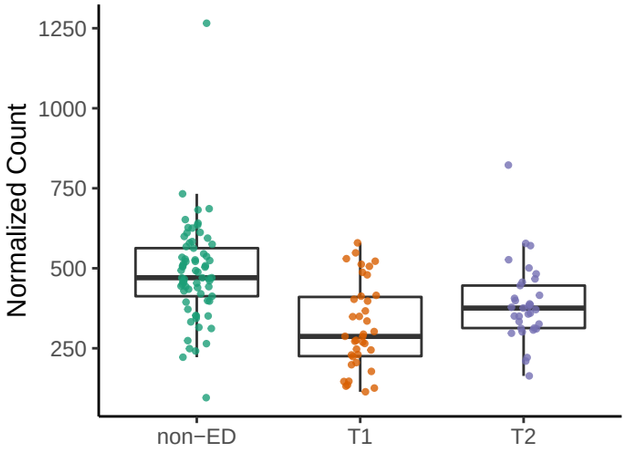
COBALSYN−PWY: adenosylcobalamin

non−ED vs. T1 adjusted p = 7.6e−06
non−ED vs. T2 adjusted p = 0.075
T1 vs. T2 adjusted p = 0.13



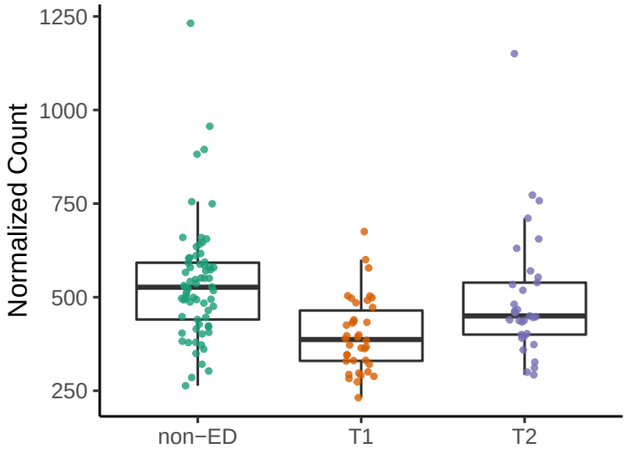
DTDPRHAMSYN−PWY: dTDP−L−rha

non−ED vs. T1 adjusted p = 1.3e−05
non−ED vs. T2 adjusted p = 0.05
T1 vs. T2 adjusted p = 0.13



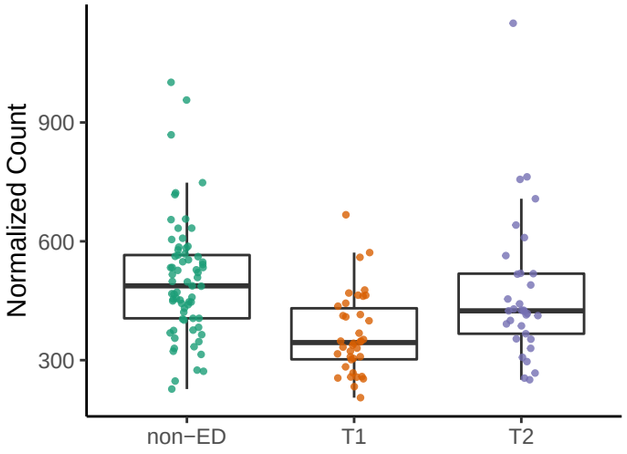
BRANCHED−CHAIN−AA−SYN−PWY:

non−ED vs. T1 adjusted p = 1.7e−05
non−ED vs. T2 adjusted p = 0.38
T1 vs. T2 adjusted p = 0.13



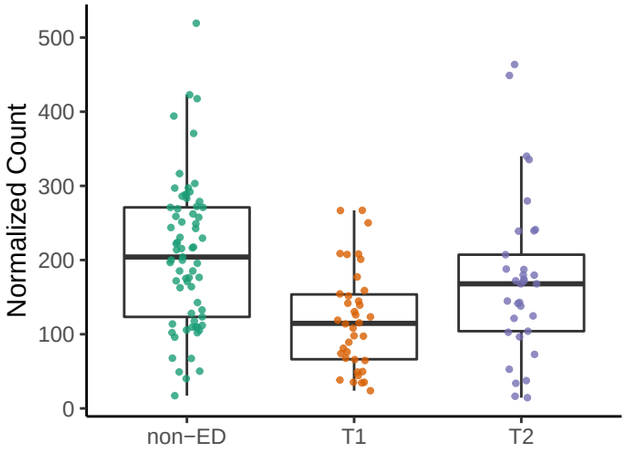
PWY−5103: L−isoleucine biosynthesis

non−ED vs. T1 adjusted p = 1.7e−05
non−ED vs. T2 adjusted p = 0.52
T1 vs. T2 adjusted p = 0.13



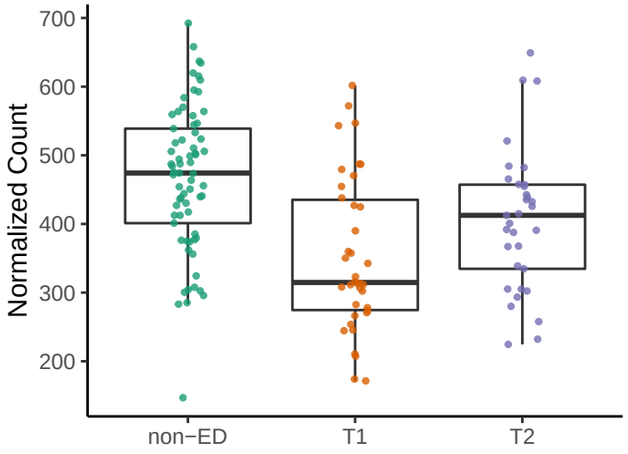
PWY−7242: D−fructuronate degradatio

non−ED vs. T1 adjusted p = 2.9e−05
non−ED vs. T2 adjusted p = 0.27
T1 vs. T2 adjusted p = 0.13



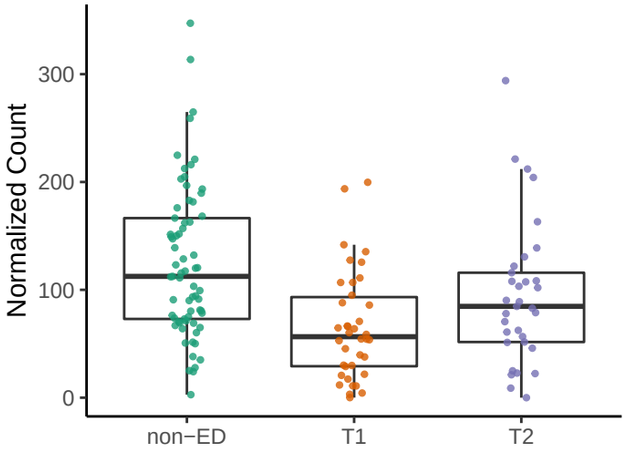
CALVIN−PWY: Calvin−Benson−Bassha

non−ED vs. T1 adjusted p = 1e−04
non−ED vs. T2 adjusted p = 0.07
T1 vs. T2 adjusted p = 0.14



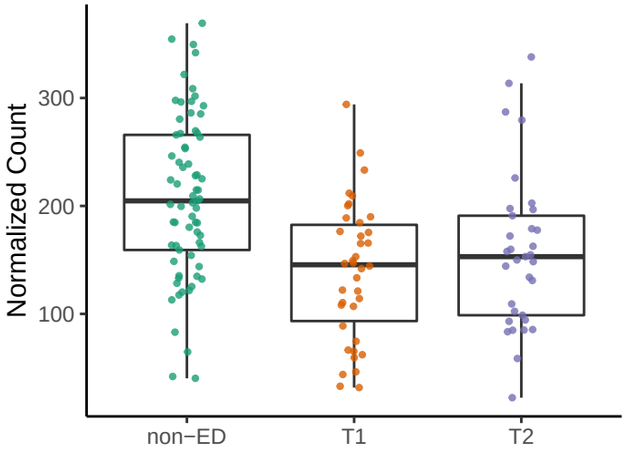
PWY−5177: glutaryl−CoA degradation

non−ED vs. T1 adjusted p = 1e−04
non−ED vs. T2 adjusted p = 0.13
T1 vs. T2 adjusted p = 0.14



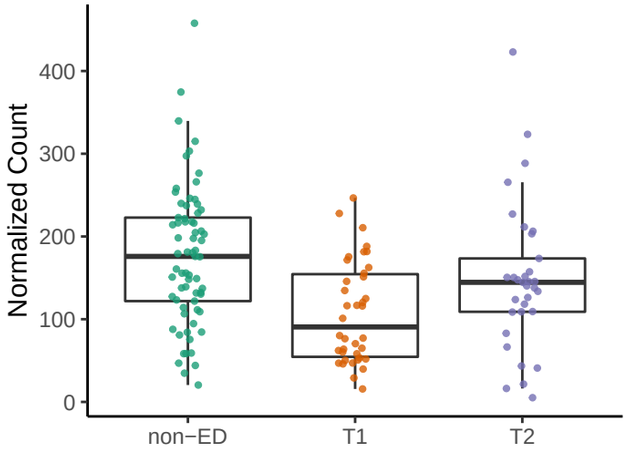
PWY−5347: superpathway of L−methio

non−ED vs. T1 adjusted p = 1e−04
non−ED vs. T2 adjusted p = 0.05
T1 vs. T2 adjusted p = 0.35



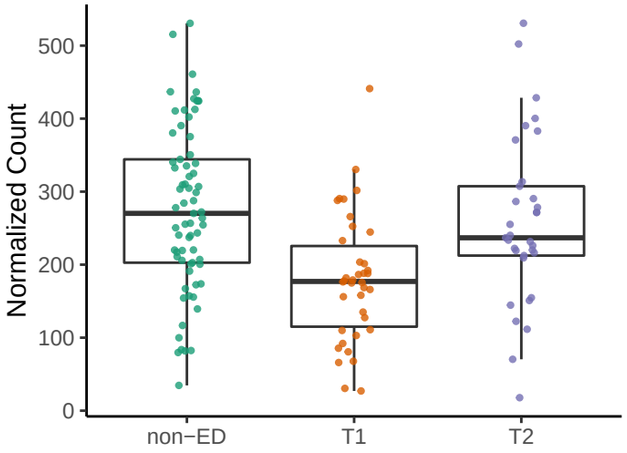
GLUCUROCAT−PWY: superpathway o

non−ED vs. T1 adjusted p = 0.00015
non−ED vs. T2 adjusted p = 0.27
T1 vs. T2 adjusted p = 0.16



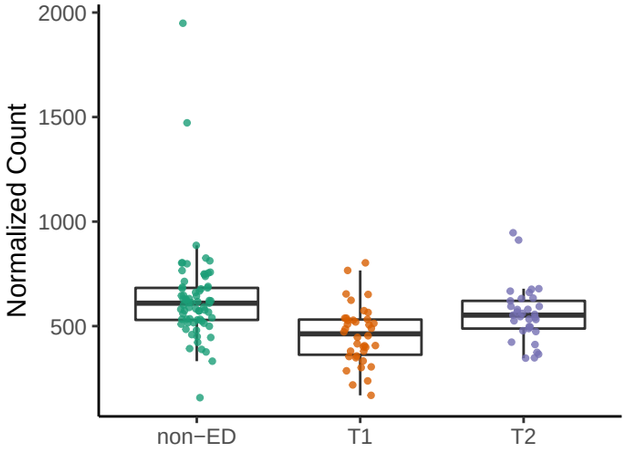
SER−GLYSYN−PWY: superpathway of

non−ED vs. T1 adjusted p = 0.00015
non−ED vs. T2 adjusted p = 0.61
T1 vs. T2 adjusted p = 0.13



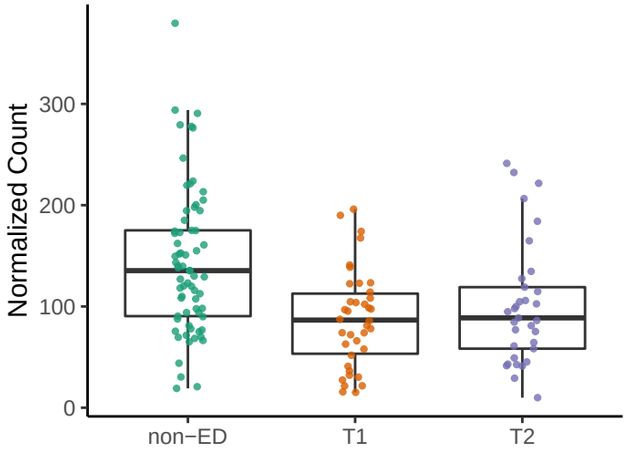
PWY−6151: S−adenosyl−L−methionin

non−ED vs. T1 adjusted p = 0.00015
non−ED vs. T2 adjusted p = 0.14
T1 vs. T2 adjusted p = 0.056



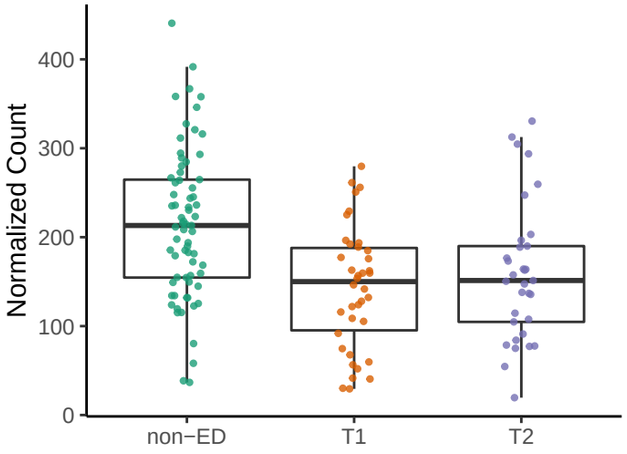
HOMOSER−METSYN−PWY: L−methio

non−ED vs. T1 adjusted p = 0.00016
non−ED vs. T2 adjusted p = 0.05
T1 vs. T2 adjusted p = 0.34



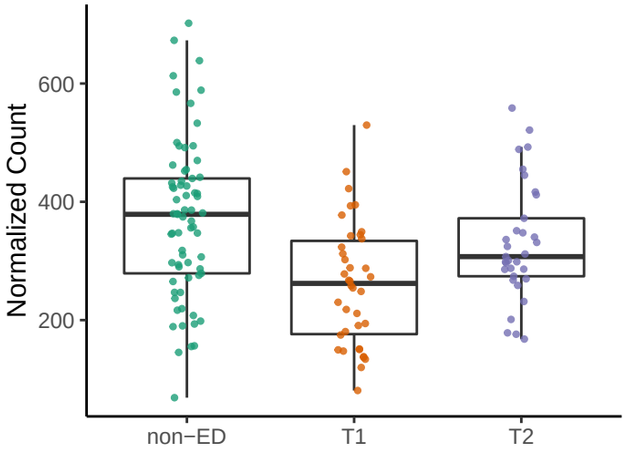
METSYN−PWY: L−homoserine and L−

non−ED vs. T1 adjusted p = 2e−04
non−ED vs. T2 adjusted p = 0.05
T1 vs. T2 adjusted p = 0.39



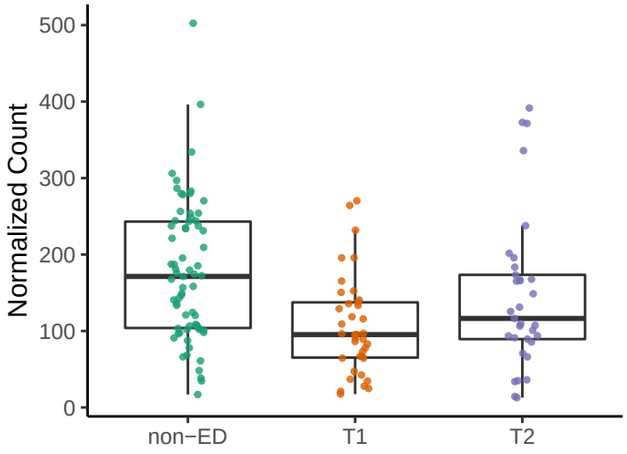
PWY−7357: thiamin formation from pyri

non−ED vs. T1 adjusted p = 2e−04
non−ED vs. T2 adjusted p = 0.2
T1 vs. T2 adjusted p = 0.13



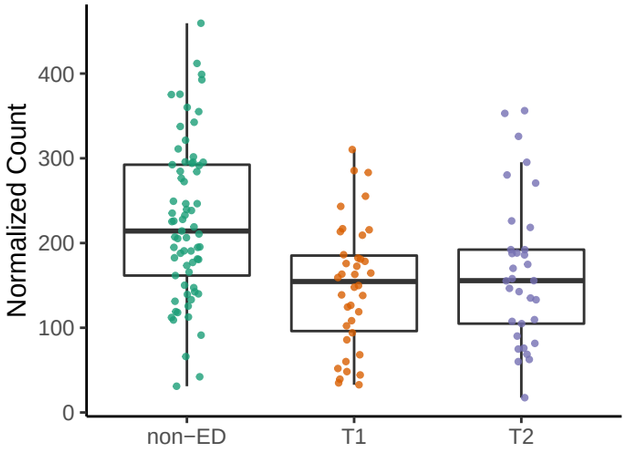
PWY−6507: 4−deoxy−L−threo−hex−4−

non−ED vs. T1 adjusted p = 0.00021
non−ED vs. T2 adjusted p = 0.26
T1 vs. T2 adjusted p = 0.19



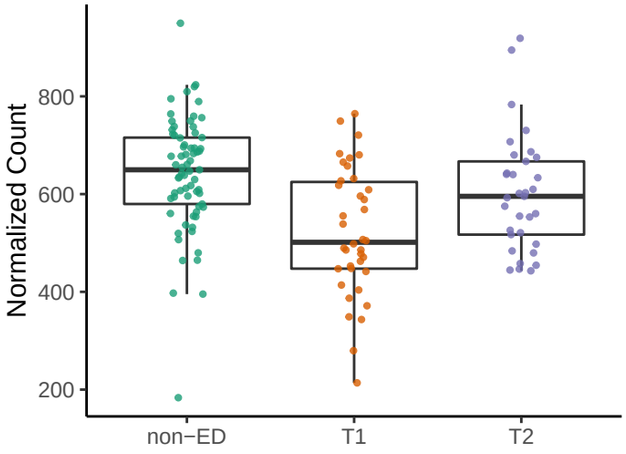
MET−SAM−PWY: superpathway of S−a

non−ED vs. T1 adjusted p = 0.00024
non−ED vs. T2 adjusted p = 0.05
T1 vs. T2 adjusted p = 0.43



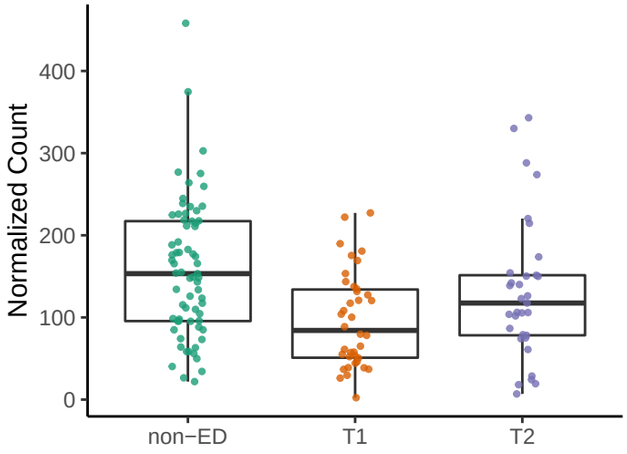
ARO−PWY: chorismate biosynthesis I

non−ED vs. T1 adjusted p = 0.00026
non−ED vs. T2 adjusted p = 0.22
T1 vs. T2 adjusted p = 0.13

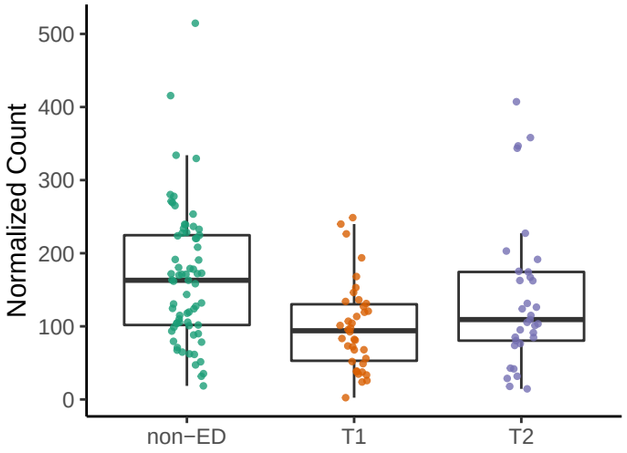


GALACT−GLUCUROCAT−PWY: super

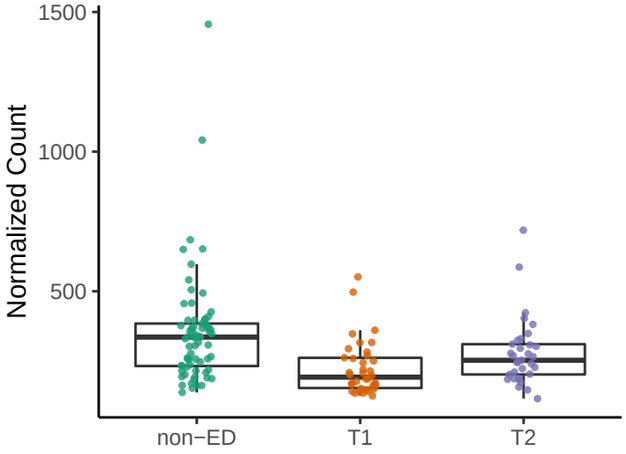
non−ED vs. T1 adjusted p = 0.00026
non−ED vs. T2 adjusted p = 0.26
T1 vs. T2 adjusted p = 0.21



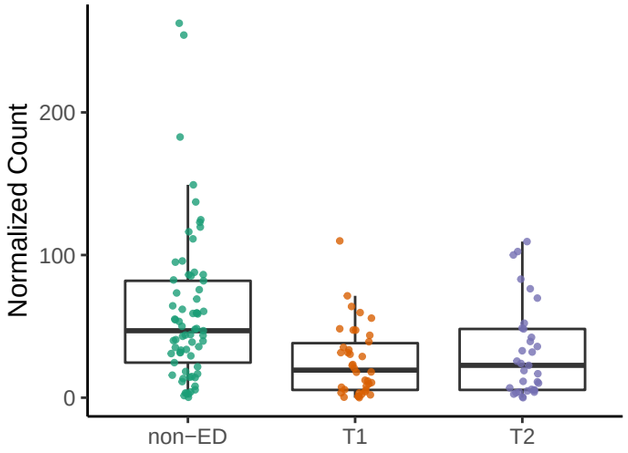
GALACTUROCAT−PWY: D−galacturon
non−ED vs. T1 adjusted p = 0.00026
non−ED vs. T2 adjusted p = 0.34
T1 vs. T2 adjusted p = 0.2



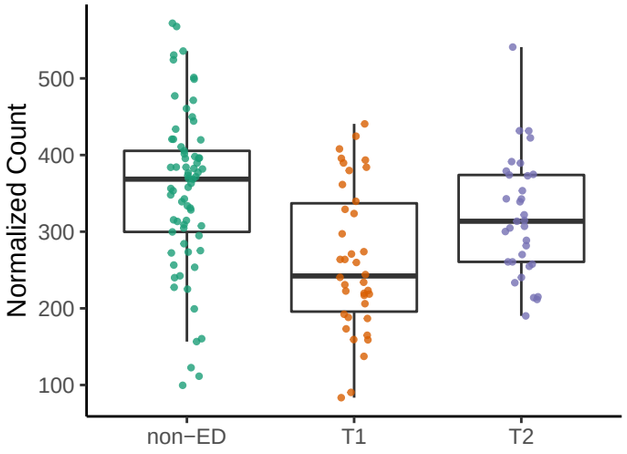
HISTSYN−PWY: L−histidine biosynthe
non−ED vs. T1 adjusted p = 0.00026
non−ED vs. T2 adjusted p = 0.12
T1 vs. T2 adjusted p = 0.13



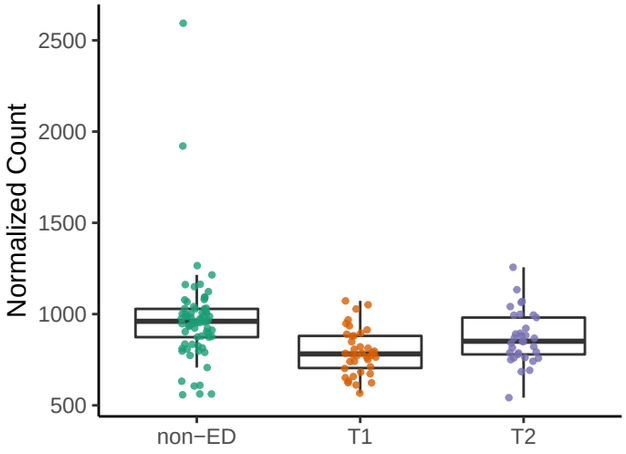
LACTOSECAT−PWY: lactose and galac
non−ED vs. T1 adjusted p = 0.00026
non−ED vs. T2 adjusted p = 0.05
T1 vs. T2 adjusted p = 0.35



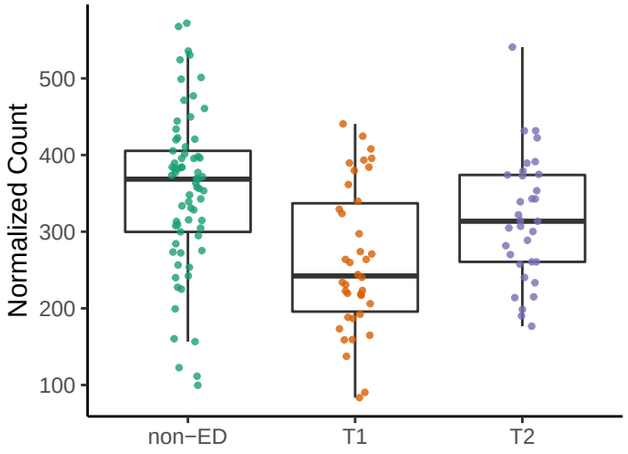
PWY−6317: galactose degradation I (L
non−ED vs. T1 adjusted p = 0.00026
non−ED vs. T2 adjusted p = 0.16
T1 vs. T2 adjusted p = 0.13



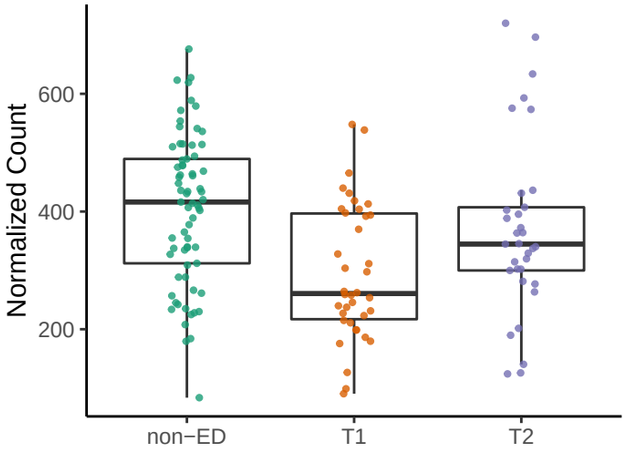
PWY−7219: adenosine ribonucleotides
non−ED vs. T1 adjusted p = 0.00026
non−ED vs. T2 adjusted p = 0.11
T1 vs. T2 adjusted p = 0.16



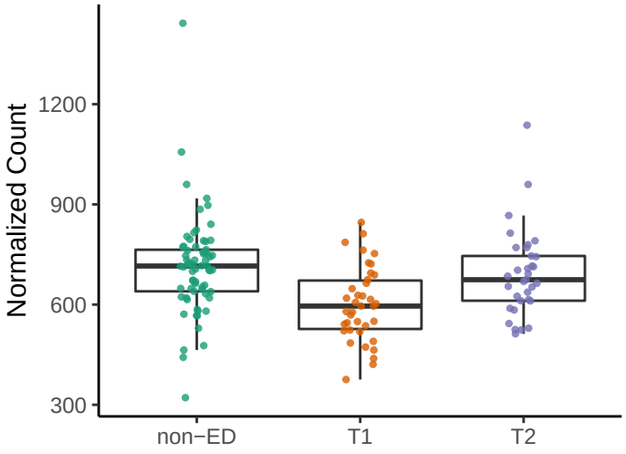
PWY66−422: D−galactose degradation
non−ED vs. T1 adjusted p = 0.00026
non−ED vs. T2 adjusted p = 0.15
T1 vs. T2 adjusted p = 0.13



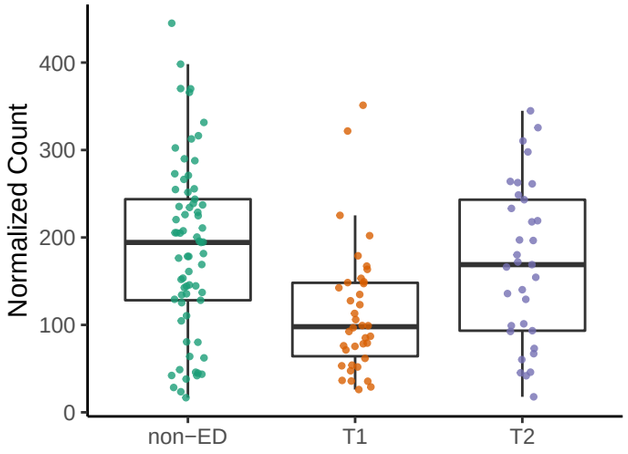
NONOXIPENT−PWY: pentose phospha
non−ED vs. T1 adjusted p = 3e−04
non−ED vs. T2 adjusted p = 0.4
T1 vs. T2 adjusted p = 0.14



PWY−5686: UMP biosynthesis
non−ED vs. T1 adjusted p = 0.00032
non−ED vs. T2 adjusted p = 0.61
T1 vs. T2 adjusted p = 0.13

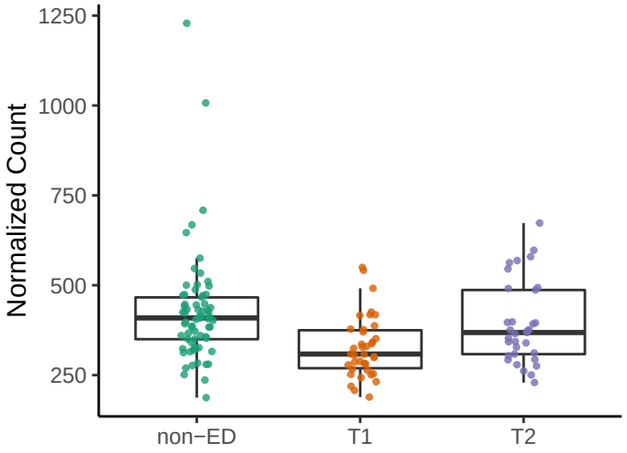


GLYCOGENSYNTH−PWY: glycogen bi
non−ED vs. T1 adjusted p = 0.00048
non−ED vs. T2 adjusted p = 0.55
T1 vs. T2 adjusted p = 0.13



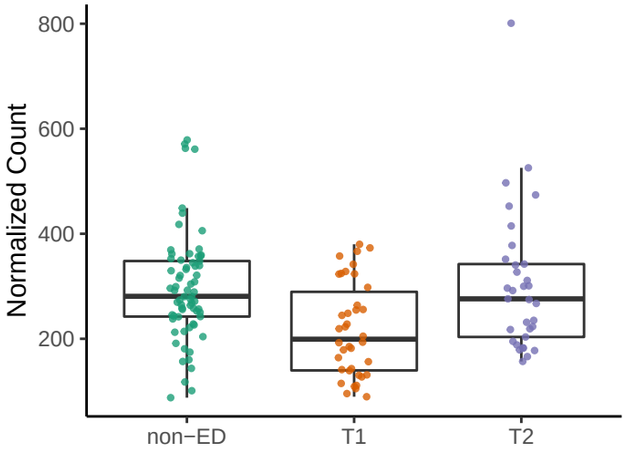
PWY−3001: superpathway of L−isoleu

non−ED vs. T1 adjusted p = 0.00049
non−ED vs. T2 adjusted p = 0.36
T1 vs. T2 adjusted p = 0.13



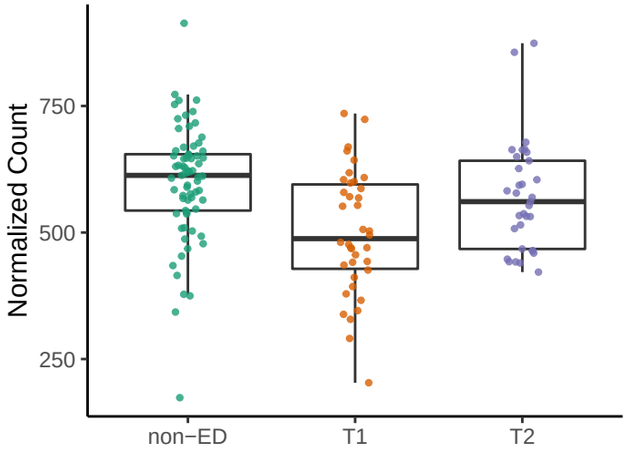
PWY−6123: inosine−5'−phosphate bios

non−ED vs. T1 adjusted p = 0.00082
non−ED vs. T2 adjusted p = 0.89
T1 vs. T2 adjusted p = 0.13



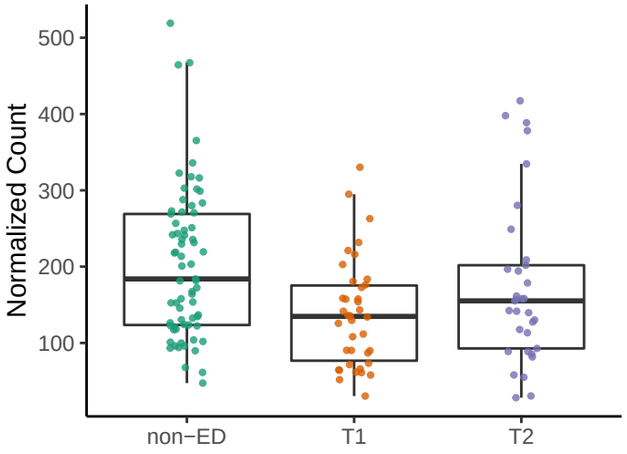
COMPLETE−ARO−PWY: superpathwa

non−ED vs. T1 adjusted p = 0.0013
non−ED vs. T2 adjusted p = 0.38
T1 vs. T2 adjusted p = 0.13



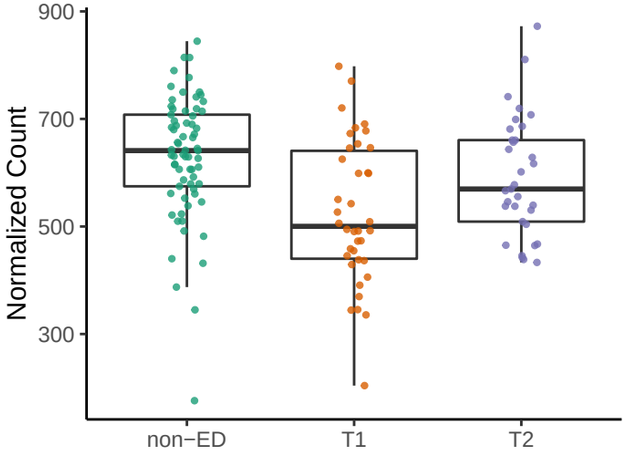
GLCMANNANAUT−PWY: superpathwa

non−ED vs. T1 adjusted p = 0.0014
non−ED vs. T2 adjusted p = 0.33
T1 vs. T2 adjusted p = 0.31



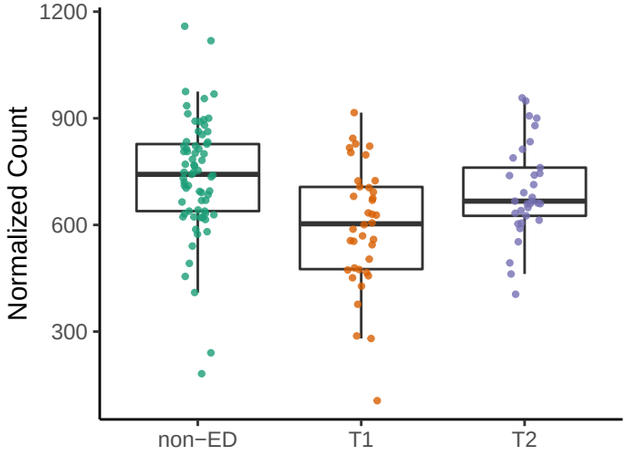
PWY−6163: chorismate biosynthesis fr

non−ED vs. T1 adjusted p = 0.0014
non−ED vs. T2 adjusted p = 0.23
T1 vs. T2 adjusted p = 0.14



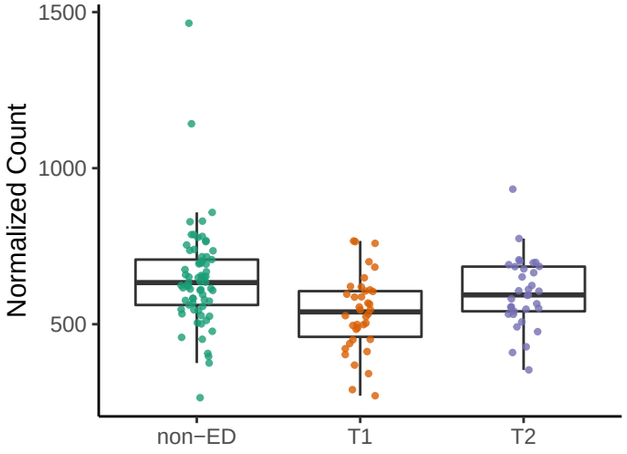
PWY−6737: starch degradation V

non−ED vs. T1 adjusted p = 0.0014
non−ED vs. T2 adjusted p = 0.33
T1 vs. T2 adjusted p = 0.13



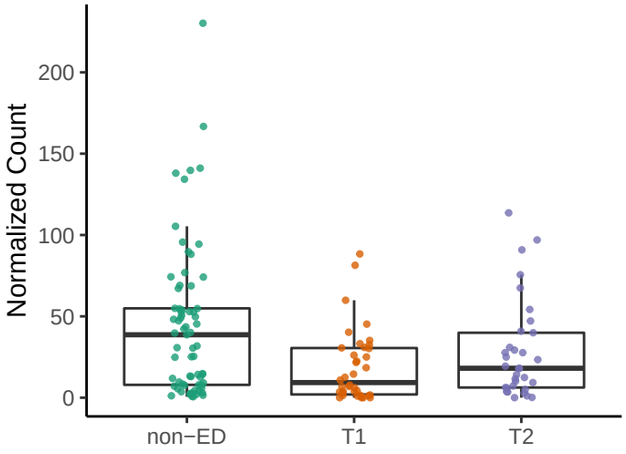
COA−PWY−1: coenzyme A biosynthes

non−ED vs. T1 adjusted p = 0.0014
non−ED vs. T2 adjusted p = 0.24
T1 vs. T2 adjusted p = 0.14



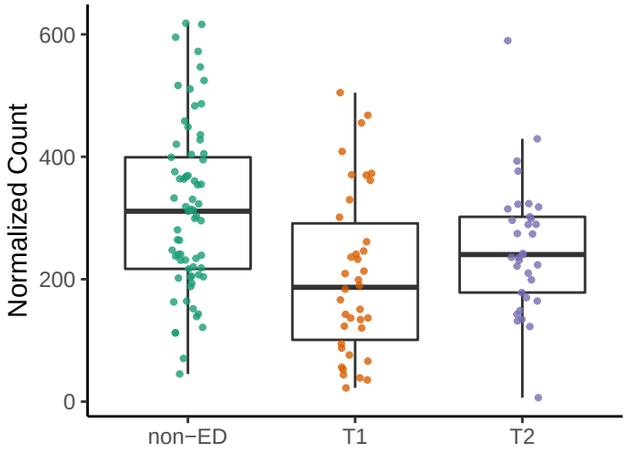
PWY−5304: superpathway of sulfur oxi

non−ED vs. T1 adjusted p = 0.0014
non−ED vs. T2 adjusted p = 0.12
T1 vs. T2 adjusted p = 0.21



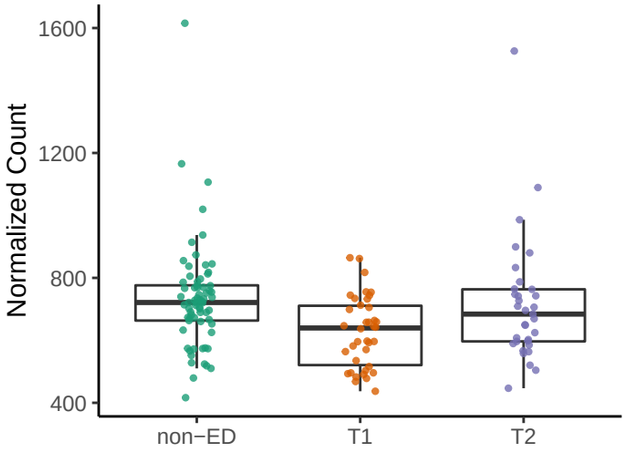
PWY−621: sucrose degradation III (suc

non−ED vs. T1 adjusted p = 0.0015
non−ED vs. T2 adjusted p = 0.091
T1 vs. T2 adjusted p = 0.32



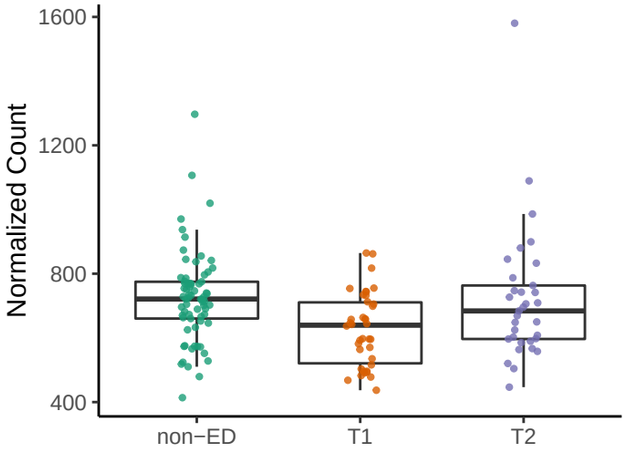
ILEUSYN−PWY: L−isoleucine biosynth

non−ED vs. T1 adjusted p = 0.0017
non−ED vs. T2 adjusted p = 0.78
T1 vs. T2 adjusted p = 0.14



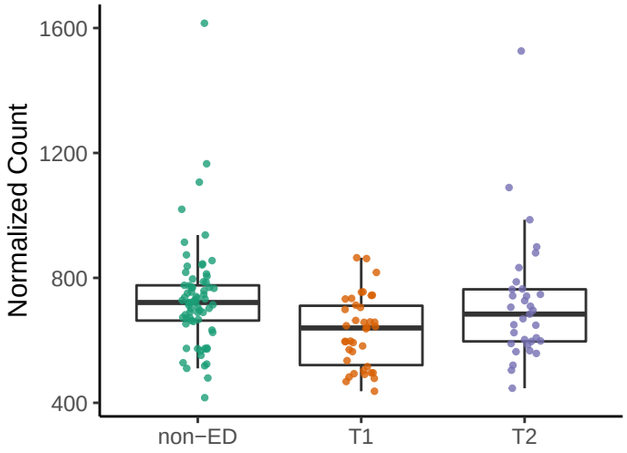
PWY−7111: pyruvate fermentation to i

non−ED vs. T1 adjusted p = 0.0017
non−ED vs. T2 adjusted p = 0.95
T1 vs. T2 adjusted p = 0.14



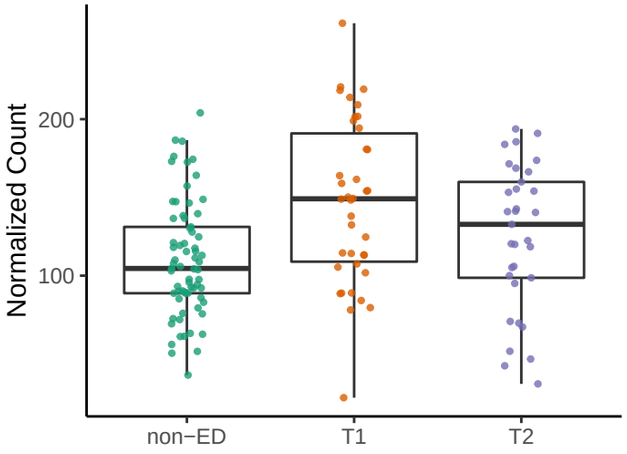
VALSYN−PWY: L−valine biosynthesis

non−ED vs. T1 adjusted p = 0.0017
non−ED vs. T2 adjusted p = 0.78
T1 vs. T2 adjusted p = 0.14



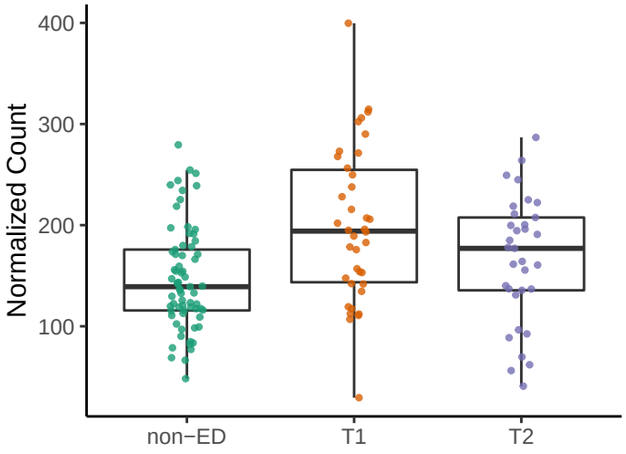
PWY−7184: pyrimidine deoxyribonucle

non−ED vs. T1 adjusted p = 0.0022
non−ED vs. T2 adjusted p = 0.22
T1 vs. T2 adjusted p = 0.18



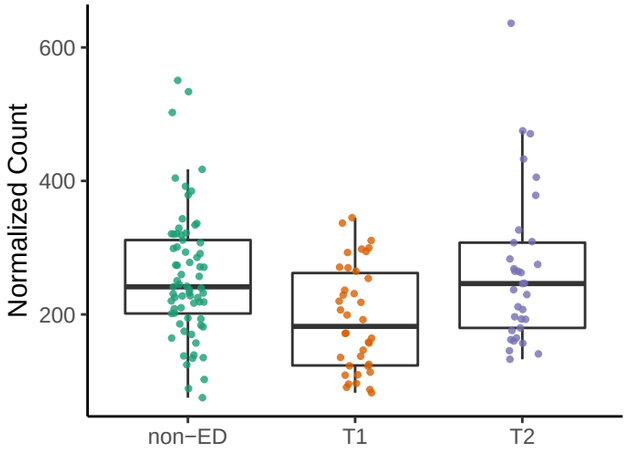
PWY−7228: superpathway of guanosin

non−ED vs. T1 adjusted p = 0.0027
non−ED vs. T2 adjusted p = 0.26
T1 vs. T2 adjusted p = 0.15



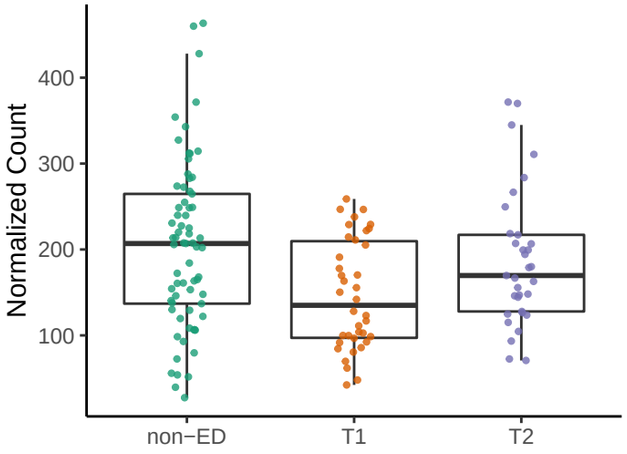
PWY−6124: inosine−5'−phosphate bios

non−ED vs. T1 adjusted p = 0.0027
non−ED vs. T2 adjusted p = 0.85
T1 vs. T2 adjusted p = 0.13



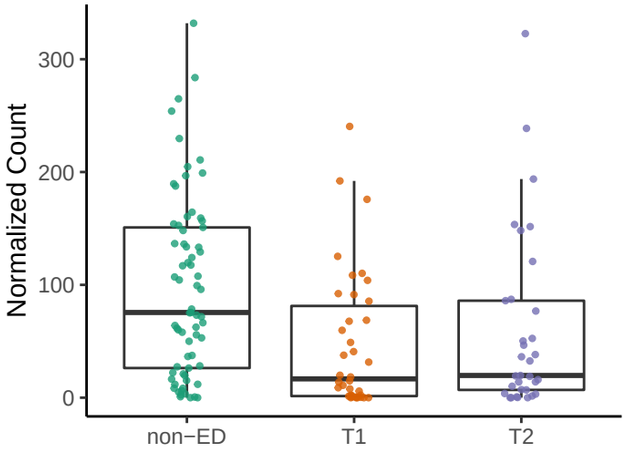
THISYNARA−PWY: superpathway of th

non−ED vs. T1 adjusted p = 0.003
non−ED vs. T2 adjusted p = 0.52
T1 vs. T2 adjusted p = 0.14



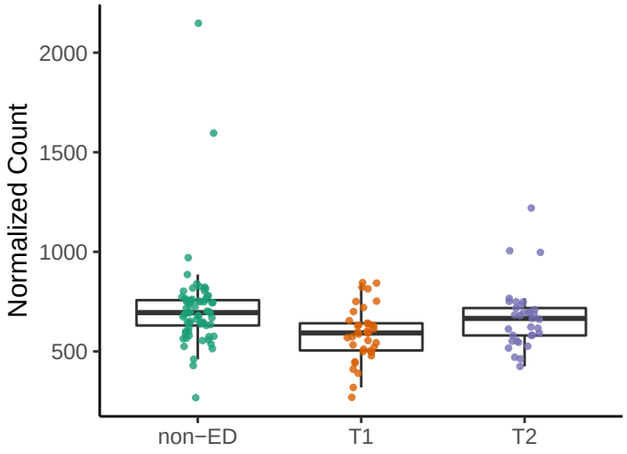
PWY−5367: petroselinate biosynthesis

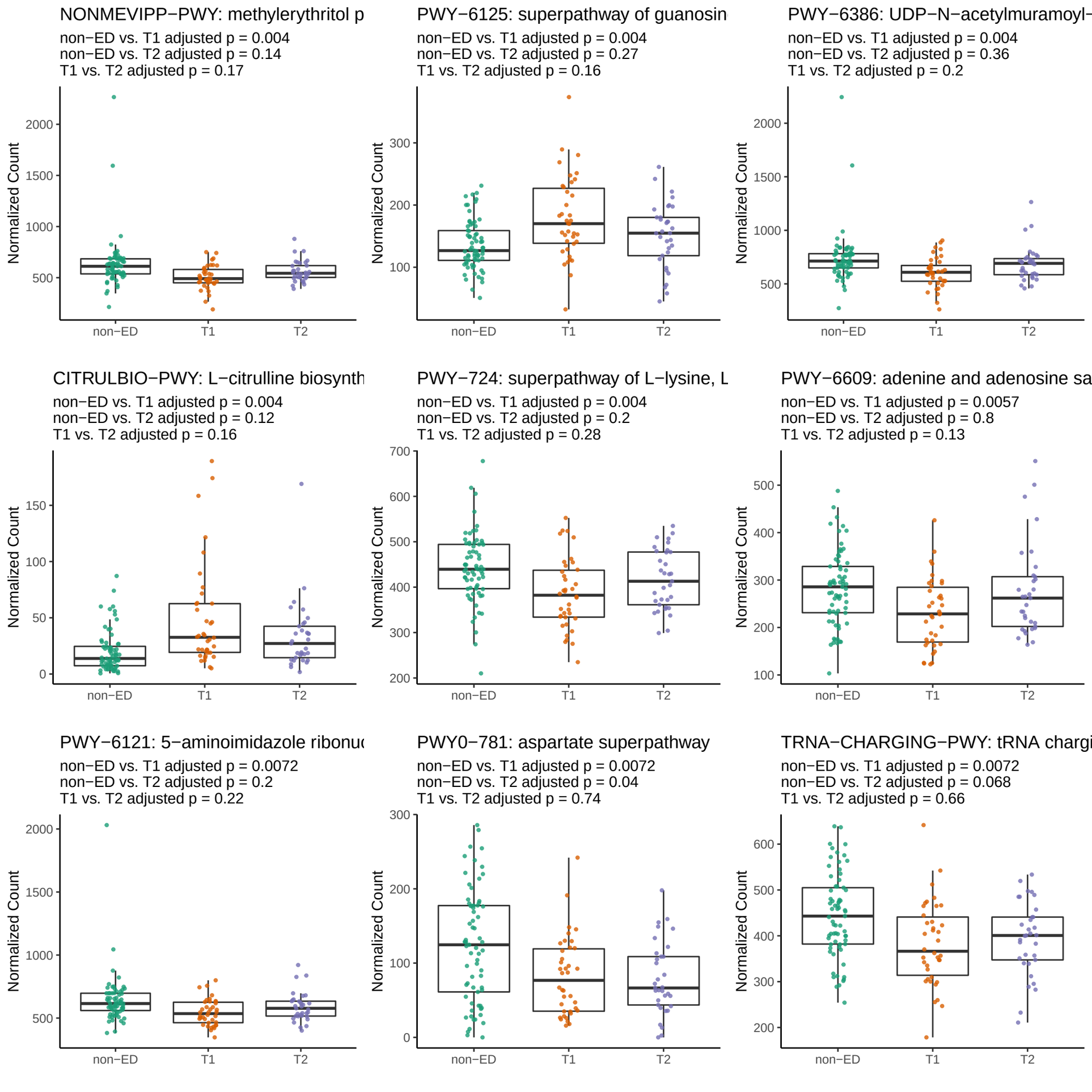
non−ED vs. T1 adjusted p = 0.0035
non−ED vs. T2 adjusted p = 0.12
T1 vs. T2 adjusted p = 0.68



PWY−6387: UDP−N−acetylmuramoyl−

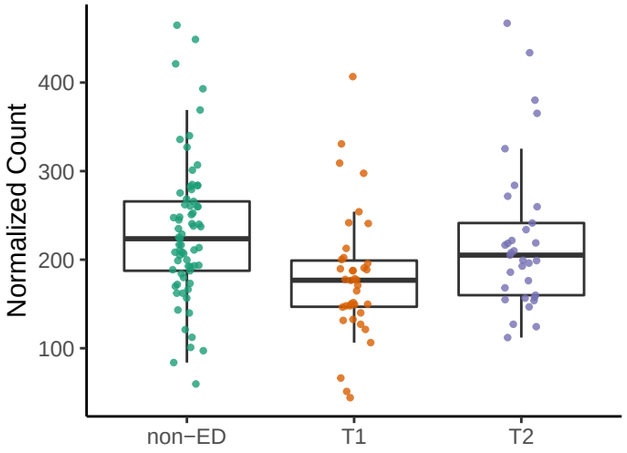
non−ED vs. T1 adjusted p = 0.0035
non−ED vs. T2 adjusted p = 0.37
T1 vs. T2 adjusted p = 0.19




NONMEVIPP−PWY: methylerythritol p
non−ED vs. T1 adjusted p = 0.004
non−ED vs. T2 adjusted p = 0.14
T1 vs. T2 adjusted p = 0.17
PWY−6125: superpathway of guanosin
non−ED vs. T1 adjusted p = 0.004
non−ED vs. T2 adjusted p = 0.27
T1 vs. T2 adjusted p = 0.16
PWY−6386: UDP−N−acetylmuramoyl−
non−ED vs. T1 adjusted p = 0.004
non−ED vs. T2 adjusted p = 0.36
T1 vs. T2 adjusted p = 0.2
CITRULBIO−PWY: L−citrulline biosynth
non−ED vs. T1 adjusted p = 0.004
non−ED vs. T2 adjusted p = 0.12
T1 vs. T2 adjusted p = 0.16
PWY−724: superpathway of L−lysine, L
non−ED vs. T1 adjusted p = 0.004
non−ED vs. T2 adjusted p = 0.2
T1 vs. T2 adjusted p = 0.28
PWY−6609: adenine and adenosine sa
non−ED vs. T1 adjusted p = 0.0057
non−ED vs. T2 adjusted p = 0.8
T1 vs. T2 adjusted p = 0.13
PWY−6121: 5−aminoimidazole ribonuc
non−ED vs. T1 adjusted p = 0.0072
non−ED vs. T2 adjusted p = 0.2
T1 vs. T2 adjusted p = 0.22
PWY0−781: aspartate superpathway
non−ED vs. T1 adjusted p = 0.0072
non−ED vs. T2 adjusted p = 0.04
T1 vs. T2 adjusted p = 0.74
TRNA−CHARGING−PWY: tRNA chargi
non−ED vs. T1 adjusted p = 0.0072
non−ED vs. T2 adjusted p = 0.068
T1 vs. T2 adjusted p = 0.66
Normalized Count
non−ED
T1
T2

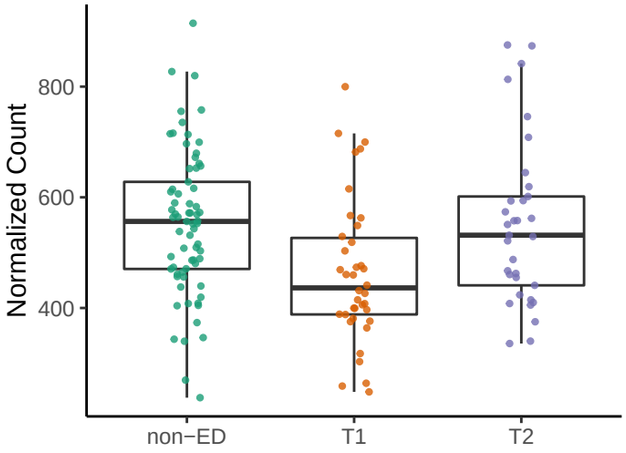
PWY−5100: pyruvate fermentation to a

non−ED vs. T1 adjusted p = 0.0074
non−ED vs. T2 adjusted p = 0.79
T1 vs. T2 adjusted p = 0.14



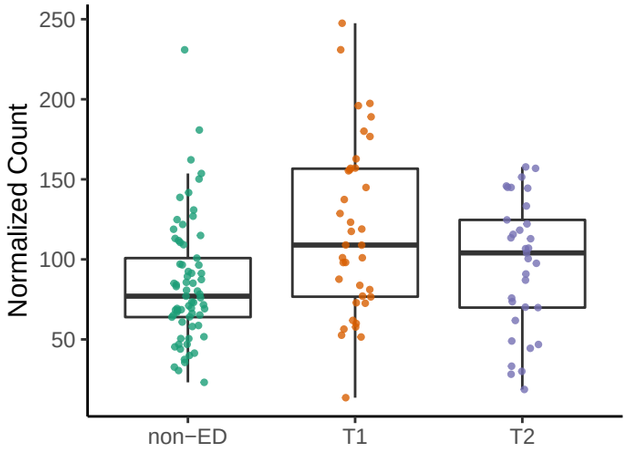
PWY−1042: glycolysis IV (plant cytosol

non−ED vs. T1 adjusted p = 0.0074
non−ED vs. T2 adjusted p = 0.98
T1 vs. T2 adjusted p = 0.13



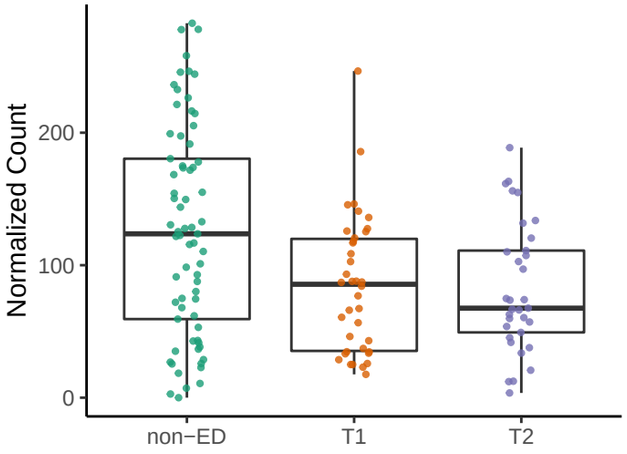
PWY−7197: pyrimidine deoxyribonucleo

non−ED vs. T1 adjusted p = 0.0074
non−ED vs. T2 adjusted p = 0.31
T1 vs. T2 adjusted p = 0.18



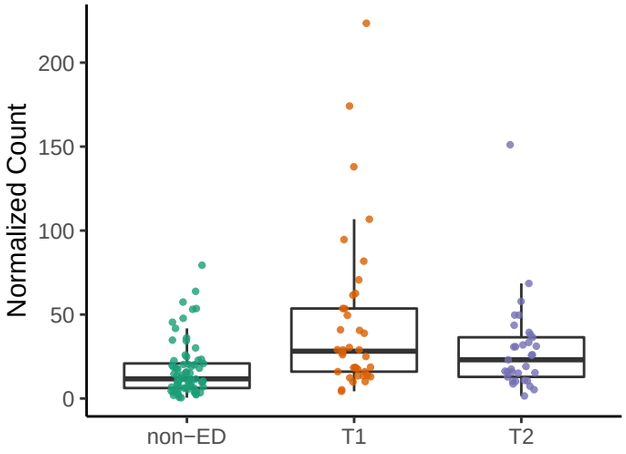
P4−PWY: superpathway of L−lysine, L-

non−ED vs. T1 adjusted p = 0.0074
non−ED vs. T2 adjusted p = 0.044
T1 vs. T2 adjusted p = 0.74



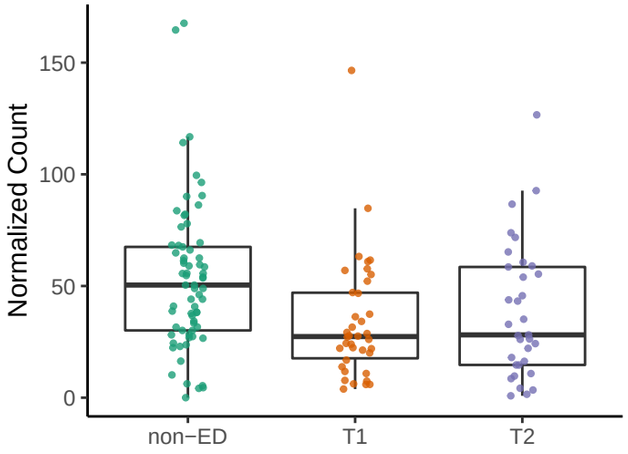
PWY−4984: urea cycle

non−ED vs. T1 adjusted p = 0.0081
non−ED vs. T2 adjusted p = 0.12
T1 vs. T2 adjusted p = 0.17



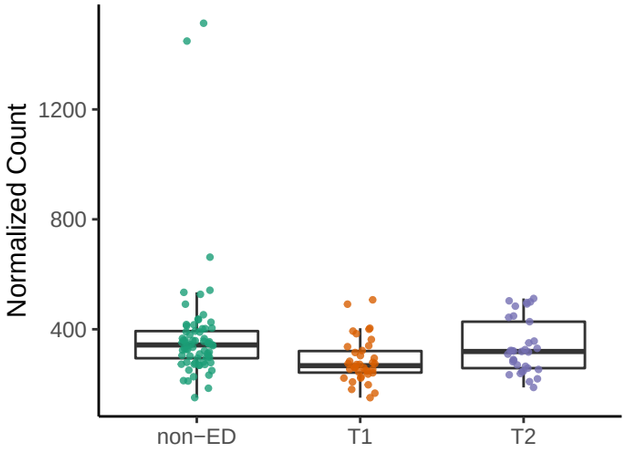
POLYAMSYN−PWY: superpathway of p

non−ED vs. T1 adjusted p = 0.0091
non−ED vs. T2 adjusted p = 0.12
T1 vs. T2 adjusted p = 0.64



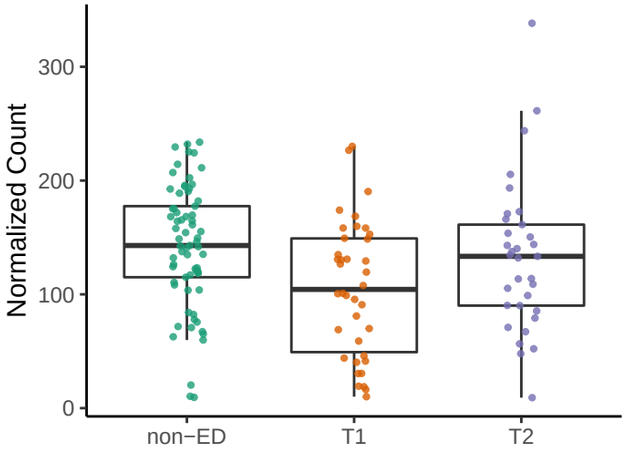
THRESYN−PWY: superpathway of L−t

non−ED vs. T1 adjusted p = 0.0097
non−ED vs. T2 adjusted p = 0.31
T1 vs. T2 adjusted p = 0.15



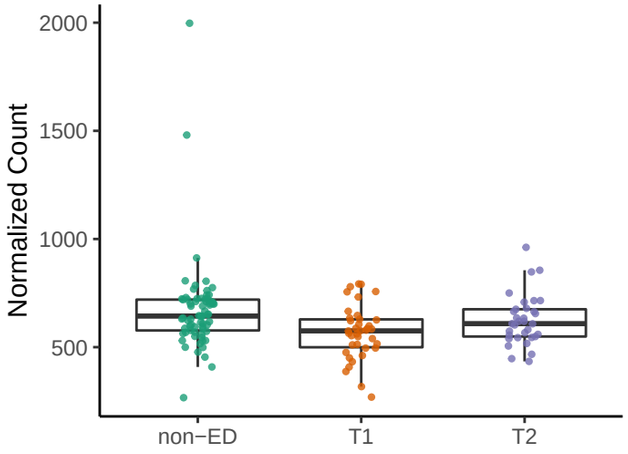
PWY−7199: pyrimidine deoxyribonucleo

non−ED vs. T1 adjusted p = 0.011
non−ED vs. T2 adjusted p = 0.62
T1 vs. T2 adjusted p = 0.23



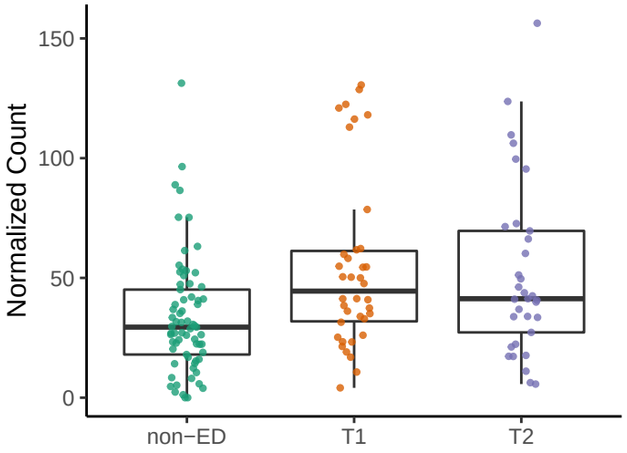
PEPTIDOGLYCANSYN−PWY: peptido

non−ED vs. T1 adjusted p = 0.012
non−ED vs. T2 adjusted p = 0.28
T1 vs. T2 adjusted p = 0.24



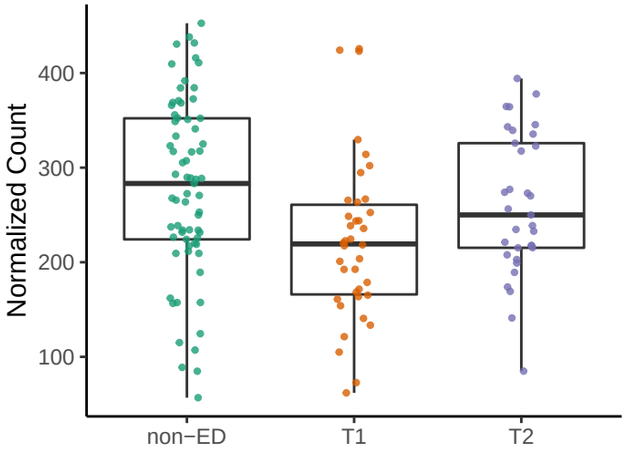
PWY66−399: gluconeogenesis III

non−ED vs. T1 adjusted p = 0.012
non−ED vs. T2 adjusted p = 0.086
T1 vs. T2 adjusted p = 0.81



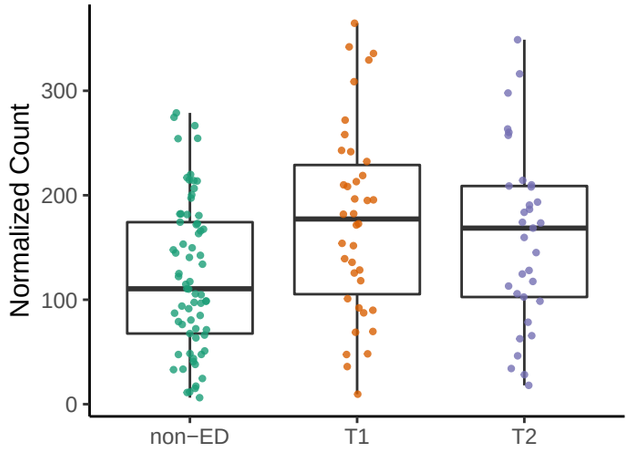
PWY−6527: stachyose degradation

non−ED vs. T1 adjusted p = 0.013
non−ED vs. T2 adjusted p = 0.44
T1 vs. T2 adjusted p = 0.23



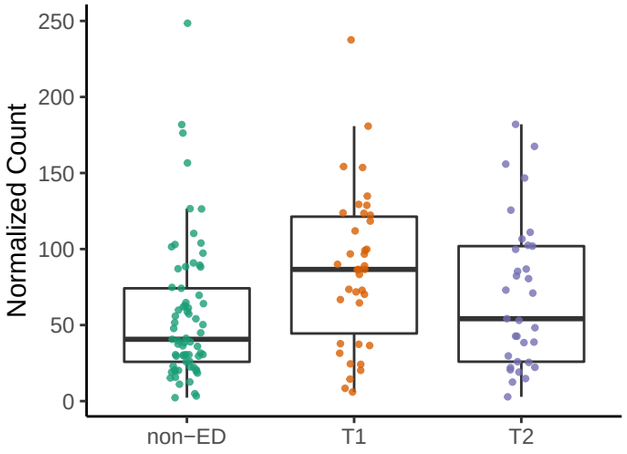
PWY−7663: gondoate biosynthesis (an

non−ED vs. T1 adjusted p = 0.015
non−ED vs. T2 adjusted p = 0.12
T1 vs. T2 adjusted p = 0.34



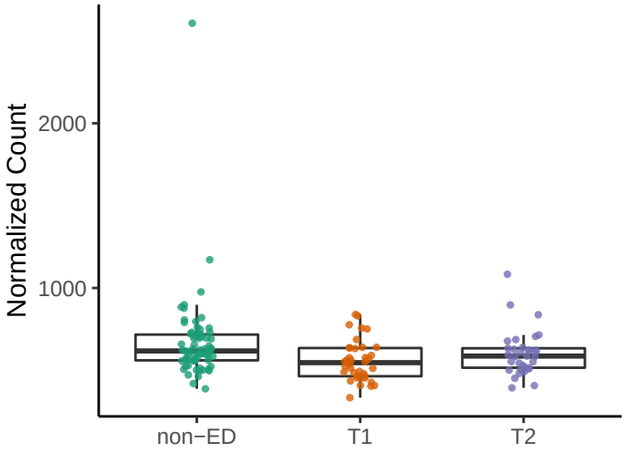
PWY−5154: L−arginine biosynthesis III

non−ED vs. T1 adjusted p = 0.016
non−ED vs. T2 adjusted p = 0.34
T1 vs. T2 adjusted p = 0.21



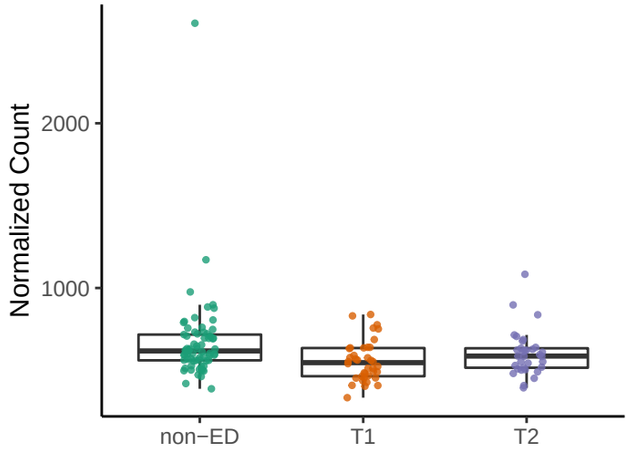
PWY−6122: 5−aminoimidazole ribonuc

non−ED vs. T1 adjusted p = 0.016
non−ED vs. T2 adjusted p = 0.21
T1 vs. T2 adjusted p = 0.24



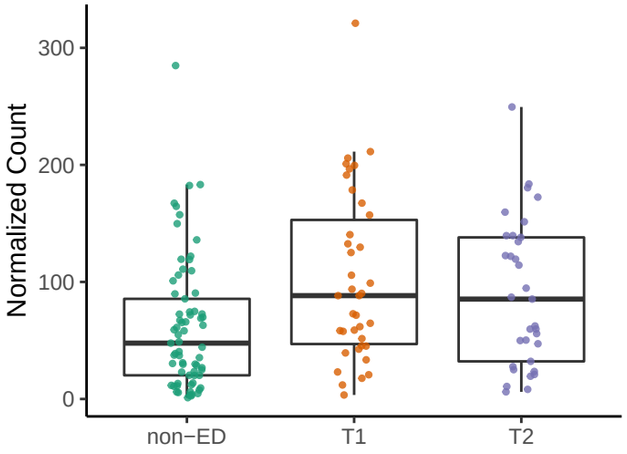
PWY−6277: superpathway of 5−amino

non−ED vs. T1 adjusted p = 0.016
non−ED vs. T2 adjusted p = 0.21
T1 vs. T2 adjusted p = 0.24



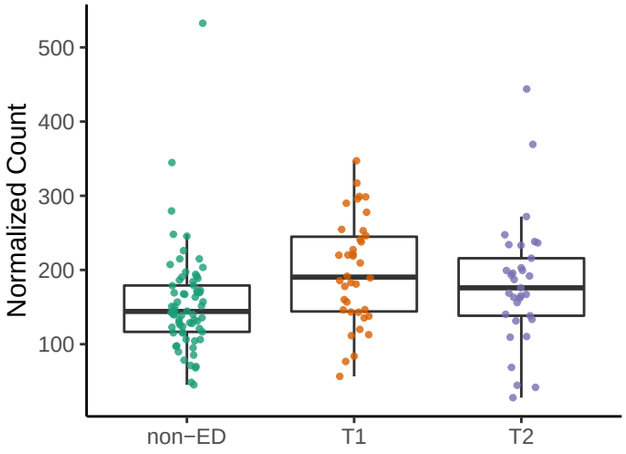
PYRIDOXSYN−PWY: pyridoxal 5'−phos

non−ED vs. T1 adjusted p = 0.016
non−ED vs. T2 adjusted p = 0.12
T1 vs. T2 adjusted p = 0.35



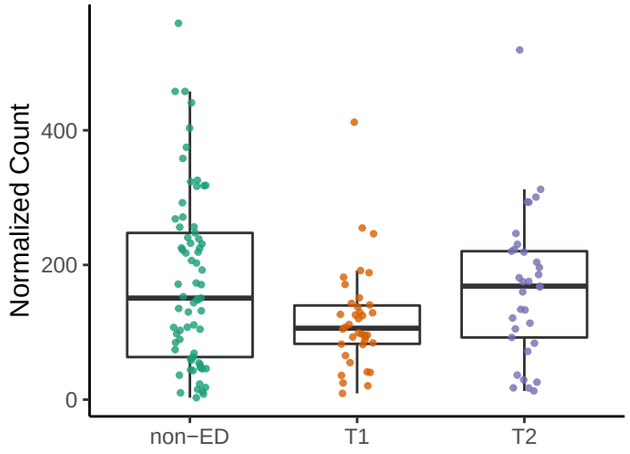
PWY−7208: superpathway of pyrimidin

non−ED vs. T1 adjusted p = 0.018
non−ED vs. T2 adjusted p = 0.26
T1 vs. T2 adjusted p = 0.61



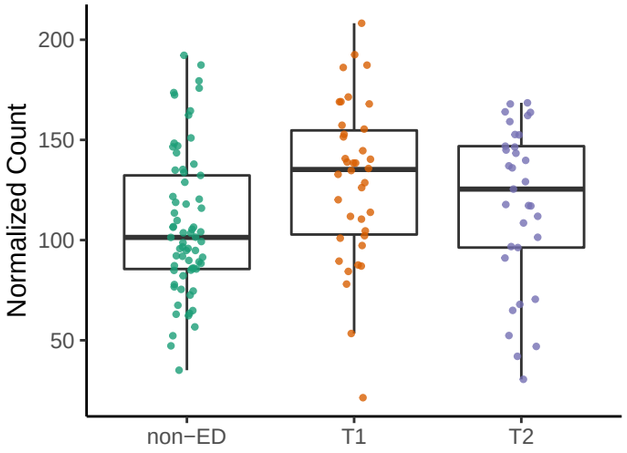
OANTIGEN−PWY: O−antigen building

non−ED vs. T1 adjusted p = 0.018
non−ED vs. T2 adjusted p = 0.78
T1 vs. T2 adjusted p = 0.14



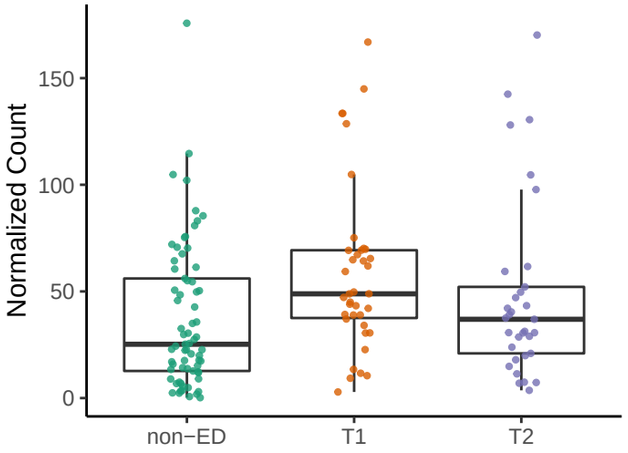
PWY−6545: pyrimidine deoxyribonucle(

non−ED vs. T1 adjusted p = 0.022
non−ED vs. T2 adjusted p = 0.37
T1 vs. T2 adjusted p = 0.27



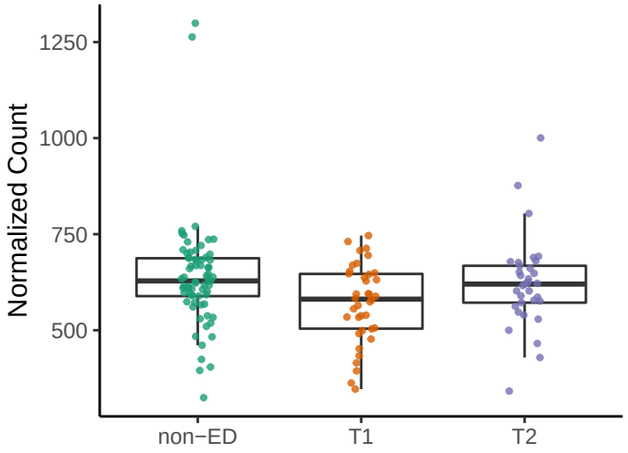
PWY−1269: CMP−3−deoxy−D−manno-

non−ED vs. T1 adjusted p = 0.025
non−ED vs. T2 adjusted p = 0.33
T1 vs. T2 adjusted p = 0.3



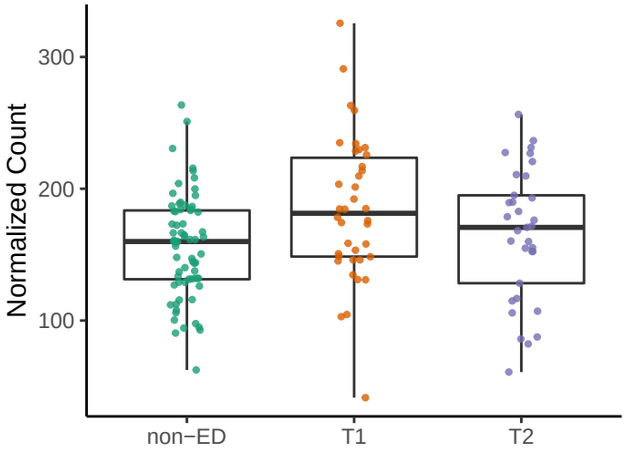
PWY−7221: guanosine ribonucleotides

non−ED vs. T1 adjusted p = 0.025
non−ED vs. T2 adjusted p = 0.67
T1 vs. T2 adjusted p = 0.2



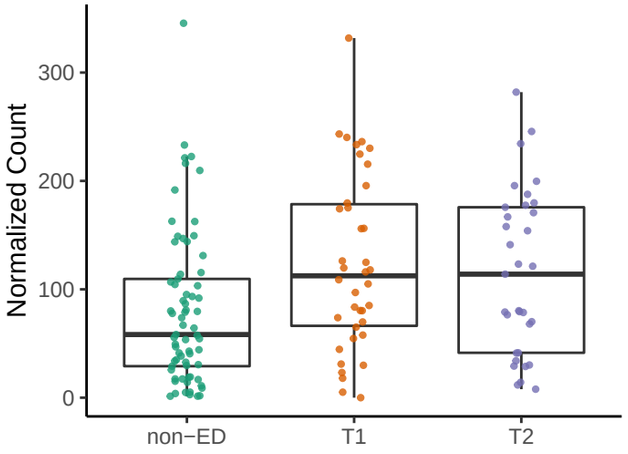
PWY0−166: superpathway of pyrimidin(

non−ED vs. T1 adjusted p = 0.025
non−ED vs. T2 adjusted p = 0.47
T1 vs. T2 adjusted p = 0.27



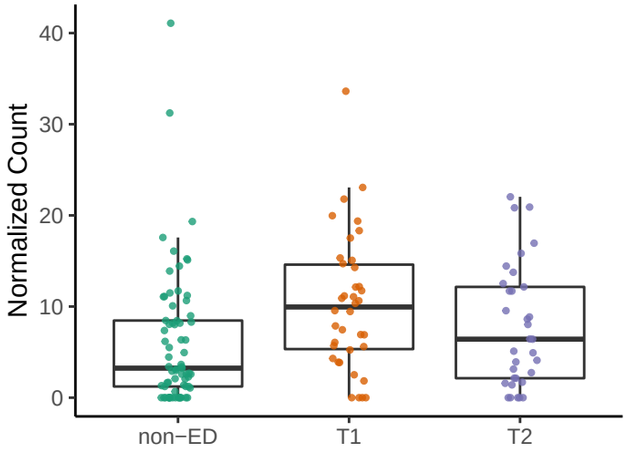
PWY0−845: superpathway of pyridoxal

non−ED vs. T1 adjusted p = 0.026
non−ED vs. T2 adjusted p = 0.12
T1 vs. T2 adjusted p = 0.55



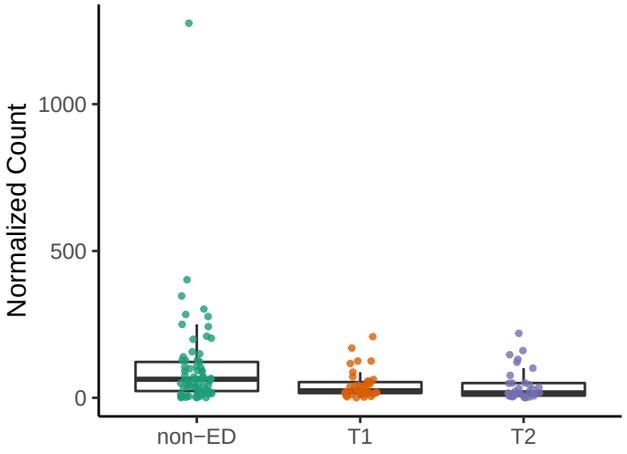
P162−PWY: L−glutamate degradation V

non−ED vs. T1 adjusted p = 0.027
non−ED vs. T2 adjusted p = 0.42
T1 vs. T2 adjusted p = 0.25



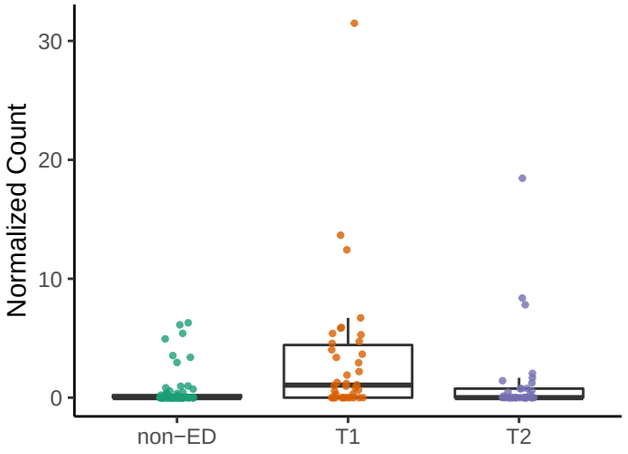
PWY−6147: 6−hydroxymethyl−dihydro

non−ED vs. T1 adjusted p = 0.03
non−ED vs. T2 adjusted p = 0.074
T1 vs. T2 adjusted p = 0.88



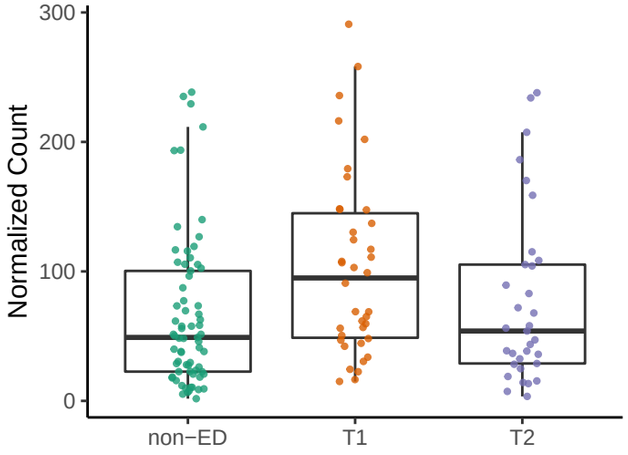
7ALPHADEHYDROX−PWY: cholate deg

non−ED vs. T1 adjusted p = 0.031
non−ED vs. T2 adjusted p = 0.37
T1 vs. T2 adjusted p = 0.26



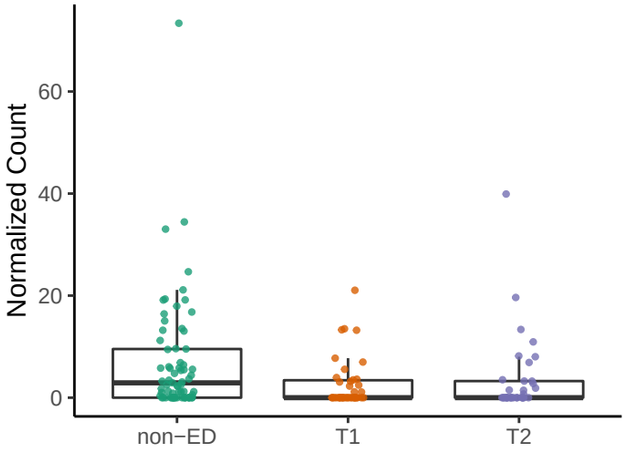
GLYCOLYSIS: glycolysis I (from glucos

non−ED vs. T1 adjusted p = 0.031
non−ED vs. T2 adjusted p = 0.59
T1 vs. T2 adjusted p = 0.22



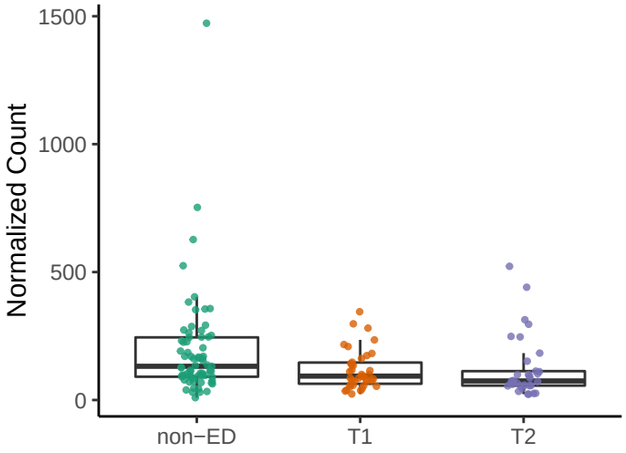
PWY−1861: formaldehyde assimilation I

non−ED vs. T1 adjusted p = 0.031
non−ED vs. T2 adjusted p = 0.21
T1 vs. T2 adjusted p = 0.39



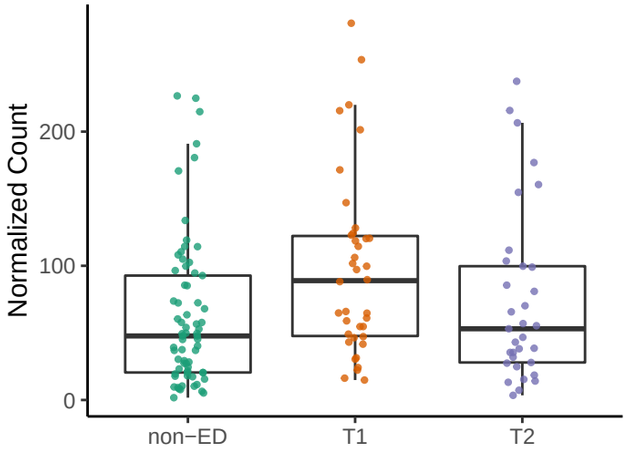
PWY−4981: L−proline biosynthesis II (

non−ED vs. T1 adjusted p = 0.031
non−ED vs. T2 adjusted p = 0.12
T1 vs. T2 adjusted p = 0.84



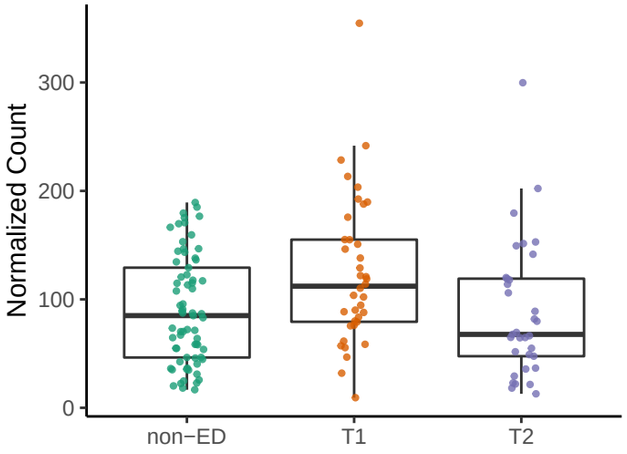
PWY−5484: glycolysis II (from fructose

non−ED vs. T1 adjusted p = 0.031
non−ED vs. T2 adjusted p = 0.52
T1 vs. T2 adjusted p = 0.24



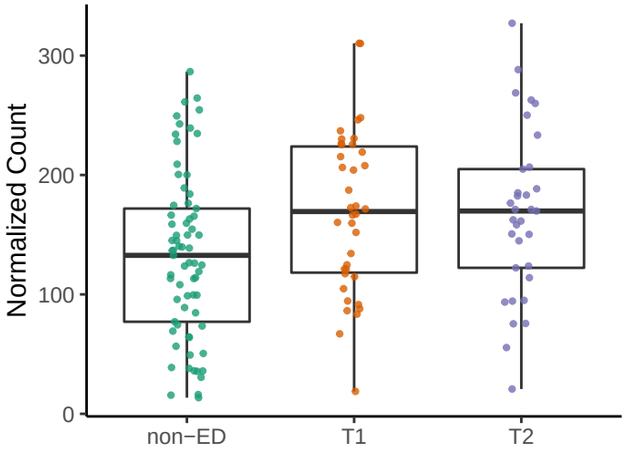
PWY66−409: superpathway of purine n

non−ED vs. T1 adjusted p = 0.031
non−ED vs. T2 adjusted p = 0.93
T1 vs. T2 adjusted p = 0.13



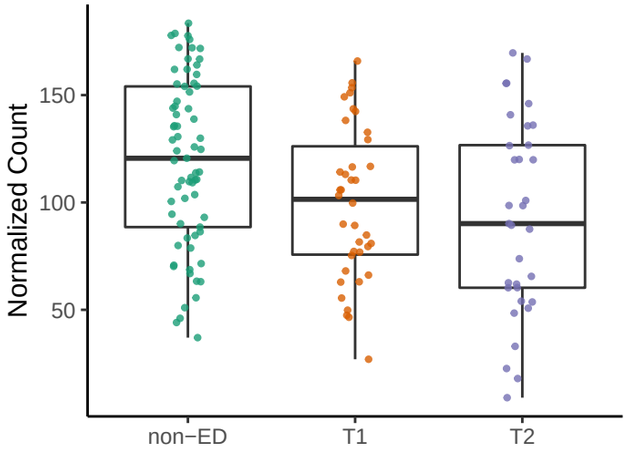
PWY−6168: flavin biosynthesis III (fung

non−ED vs. T1 adjusted p = 0.034
non−ED vs. T2 adjusted p = 0.11
T1 vs. T2 adjusted p = 0.68



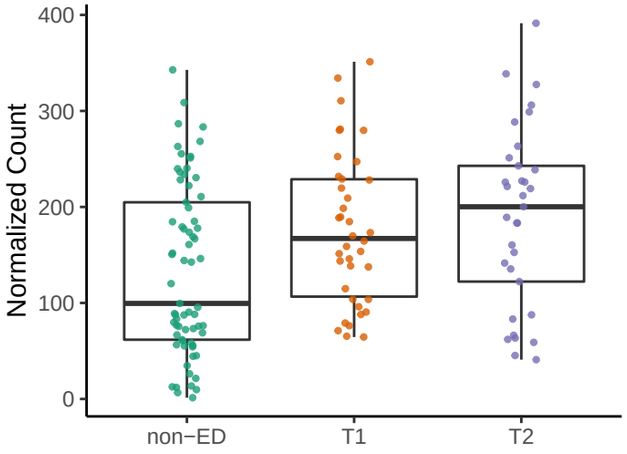
PRPP−PWY: superpathway of histidine

non−ED vs. T1 adjusted p = 0.038
non−ED vs. T2 adjusted p = 0.068
T1 vs. T2 adjusted p = 0.27



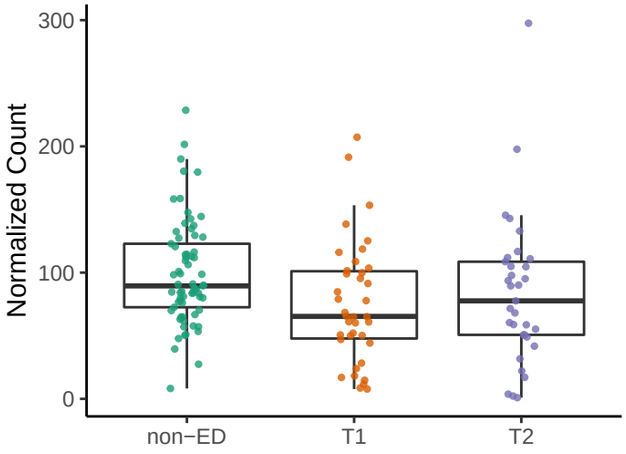
PWY−6897: thiamin salvage II

non−ED vs. T1 adjusted p = 0.041
non−ED vs. T2 adjusted p = 0.068
T1 vs. T2 adjusted p = 0.68



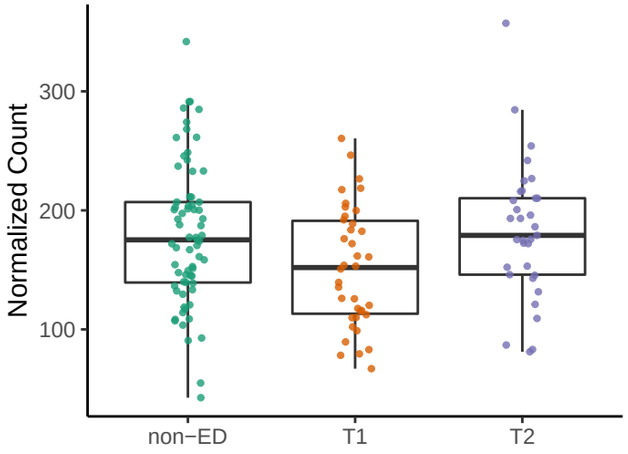
PWY−6305: putrescine biosynthesis IV

non−ED vs. T1 adjusted p = 0.041
non−ED vs. T2 adjusted p = 0.31
T1 vs. T2 adjusted p = 0.39



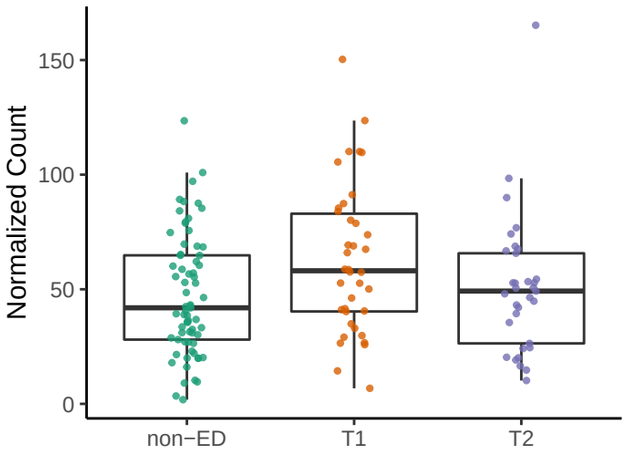
PYRIDNUCSYN−PWY: NAD biosynthe

non−ED vs. T1 adjusted p = 0.044
non−ED vs. T2 adjusted p = 0.89
T1 vs. T2 adjusted p = 0.16



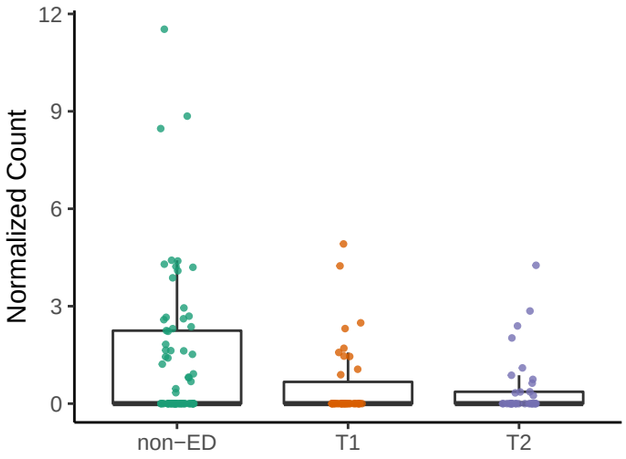
P42−PWY: incomplete reductive TCA c

non−ED vs. T1 adjusted p = 0.05
non−ED vs. T2 adjusted p = 0.73
T1 vs. T2 adjusted p = 0.18



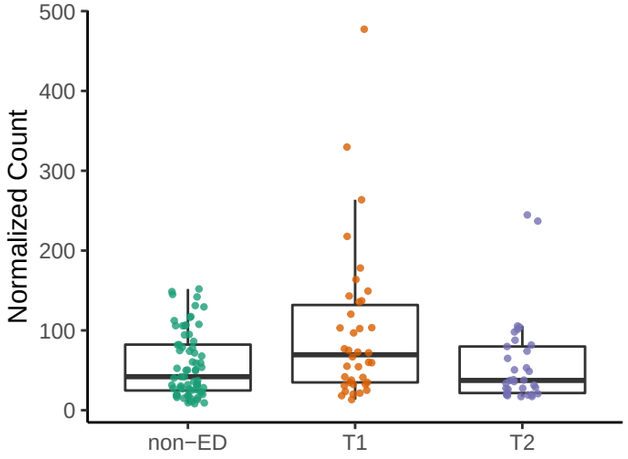
RUMP−PWY: formaldehyde oxidation I

non−ED vs. T1 adjusted p = 0.053
non−ED vs. T2 adjusted p = 0.068
T1 vs. T2 adjusted p = 0.73



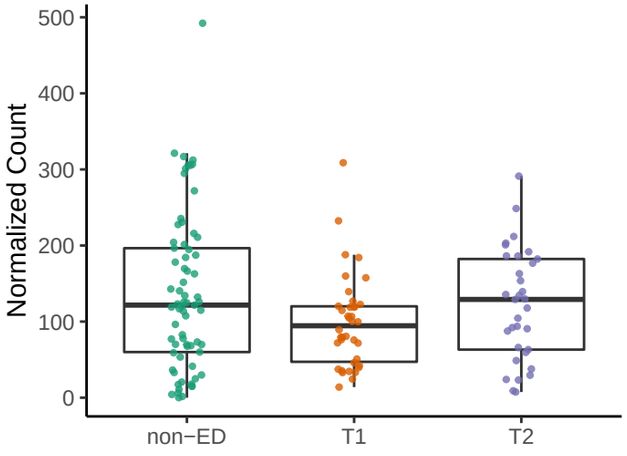
PWY0−1297: superpathway of purine d

non−ED vs. T1 adjusted p = 0.054
non−ED vs. T2 adjusted p = 0.95
T1 vs. T2 adjusted p = 0.056



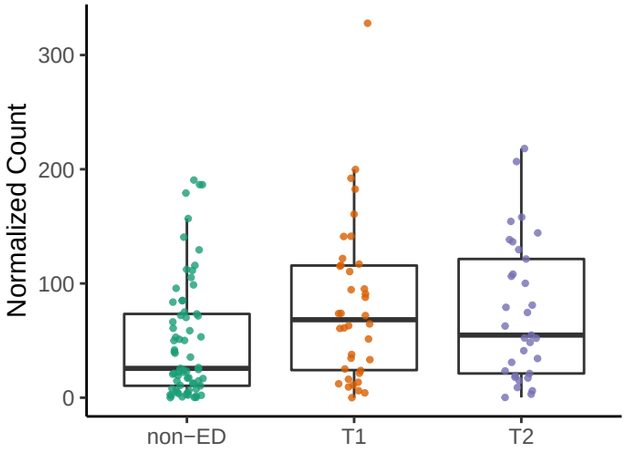
DAPLYSINESYN−PWY: L−lysine biosyr

non−ED vs. T1 adjusted p = 0.066
non−ED vs. T2 adjusted p = 0.55
T1 vs. T2 adjusted p = 0.29



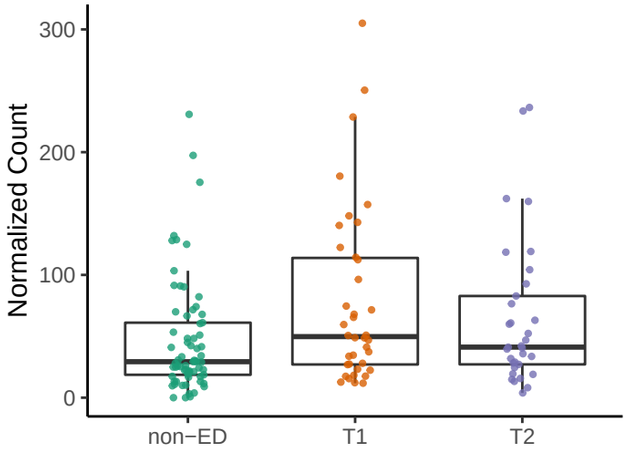
ARGININE−SYN4−PWY: L−ornithine d

non−ED vs. T1 adjusted p = 0.068
non−ED vs. T2 adjusted p = 0.14
T1 vs. T2 adjusted p = 0.66



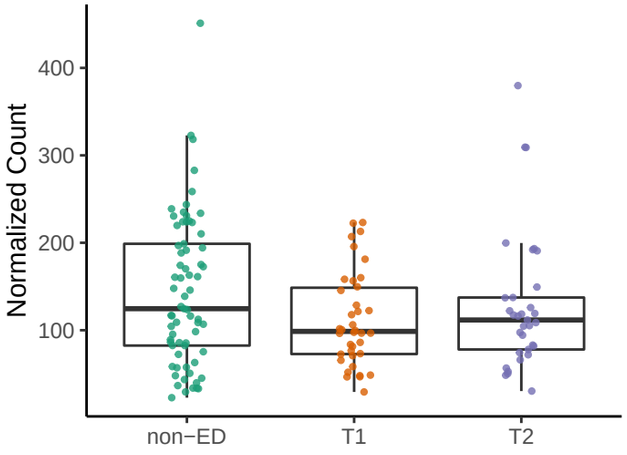
PWY66−400: glycolysis VI (metazoan)

non−ED vs. T1 adjusted p = 0.068
non−ED vs. T2 adjusted p = 0.26
T1 vs. T2 adjusted p = 0.37



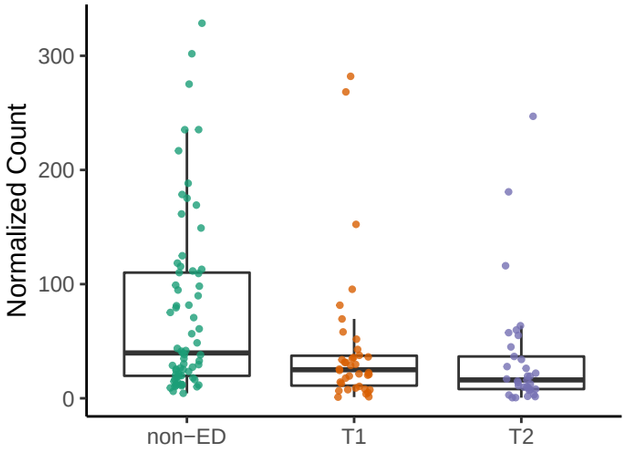
PWY0−1586: peptidoglycan maturation

non−ED vs. T1 adjusted p = 0.068
non−ED vs. T2 adjusted p = 0.53
T1 vs. T2 adjusted p = 0.4



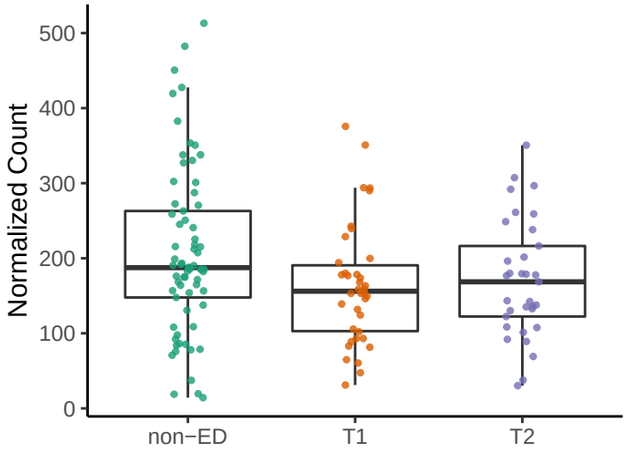
PENTOSE−P−PWY: pentose phosphat

non−ED vs. T1 adjusted p = 0.081
non−ED vs. T2 adjusted p = 0.05
T1 vs. T2 adjusted p = 0.45



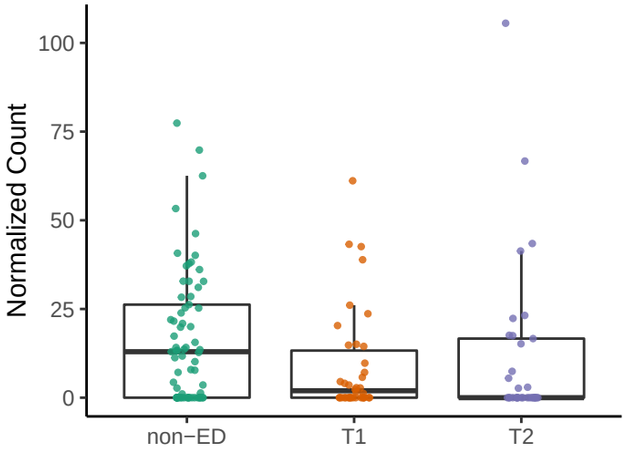
TRPSYN−PWY: L−tryptophan biosynth

non−ED vs. T1 adjusted p = 0.082
non−ED vs. T2 adjusted p = 0.15
T1 vs. T2 adjusted p = 0.86



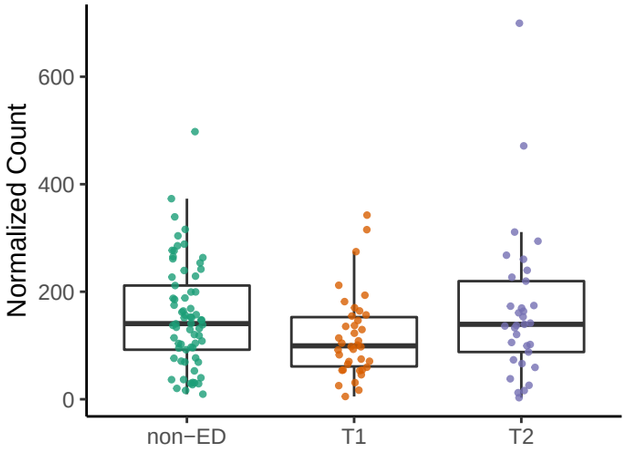
PWY−7209: superpathway of pyrimidin

non−ED vs. T1 adjusted p = 0.085
non−ED vs. T2 adjusted p = 0.44
T1 vs. T2 adjusted p = 0.61



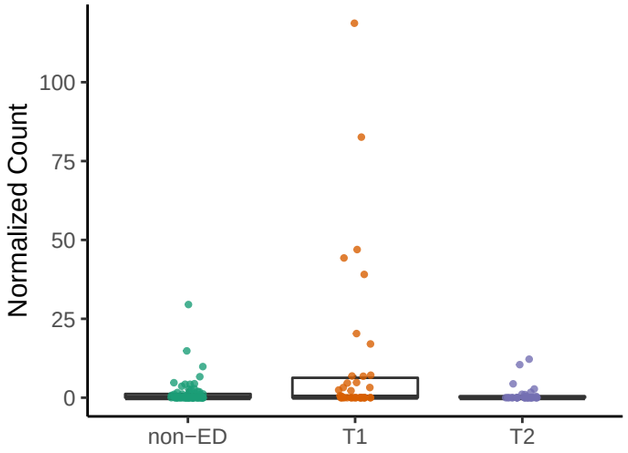
PWY−7234: inosine−5'−phosphate bios

non−ED vs. T1 adjusted p = 0.086
non−ED vs. T2 adjusted p = 0.78
T1 vs. T2 adjusted p = 0.13



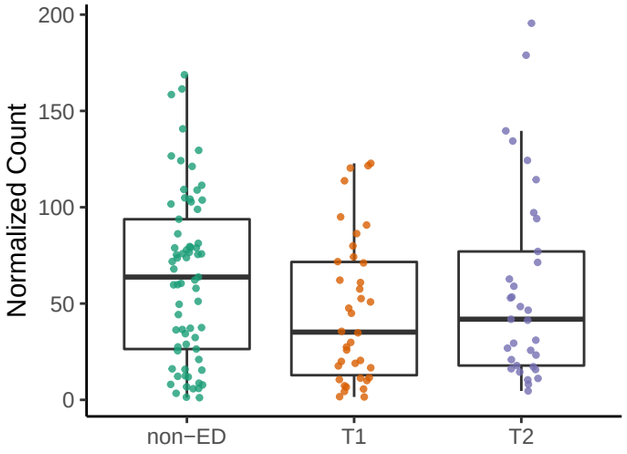
PWY−7013: L−1,2−propanediol degrad

non−ED vs. T1 adjusted p = 0.091
non−ED vs. T2 adjusted p = 0.62
T1 vs. T2 adjusted p = 0.14



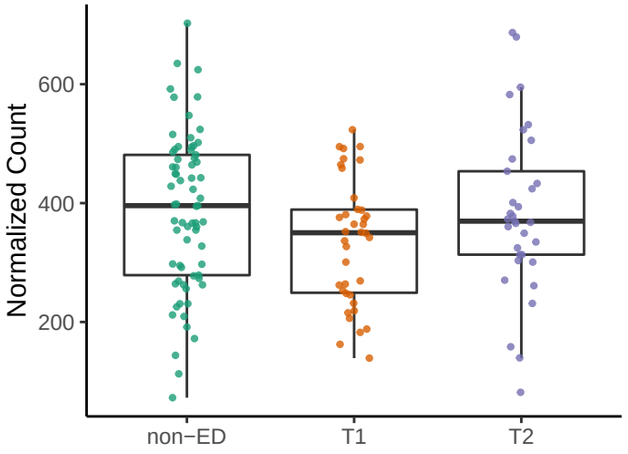
PWY−5989: stearate biosynthesis II (ba

non−ED vs. T1 adjusted p = 0.093
non−ED vs. T2 adjusted p = 0.71
T1 vs. T2 adjusted p = 0.44



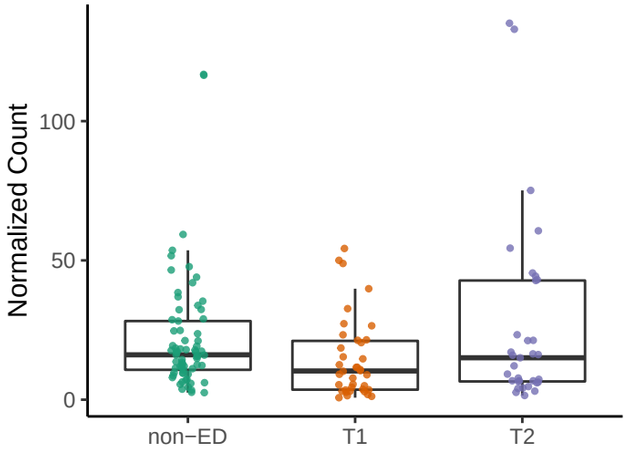
PWY−6385: peptidoglycan biosynthesis

non−ED vs. T1 adjusted p = 0.093
non−ED vs. T2 adjusted p = 0.96
T1 vs. T2 adjusted p = 0.22



PWY−6588: pyruvate fermentation to a

non−ED vs. T1 adjusted p = 0.093
non−ED vs. T2 adjusted p = 0.64
T1 vs. T2 adjusted p = 0.17



FOLSYN−PWY: superpathway of tetrahydrofolate biosynthesis and salvage

non−ED vs. T1 adjusted p = 0.095
non−ED vs. T2 adjusted p = 0.32
T1 vs. T2 adjusted p = 0.52