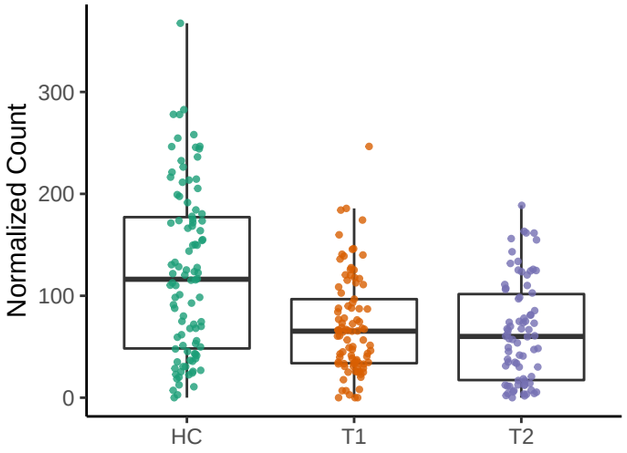
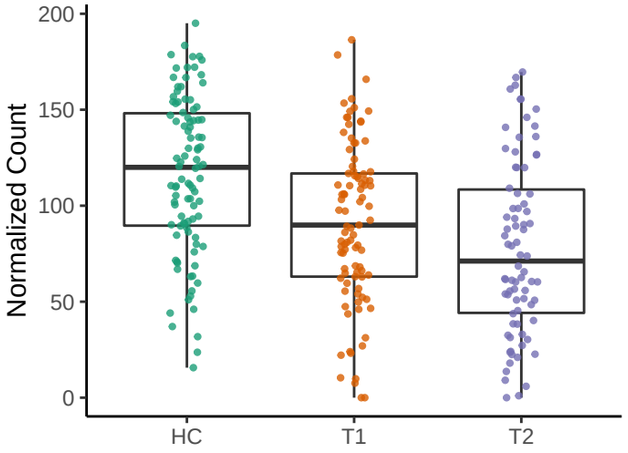
P4−PWY: superpathway of L−lysine, L-

HC vs. T1 adjusted p = 3.1e−06
HC vs. T2 adjusted p = 3.9e−06
T1 vs. T2 adjusted p = 0.59



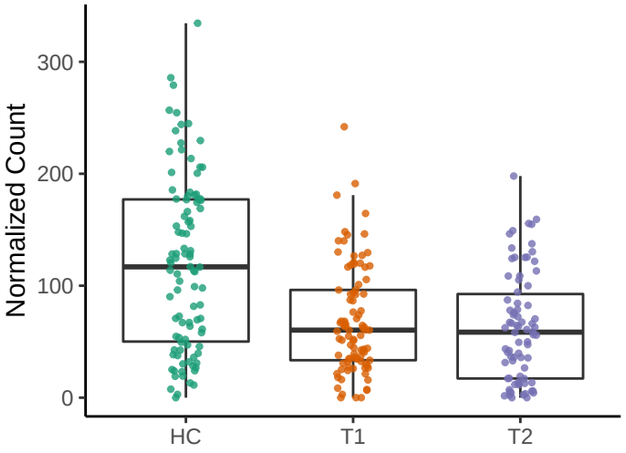
PRPP−PWY: superpathway of histidine

HC vs. T1 adjusted p = 0.00022
HC vs. T2 adjusted p = 3.9e−06
T1 vs. T2 adjusted p = 0.029



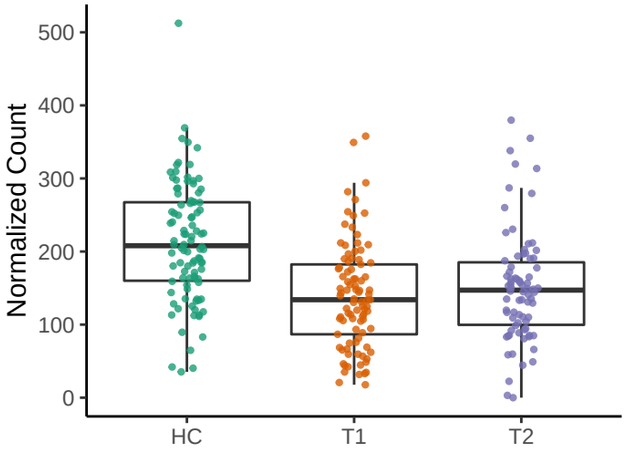
PWY0−781: aspartate superpathway

HC vs. T1 adjusted p = 1.8e−06
HC vs. T2 adjusted p = 3.9e−06
T1 vs. T2 adjusted p = 0.6



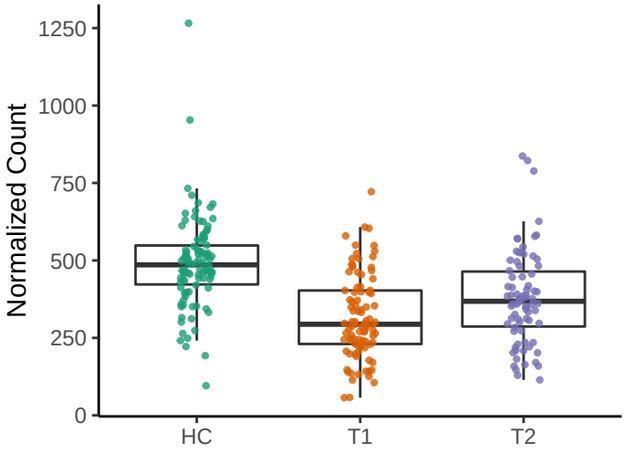
PWY−5347: superpathway of L−methio

HC vs. T1 adjusted p = 9.2e−09
HC vs. T2 adjusted p = 4.5e−05
T1 vs. T2 adjusted p = 0.24



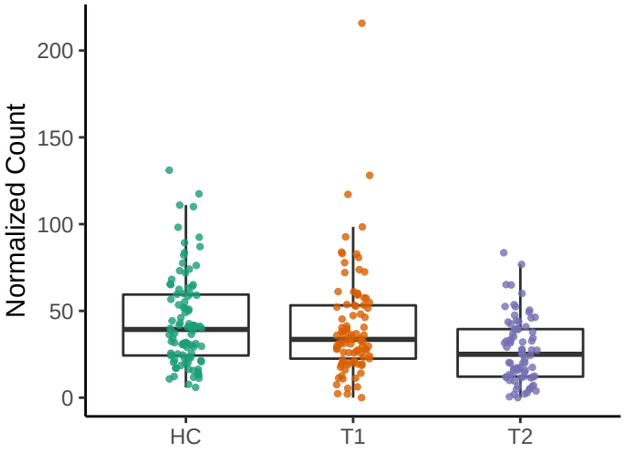
DTDPRHAMSYN−PWY: dTDP−L−rhaı

HC vs. T1 adjusted p = 6.6e−12
HC vs. T2 adjusted p = 0.00014
T1 vs. T2 adjusted p = 0.011



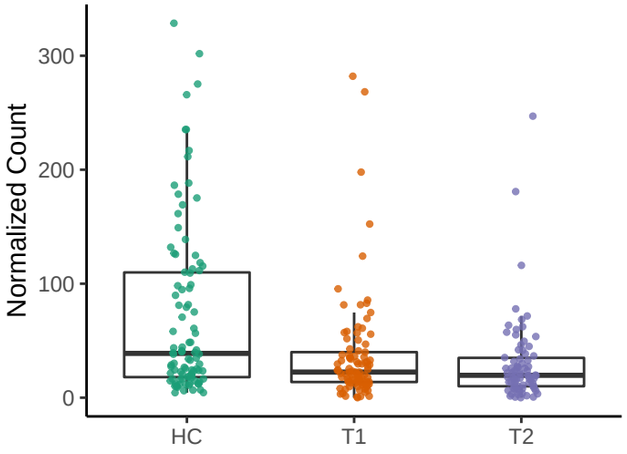
TCA: TCA cycle I (prokaryotic)

HC vs. T1 adjusted p = 0.6
HC vs. T2 adjusted p = 0.00014
T1 vs. T2 adjusted p = 0.0069



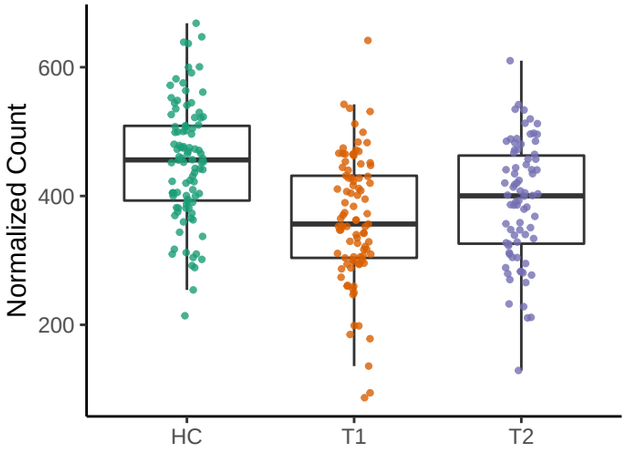
PENTOSE−P−PWY: pentose phosphat

HC vs. T1 adjusted p = 0.0011
HC vs. T2 adjusted p = 0.00017
T1 vs. T2 adjusted p = 0.29



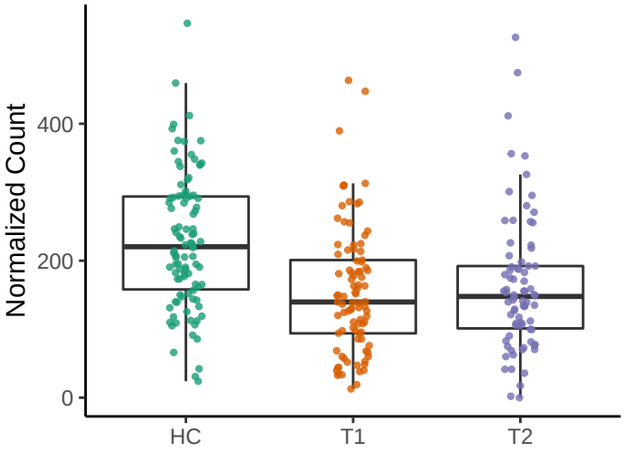
TRNA−CHARGING−PWY: tRNA chargi

HC vs. T1 adjusted p = 1.9e−08
HC vs. T2 adjusted p = 0.00067
T1 vs. T2 adjusted p = 0.078



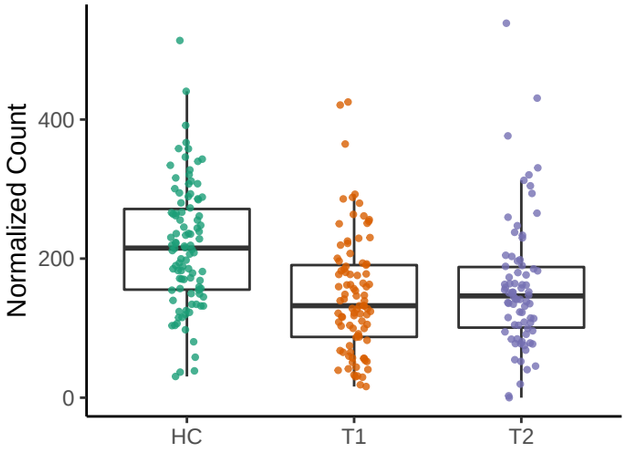
MET−SAM−PWY: superpathway of S−a

HC vs. T1 adjusted p = 3.5e−06
HC vs. T2 adjusted p = 0.00095
T1 vs. T2 adjusted p = 0.49



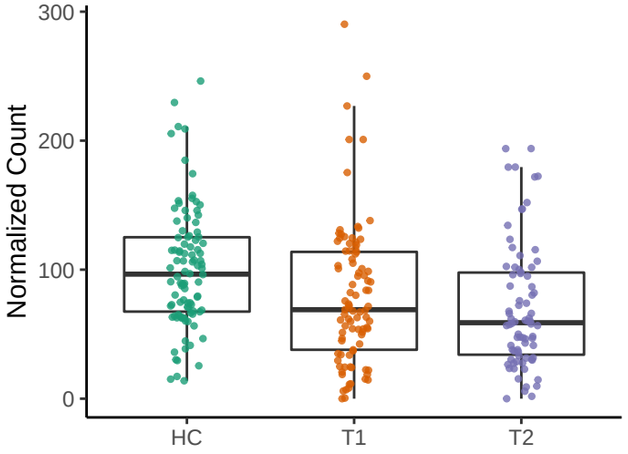
METSYN−PWY: L−homoserine and L−

HC vs. T1 adjusted p = 9.3e−07
HC vs. T2 adjusted p = 0.00095
T1 vs. T2 adjusted p = 0.32



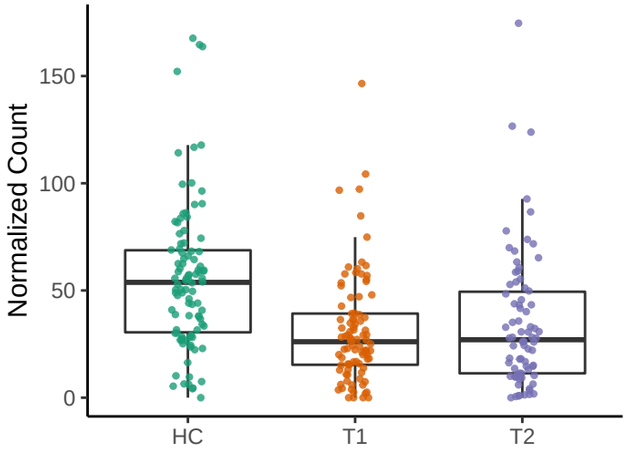
HSERMETANA−PWY: L−methionine bi

HC vs. T1 adjusted p = 0.014
HC vs. T2 adjusted p = 0.0013
T1 vs. T2 adjusted p = 0.55



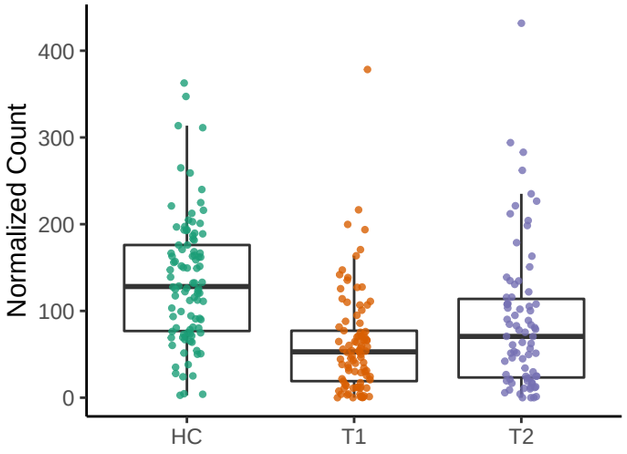
POLYAMSYN−PWY: superpathway of p

HC vs. T1 adjusted p = 1.1e−06
HC vs. T2 adjusted p = 0.0013
T1 vs. T2 adjusted p = 0.43



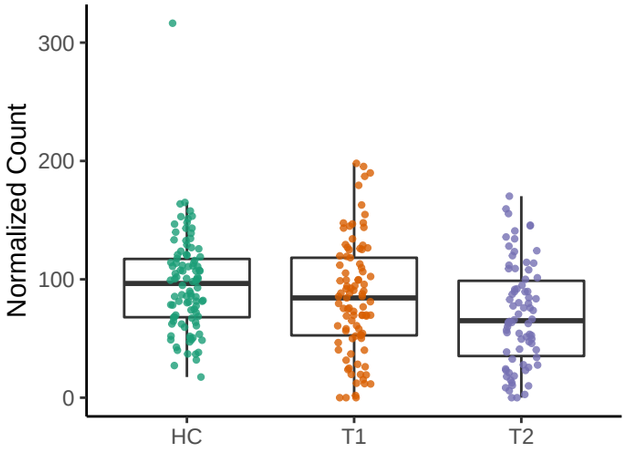
PWY−5177: glutaryl−CoA degradation

HC vs. T1 adjusted p = 1.8e−10
HC vs. T2 adjusted p = 0.0023
T1 vs. T2 adjusted p = 0.019



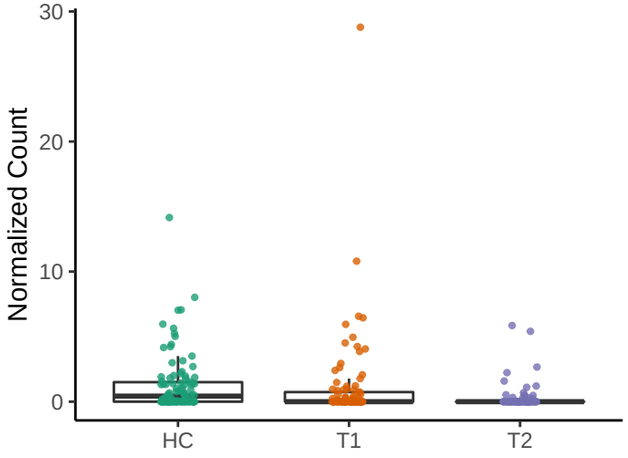
PWY−7211: superpathway of pyrimidin

HC vs. T1 adjusted p = 0.17
HC vs. T2 adjusted p = 0.0023
T1 vs. T2 adjusted p = 0.018



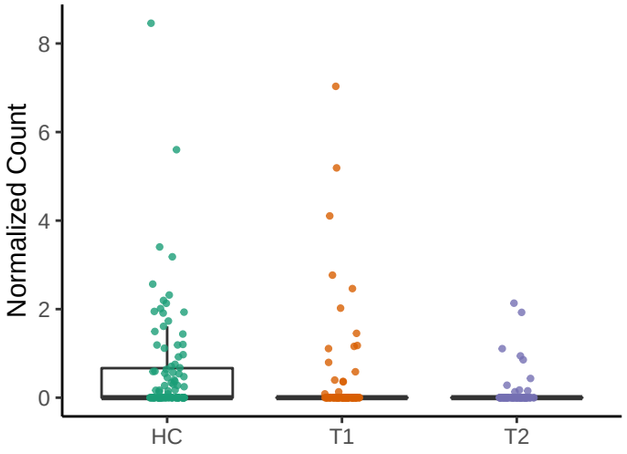
PWY490−3: nitrate reduction VI (assimil

HC vs. T1 adjusted p = 0.78
HC vs. T2 adjusted p = 0.0027
T1 vs. T2 adjusted p = 0.14



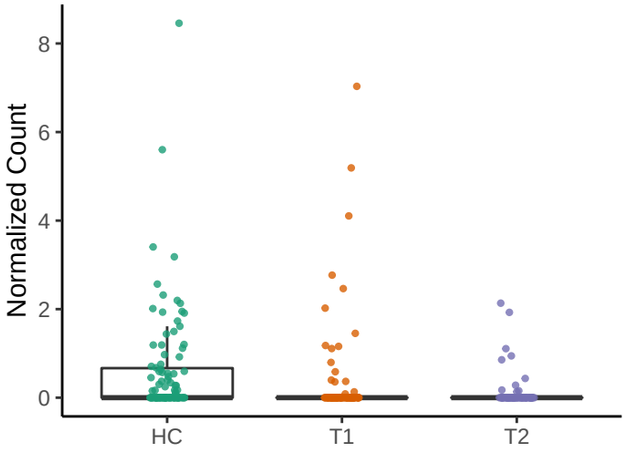
PWY−5692: allantoin degradation to glyo

HC vs. T1 adjusted p = 0.18
HC vs. T2 adjusted p = 0.0032
T1 vs. T2 adjusted p = 0.074



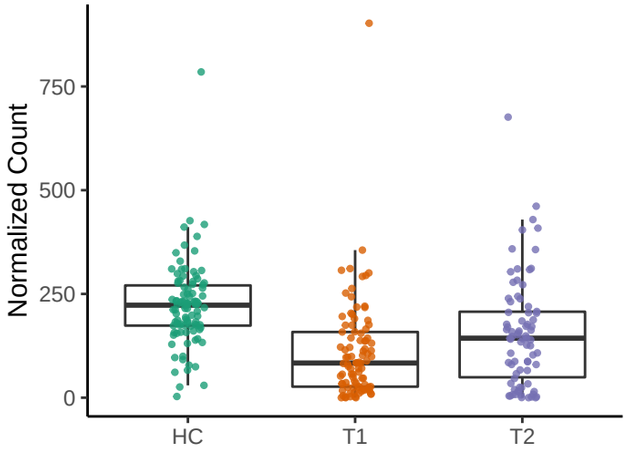
URDEGR−PWY: superpathway of allanto

HC vs. T1 adjusted p = 0.18
HC vs. T2 adjusted p = 0.0032
T1 vs. T2 adjusted p = 0.074



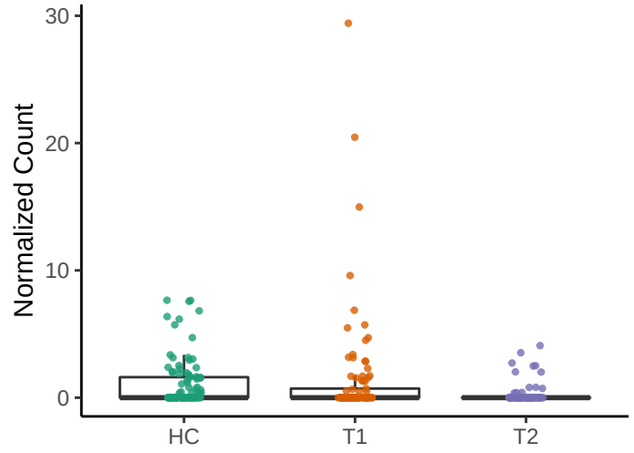
COBALSYN−PWY: adenosylcobalamin

HC vs. T1 adjusted p = 4.3e−09
HC vs. T2 adjusted p = 0.0041
T1 vs. T2 adjusted p = 0.0069



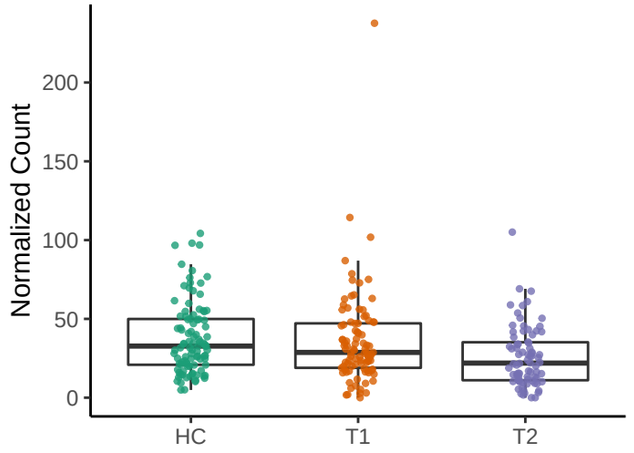
PWY−6263: superpathway of menaquin

HC vs. T1 adjusted p = 0.57
HC vs. T2 adjusted p = 0.0041
T1 vs. T2 adjusted p = 0.056



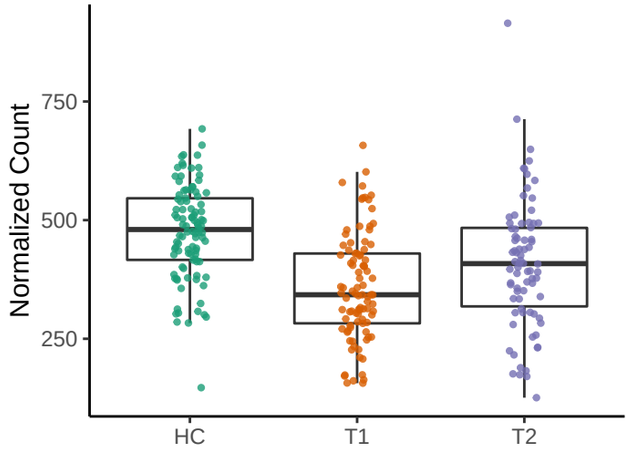
PWY−6969: TCA cycle V (2−oxoglutara

HC vs. T1 adjusted p = 0.79
HC vs. T2 adjusted p = 0.0041
T1 vs. T2 adjusted p = 0.019



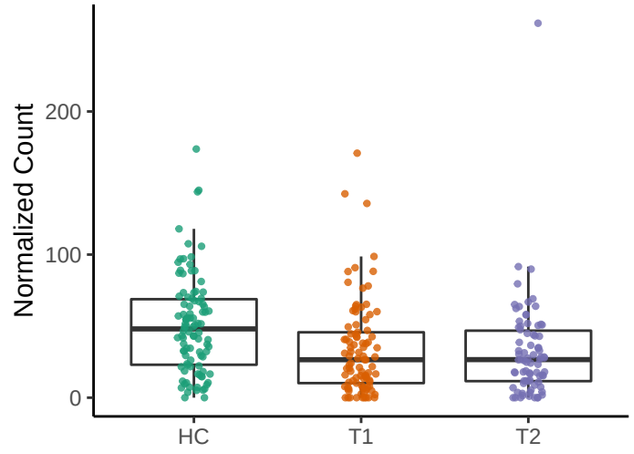
CALVIN−PWY: Calvin−Benson−Bassha

HC vs. T1 adjusted p = 1.5e−10
HC vs. T2 adjusted p = 0.0048
T1 vs. T2 adjusted p = 0.018



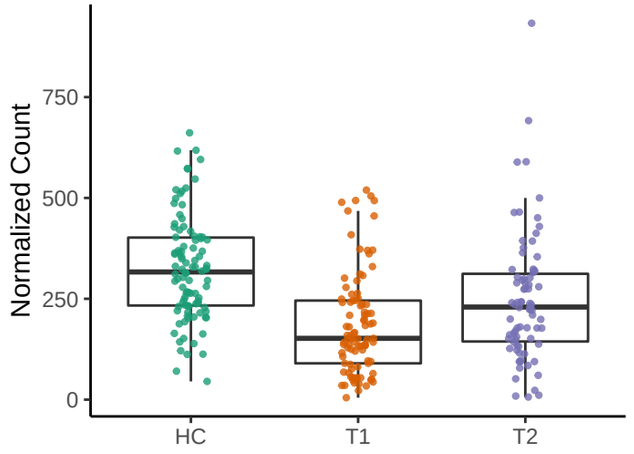
ARG+POLYAMINE−SYN: superpathwa

HC vs. T1 adjusted p = 0.0033
HC vs. T2 adjusted p = 0.0099
T1 vs. T2 adjusted p = 0.81



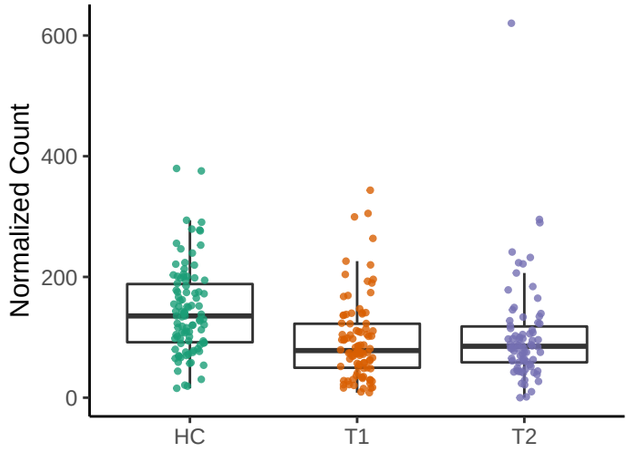
PWY−621: sucrose degradation III (suc

HC vs. T1 adjusted p = 1.8e−10
HC vs. T2 adjusted p = 0.0099
T1 vs. T2 adjusted p = 0.018



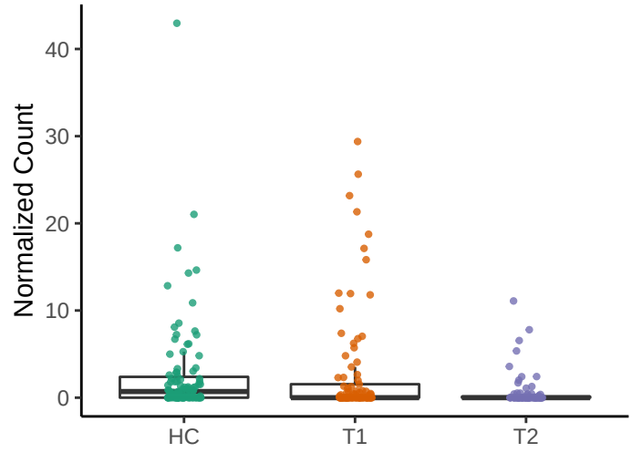
HOMOSER−METSYN−PWY: L−methio

HC vs. T1 adjusted p = 1.9e−05
HC vs. T2 adjusted p = 0.01
T1 vs. T2 adjusted p = 0.32



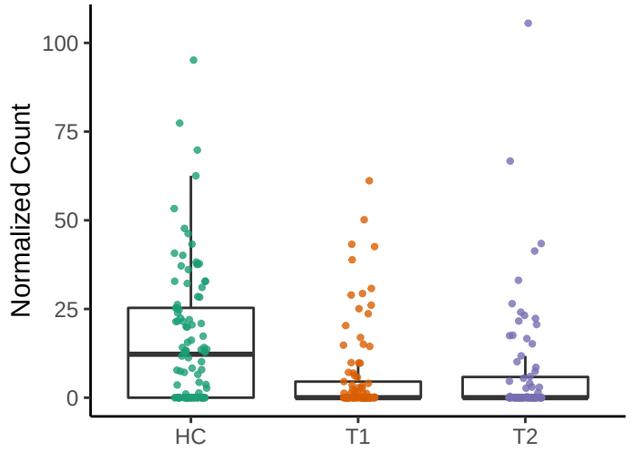
KETOGLUCONMET−PWY: ketoglucona

HC vs. T1 adjusted p = 0.91
HC vs. T2 adjusted p = 0.011
T1 vs. T2 adjusted p = 0.018



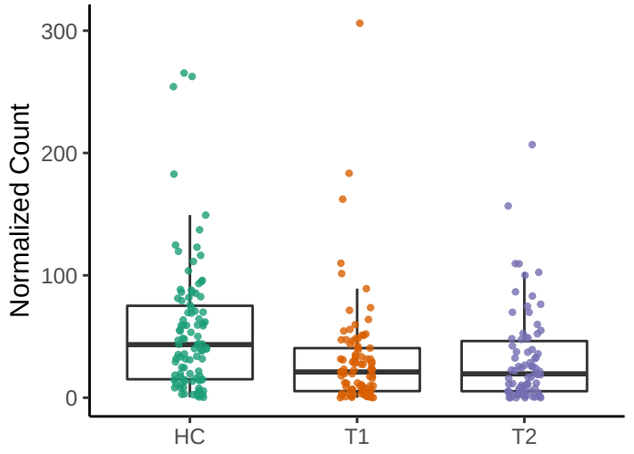
PWY−7209: superpathway of pyrimidin

HC vs. T1 adjusted p = 0.00014
HC vs. T2 adjusted p = 0.011
T1 vs. T2 adjusted p = 0.57



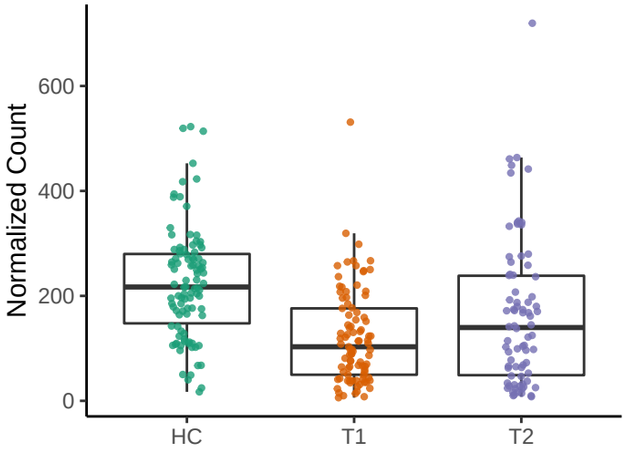
LACTOSECAT−PWY: lactose and galac

HC vs. T1 adjusted p = 0.0029
HC vs. T2 adjusted p = 0.012
T1 vs. T2 adjusted p = 0.54



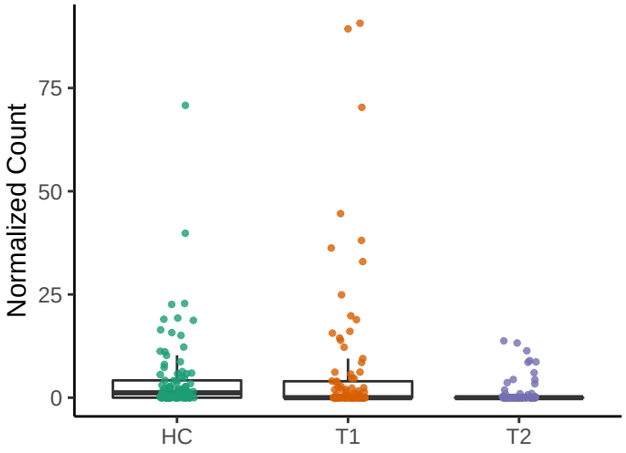
PWY−7242: D−fructuronate degradatio

HC vs. T1 adjusted p = 9.7e−10
HC vs. T2 adjusted p = 0.019
T1 vs. T2 adjusted p = 0.027



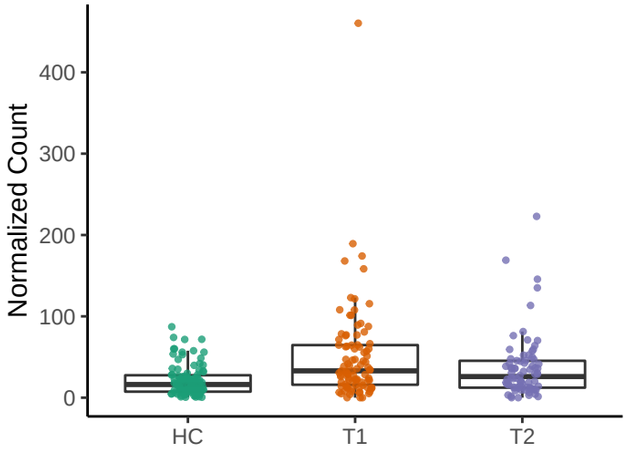
PWY−7204: pyridoxal 5'−phosphate salv

HC vs. T1 adjusted p = 0.39
HC vs. T2 adjusted p = 0.019
T1 vs. T2 adjusted p = 0.023



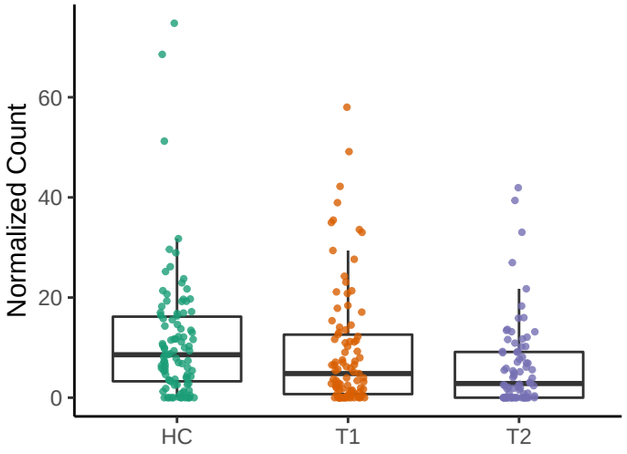
CITRULBIO−PWY: L−citrulline biosynth

HC vs. T1 adjusted p = 9.4e−05
HC vs. T2 adjusted p = 0.021
T1 vs. T2 adjusted p = 0.11



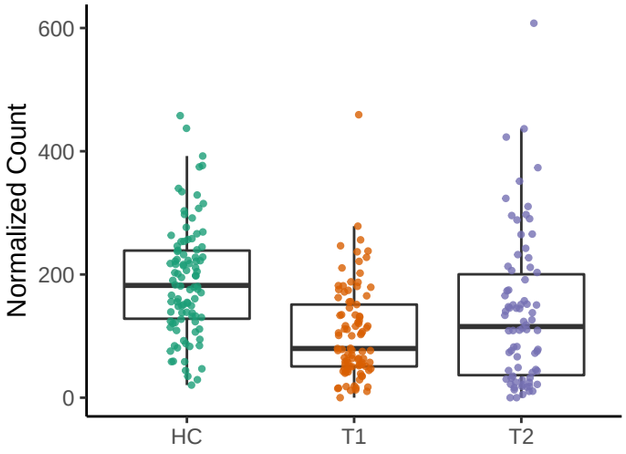
FAO−PWY: fatty acid β−oxidation I

HC vs. T1 adjusted p = 0.42
HC vs. T2 adjusted p = 0.021
T1 vs. T2 adjusted p = 0.065



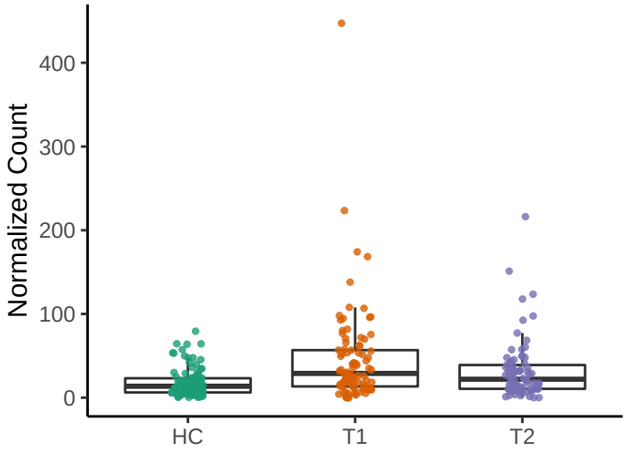
GLUCUROCAT−PWY: superpathway o

HC vs. T1 adjusted p = 2.4e−09
HC vs. T2 adjusted p = 0.021
T1 vs. T2 adjusted p = 0.033



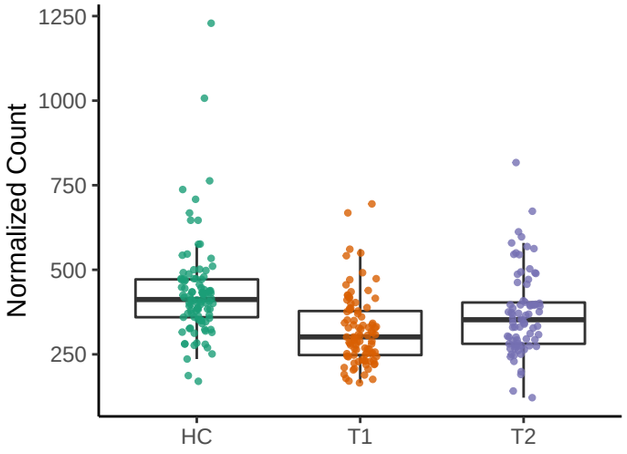
PWY−4984: urea cycle

HC vs. T1 adjusted p = 0.00019
HC vs. T2 adjusted p = 0.021
T1 vs. T2 adjusted p = 0.12



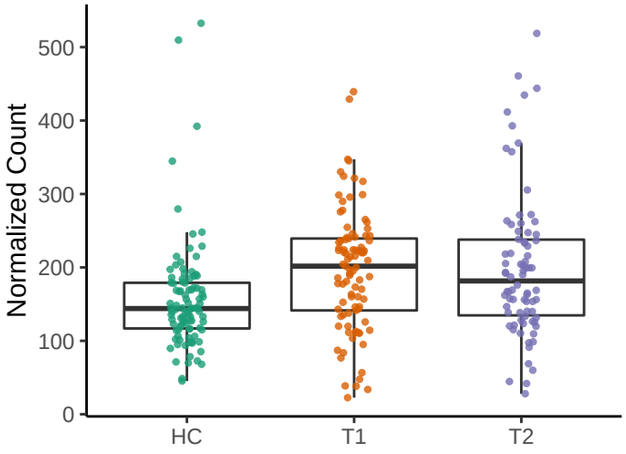
PWY−3001: superpathway of L−isoleu

HC vs. T1 adjusted p = 8.9e−08
HC vs. T2 adjusted p = 0.021
T1 vs. T2 adjusted p = 0.0069



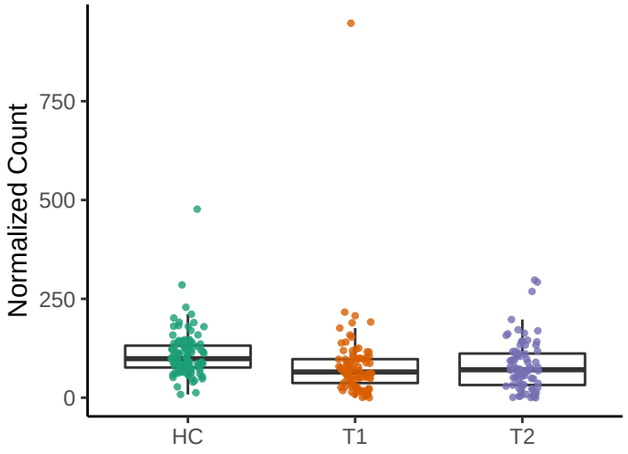
PWY−7208: superpathway of pyrimidin

HC vs. T1 adjusted p = 0.0033
HC vs. T2 adjusted p = 0.021
T1 vs. T2 adjusted p = 0.88



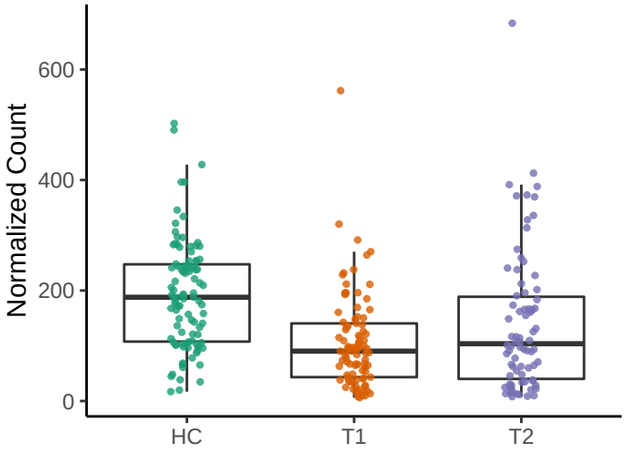
PWY−6305: putrescine biosynthesis IV

HC vs. T1 adjusted p = 0.046
HC vs. T2 adjusted p = 0.024
T1 vs. T2 adjusted p = 0.12



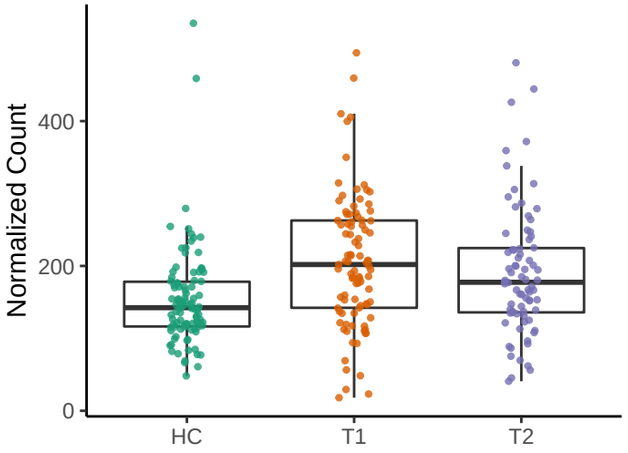
PWY−6507: 4−deoxy−L−threo−hex−4−

HC vs. T1 adjusted p = 1.8e−08
HC vs. T2 adjusted p = 0.024
T1 vs. T2 adjusted p = 0.042



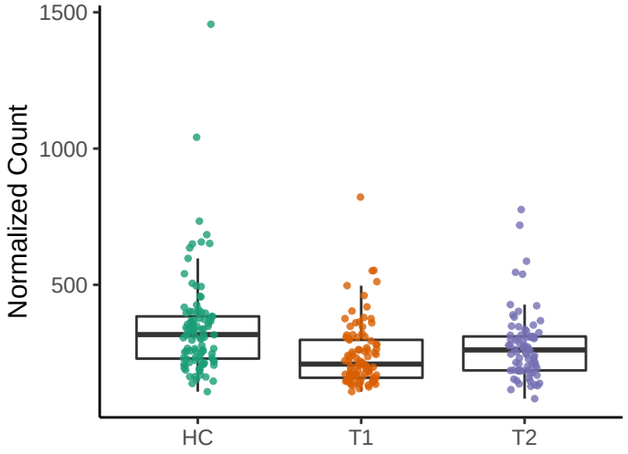
PWY−7228: superpathway of guanosin

HC vs. T1 adjusted p = 0.00011
HC vs. T2 adjusted p = 0.025
T1 vs. T2 adjusted p = 0.15



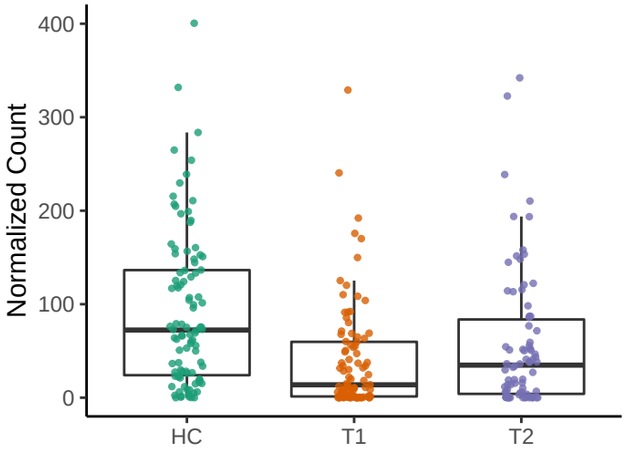
HISTSYN−PWY: L−histidine biosynthe

HC vs. T1 adjusted p = 0.00011
HC vs. T2 adjusted p = 0.025
T1 vs. T2 adjusted p = 0.027



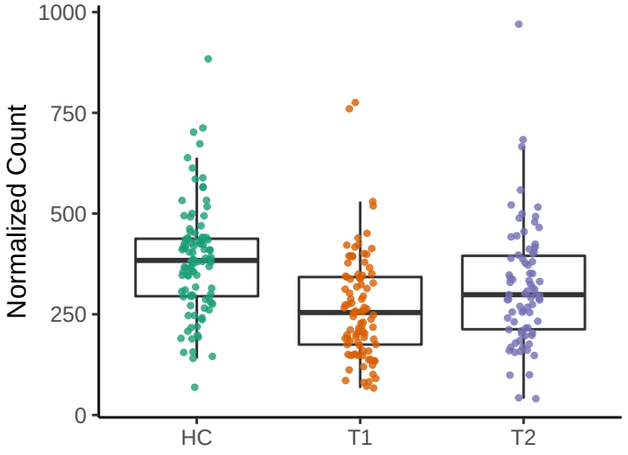
PWY−5367: petroselinate biosynthesis

HC vs. T1 adjusted p = 9.1e−06
HC vs. T2 adjusted p = 0.025
T1 vs. T2 adjusted p = 0.29



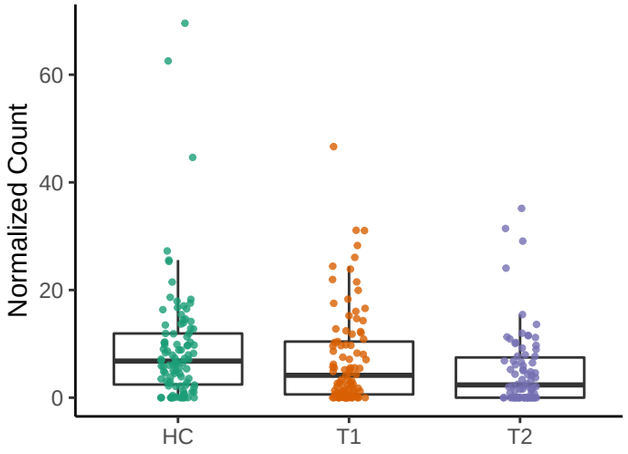
PWY−7357: thiamin formation from py

HC vs. T1 adjusted p = 9.8e−08
HC vs. T2 adjusted p = 0.025
T1 vs. T2 adjusted p = 0.011



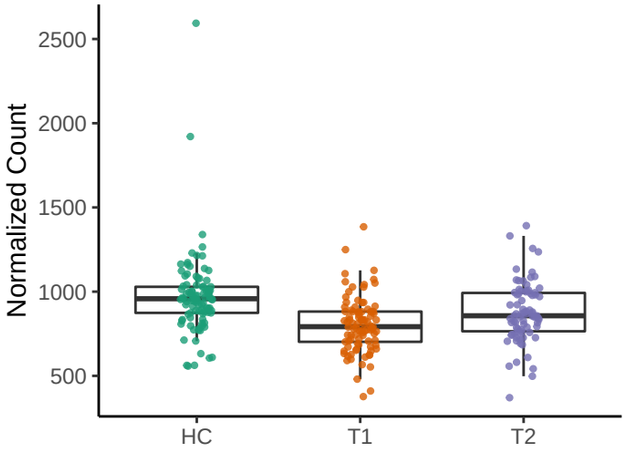
PWY−5136: fatty acid β−oxidation

HC vs. T1 adjusted p = 0.29
HC vs. T2 adjusted p = 0.027
T1 vs. T2 adjusted p = 0.096



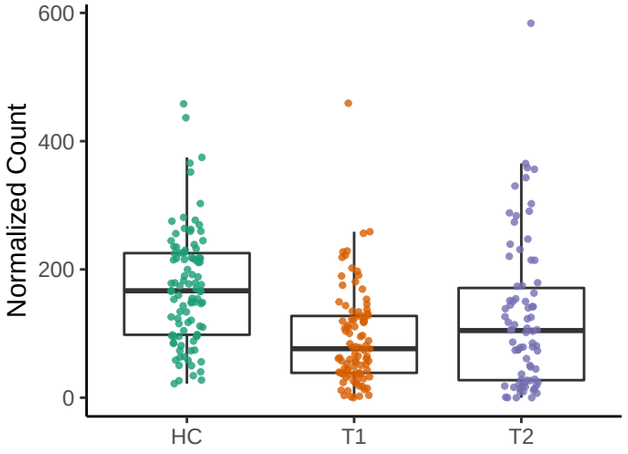
PWY−7219: adenosine ribonucleotides

HC vs. T1 adjusted p = 1.1e−06
HC vs. T2 adjusted p = 0.028
T1 vs. T2 adjusted p = 0.02



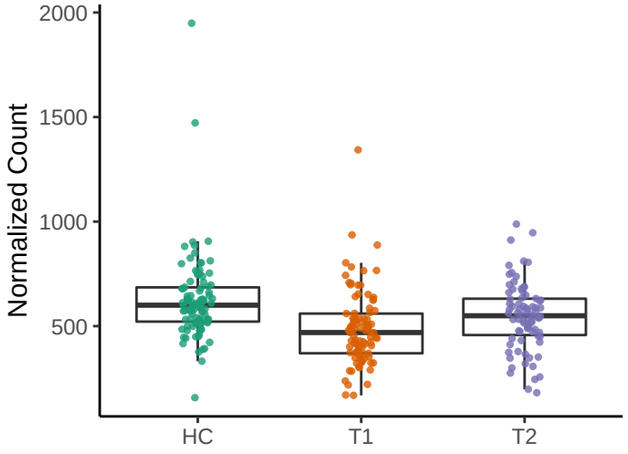
GALACT−GLUCUROCAT−PWY: super

HC vs. T1 adjusted p = 9.2e−09
HC vs. T2 adjusted p = 0.028
T1 vs. T2 adjusted p = 0.033



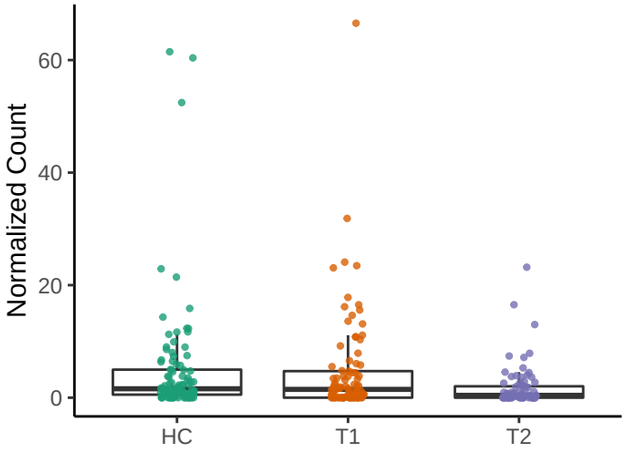
PWY−6151: S−adenosyl−L−methionin

HC vs. T1 adjusted p = 1.5e−05
HC vs. T2 adjusted p = 0.033
T1 vs. T2 adjusted p = 0.011



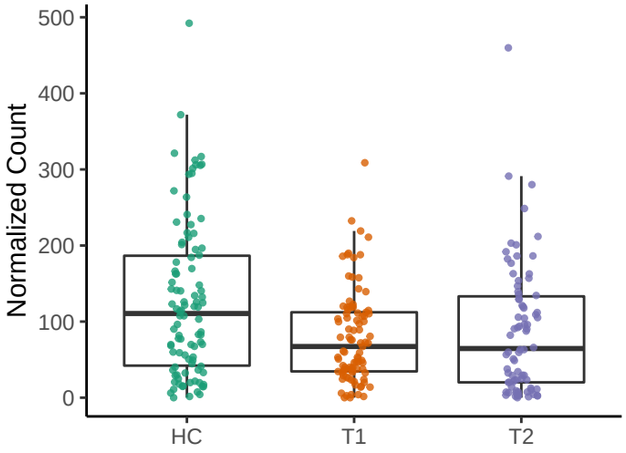
REDCITCYC: TCA cycle VIII (helicobact

HC vs. T1 adjusted p = 0.87
HC vs. T2 adjusted p = 0.037
T1 vs. T2 adjusted p = 0.019



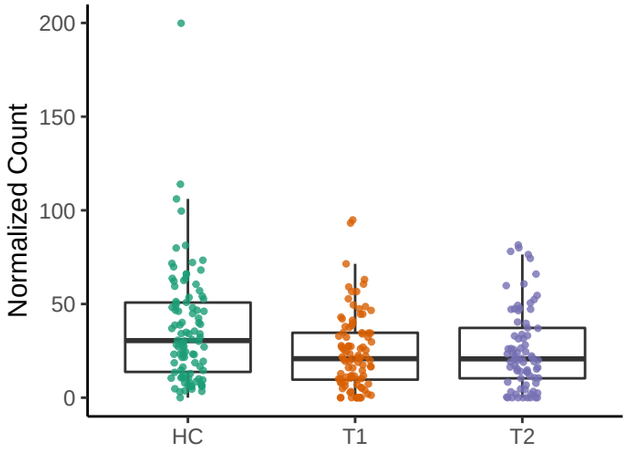
DAPLYSINESYN−PWY: L−lysine biosyr

HC vs. T1 adjusted p = 0.00074
HC vs. T2 adjusted p = 0.041
T1 vs. T2 adjusted p = 0.37



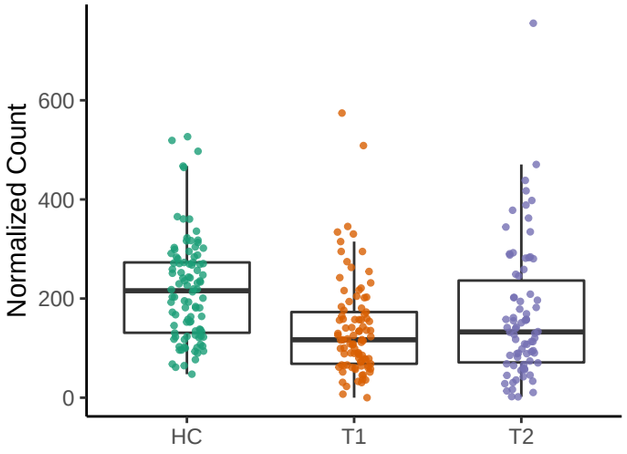
FOLSYN−PWY: superpathway of tetrah

HC vs. T1 adjusted p = 0.0048
HC vs. T2 adjusted p = 0.041
T1 vs. T2 adjusted p = 0.54



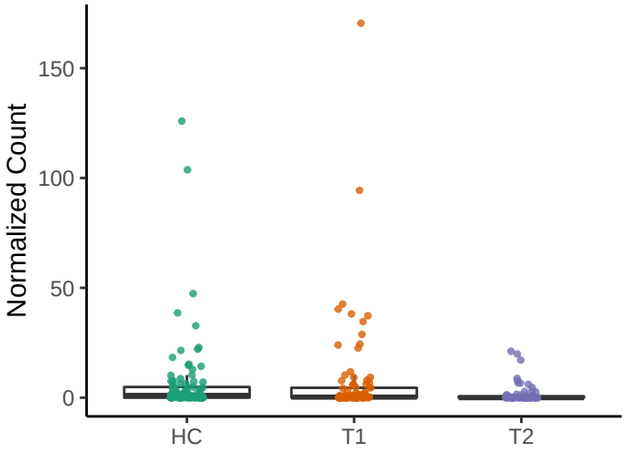
GLCMANNANAUT−PWY: superpathwa

HC vs. T1 adjusted p = 2.8e−06
HC vs. T2 adjusted p = 0.041
T1 vs. T2 adjusted p = 0.071



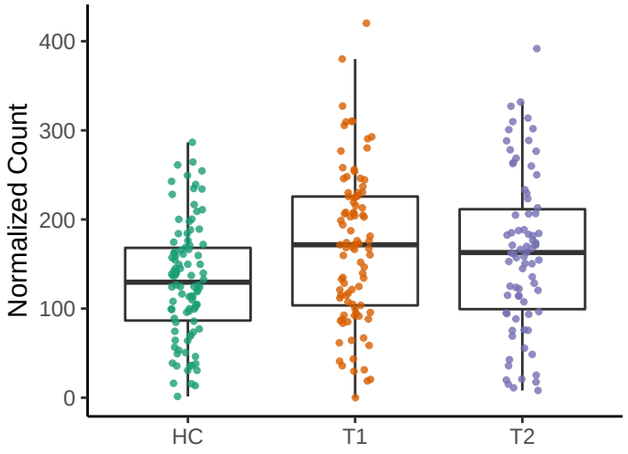
GLUCOSE1PMETAB−PWY: glucose ar

HC vs. T1 adjusted p = 0.86
HC vs. T2 adjusted p = 0.041
T1 vs. T2 adjusted p = 0.029



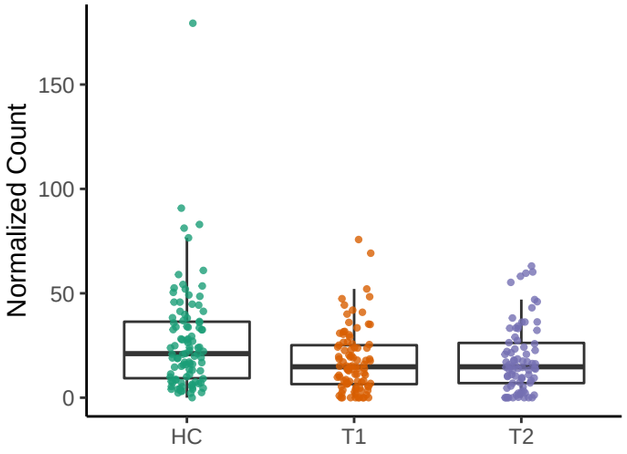
PWY−6168: flavin biosynthesis III (fung

HC vs. T1 adjusted p = 0.0021
HC vs. T2 adjusted p = 0.041
T1 vs. T2 adjusted p = 0.36



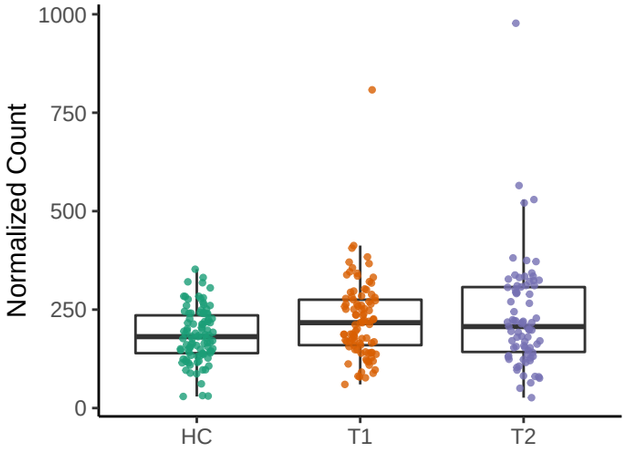
PWY−6612: superpathway of tetrahydr

HC vs. T1 adjusted p = 0.0063
HC vs. T2 adjusted p = 0.041
T1 vs. T2 adjusted p = 0.53



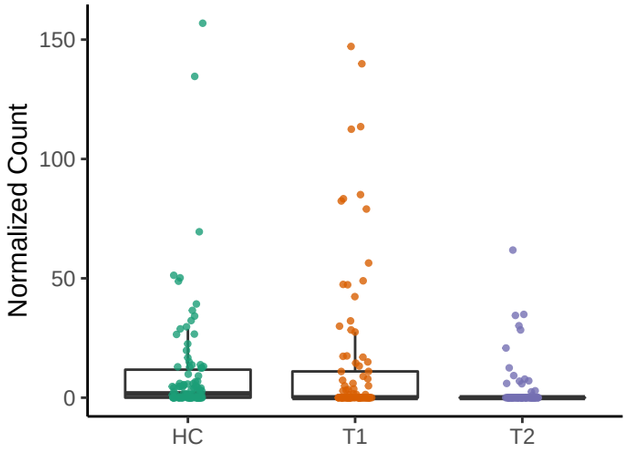
PWY−6703: preQ0 biosynthesis

HC vs. T1 adjusted p = 0.0069
HC vs. T2 adjusted p = 0.041
T1 vs. T2 adjusted p = 0.99



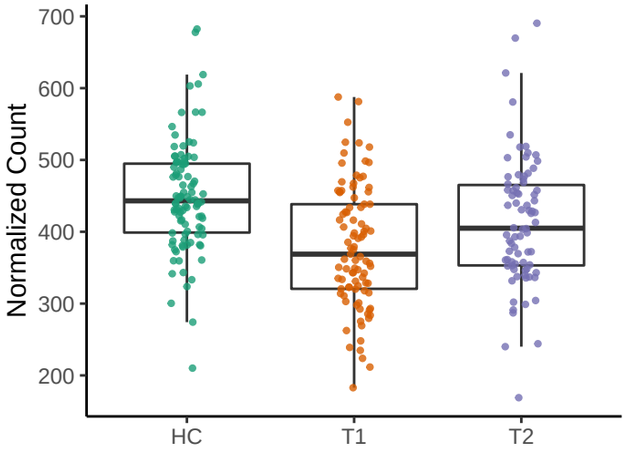
PWY−6895: superpathway of thiamin d

HC vs. T1 adjusted p = 0.46
HC vs. T2 adjusted p = 0.041
T1 vs. T2 adjusted p = 0.018



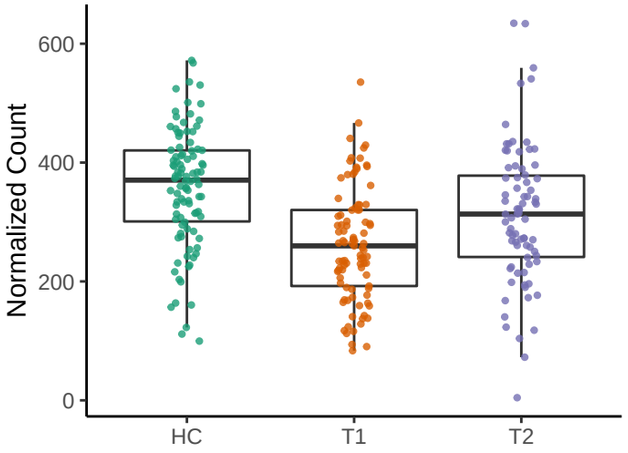
PWY−724: superpathway of L−lysine, L

HC vs. T1 adjusted p = 2.6e−07
HC vs. T2 adjusted p = 0.041
T1 vs. T2 adjusted p = 0.024



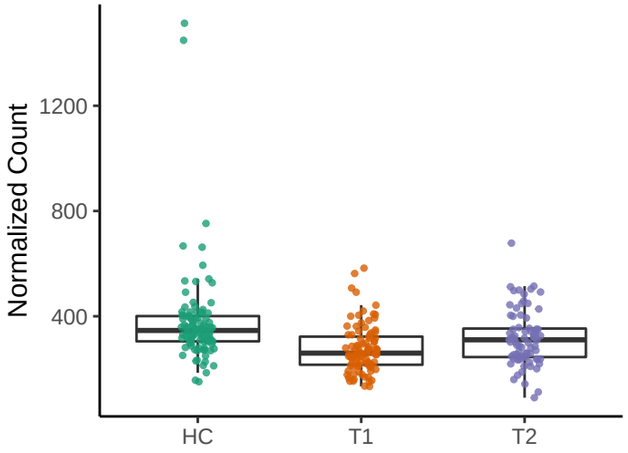
PWY66−422: D−galactose degradation

HC vs. T1 adjusted p = 5.3e−09
HC vs. T2 adjusted p = 0.041
T1 vs. T2 adjusted p = 0.008



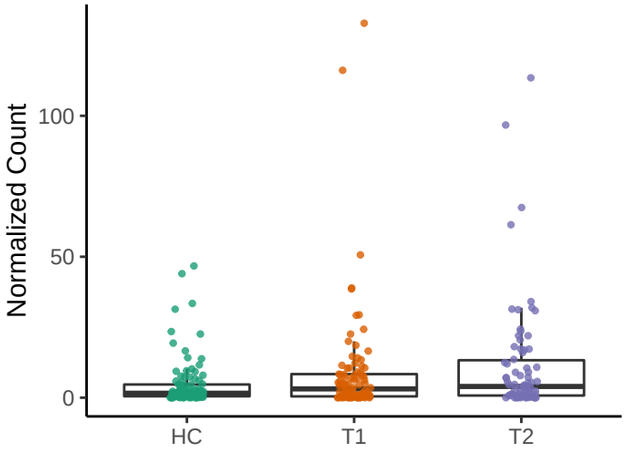
THRESYN−PWY: superpathway of L−t

HC vs. T1 adjusted p = 2.8e−05
HC vs. T2 adjusted p = 0.041
T1 vs. T2 adjusted p = 0.016



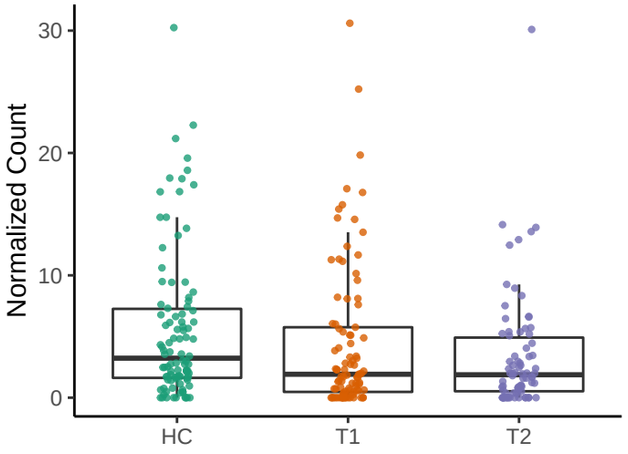
PWY−5384: sucrose degradation IV (su

HC vs. T1 adjusted p = 0.13
HC vs. T2 adjusted p = 0.042
T1 vs. T2 adjusted p = 0.34



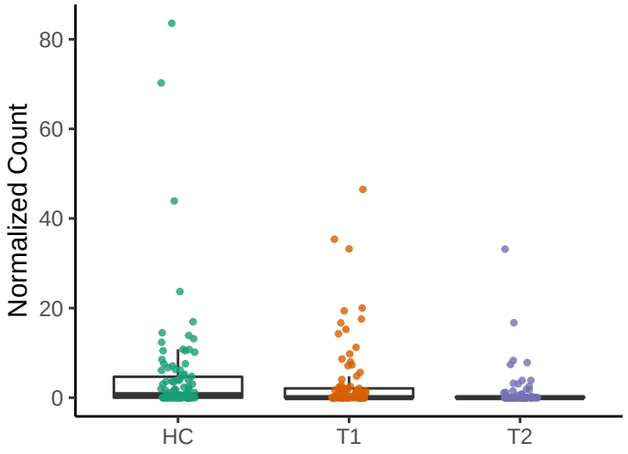
PWY−7115: C4 photosynthetic carbon a

HC vs. T1 adjusted p = 0.32
HC vs. T2 adjusted p = 0.042
T1 vs. T2 adjusted p = 0.2



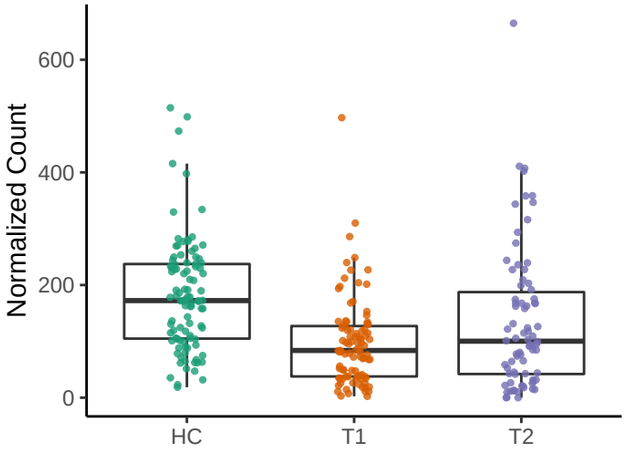
PWY−561: superpathway of glyoxylate c

HC vs. T1 adjusted p = 0.47
HC vs. T2 adjusted p = 0.043
T1 vs. T2 adjusted p = 0.058



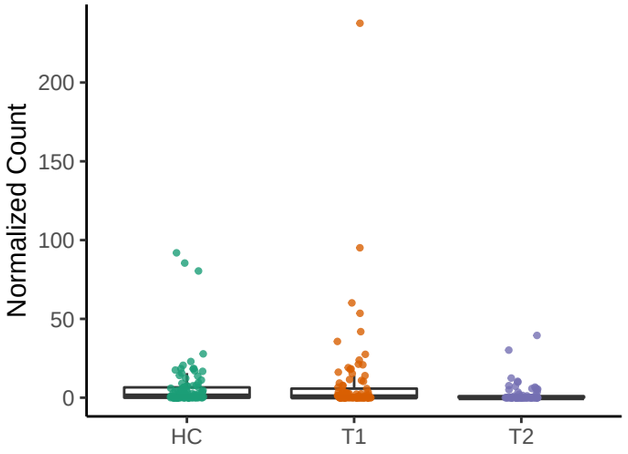
GALACTUROCAT−PWY: D−galacturon

HC vs. T1 adjusted p = 1.9e−08
HC vs. T2 adjusted p = 0.045
T1 vs. T2 adjusted p = 0.029



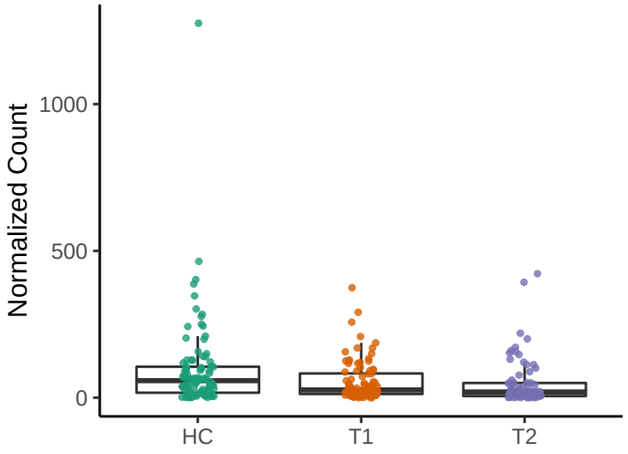
P105−PWY: TCA cycle IV (2−oxoglutar

HC vs. T1 adjusted p = 0.58
HC vs. T2 adjusted p = 0.047
T1 vs. T2 adjusted p = 0.029



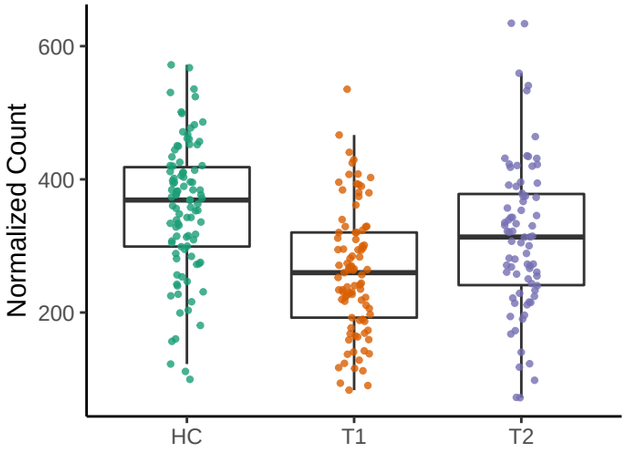
PWY−6147: 6−hydroxymethyl−dihydro

HC vs. T1 adjusted p = 0.048
HC vs. T2 adjusted p = 0.047
T1 vs. T2 adjusted p = 0.82



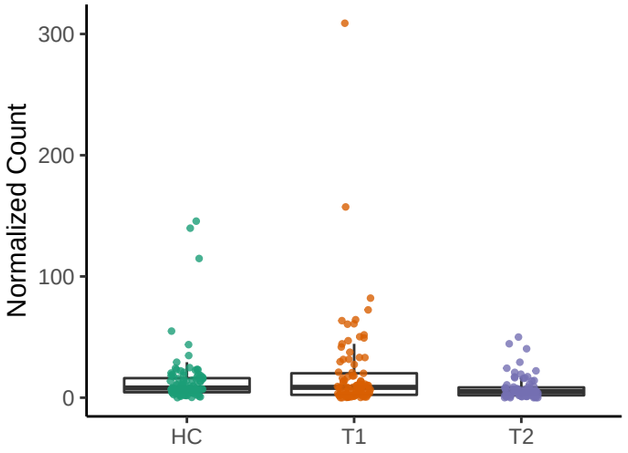
PWY−6317: galactose degradation I (L

HC vs. T1 adjusted p = 6e−09
HC vs. T2 adjusted p = 0.048
T1 vs. T2 adjusted p = 0.0069



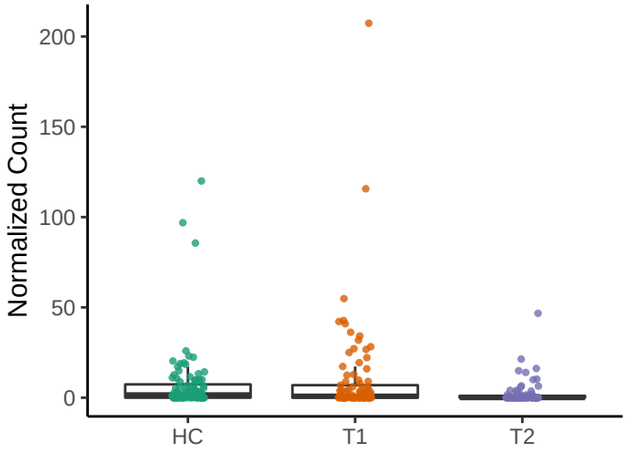
PWY0−1061: superpathway of L−alanir

HC vs. T1 adjusted p = 0.4
HC vs. T2 adjusted p = 0.049
T1 vs. T2 adjusted p = 0.018



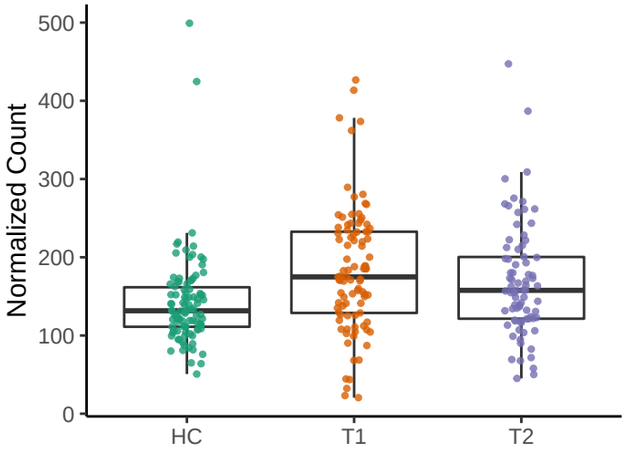
GLYCOLYSIS−TCA−GLYOX−BYPASS:

HC vs. T1 adjusted p = 0.55
HC vs. T2 adjusted p = 0.049
T1 vs. T2 adjusted p = 0.019



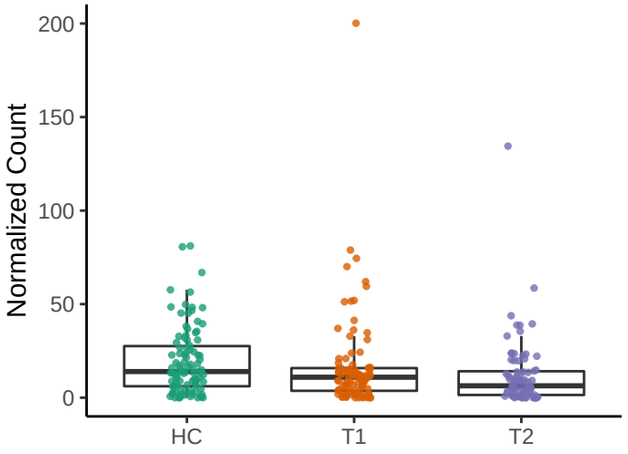
PWY−6125: superpathway of guanosin

HC vs. T1 adjusted p = 0.00052
HC vs. T2 adjusted p = 0.049
T1 vs. T2 adjusted p = 0.12



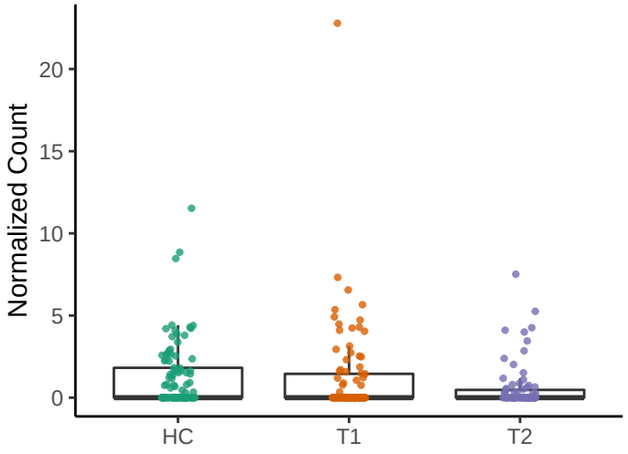
FUCCAT−PWY: fucose degradation

HC vs. T1 adjusted p = 0.56
HC vs. T2 adjusted p = 0.052
T1 vs. T2 adjusted p = 0.3



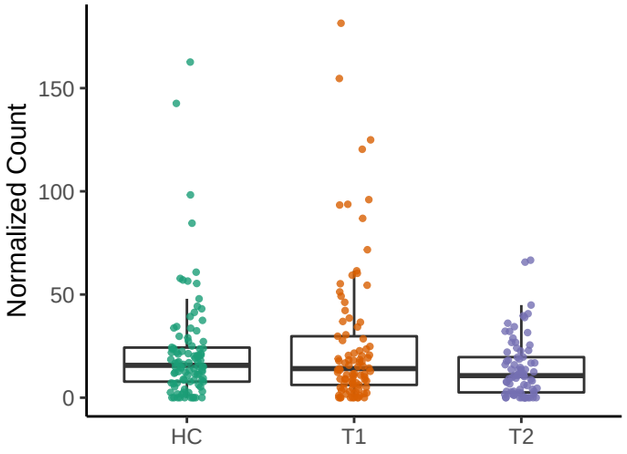
RUMP−PWY: formaldehyde oxidation I

HC vs. T1 adjusted p = 0.89
HC vs. T2 adjusted p = 0.052
T1 vs. T2 adjusted p = 0.2



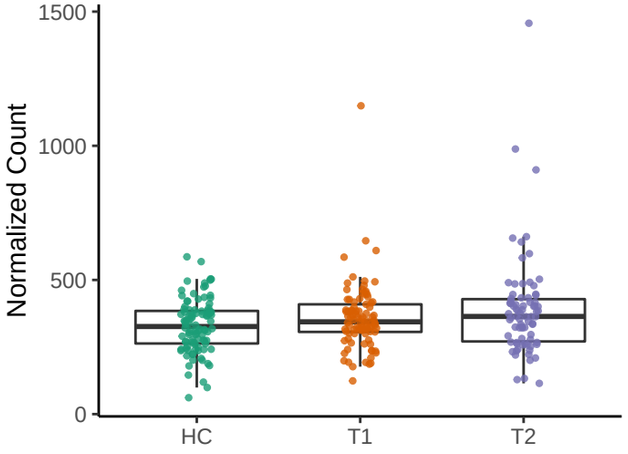
THISYN−PWY: superpathway of thiami

HC vs. T1 adjusted p = 0.5
HC vs. T2 adjusted p = 0.052
T1 vs. T2 adjusted p = 0.019



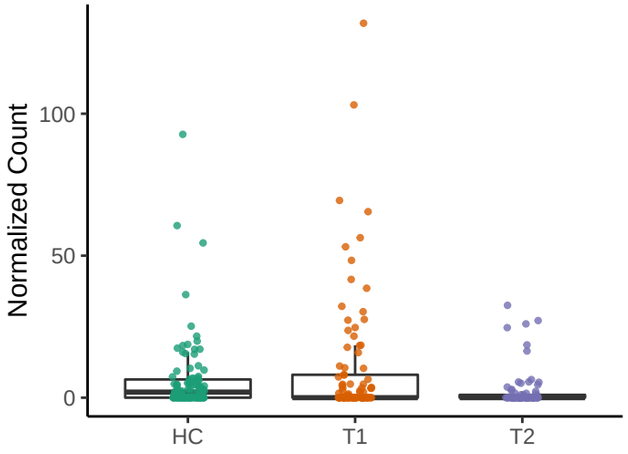
PANTO−PWY: phosphopantothenate b

HC vs. T1 adjusted p = 0.14
HC vs. T2 adjusted p = 0.054
T1 vs. T2 adjusted p = 0.5

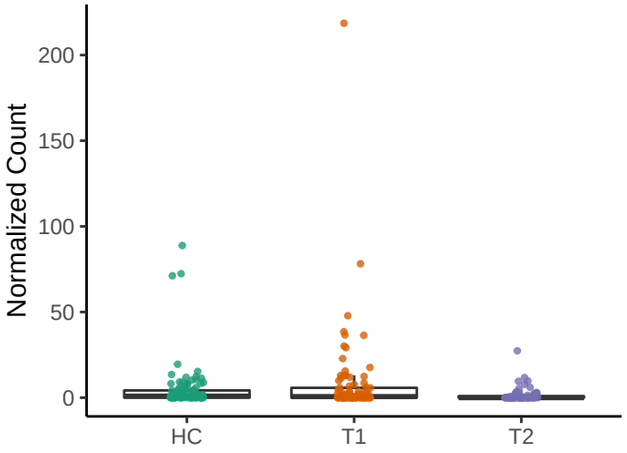


PWY−5838: superpathway of menaquin

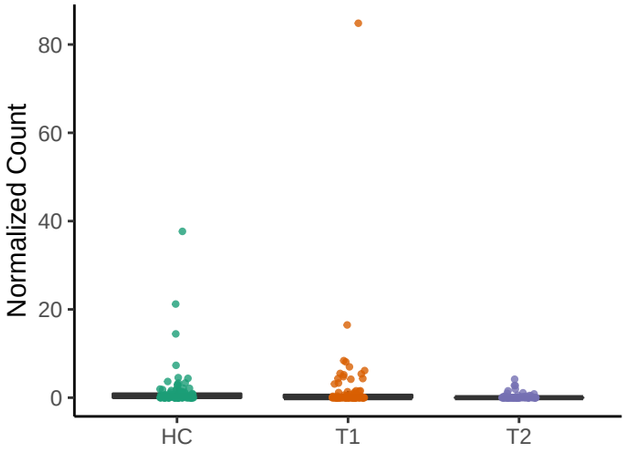
HC vs. T1 adjusted p = 0.23
HC vs. T2 adjusted p = 0.054
T1 vs. T2 adjusted p = 0.016



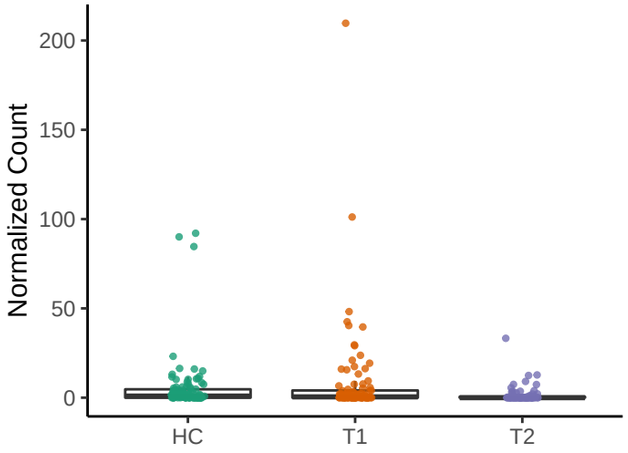
GLYOXYLATE−BYPASS: glyoxylate cyc
HC vs. T1 adjusted p = 0.45
HC vs. T2 adjusted p = 0.056
T1 vs. T2 adjusted p = 0.018



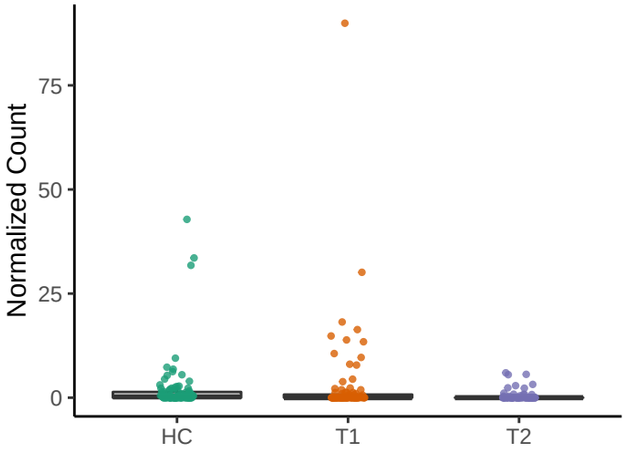
PWY−5705: allantoin degradation to gly
HC vs. T1 adjusted p = 0.69
HC vs. T2 adjusted p = 0.057
T1 vs. T2 adjusted p = 0.023



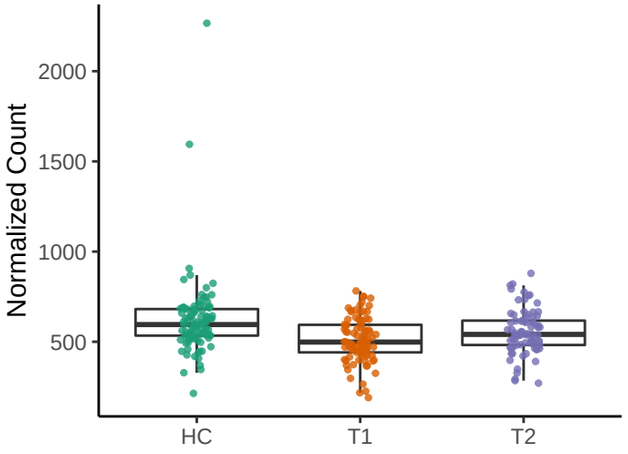
TCA−GLYOX−BYPASS: superpathway
HC vs. T1 adjusted p = 0.5
HC vs. T2 adjusted p = 0.06
T1 vs. T2 adjusted p = 0.021



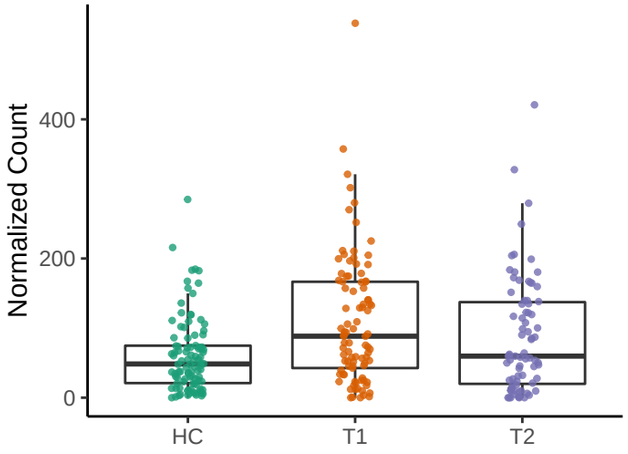
PWY−2723: trehalose degradation V
HC vs. T1 adjusted p = 0.65
HC vs. T2 adjusted p = 0.064
T1 vs. T2 adjusted p = 0.027



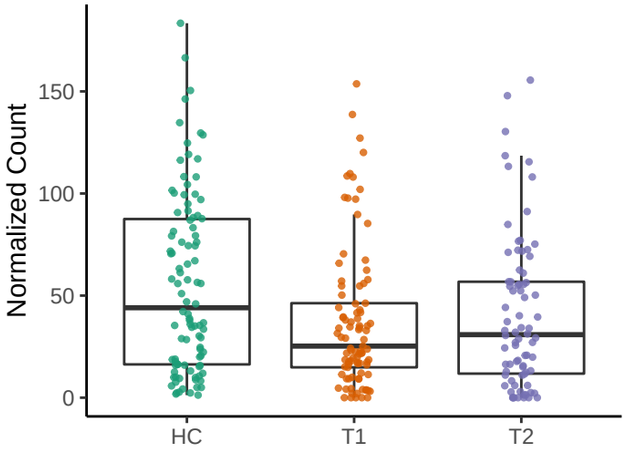
NONMEVIPP−PWY: methylerythritol p
HC vs. T1 adjusted p = 0.00025
HC vs. T2 adjusted p = 0.065
T1 vs. T2 adjusted p = 0.047



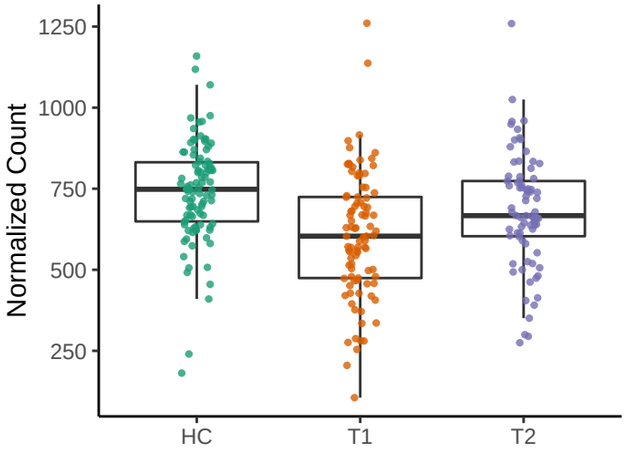
PYRIDOXSYN−PWY: pyridoxal 5'−phos
HC vs. T1 adjusted p = 0.00011
HC vs. T2 adjusted p = 0.067
T1 vs. T2 adjusted p = 0.042



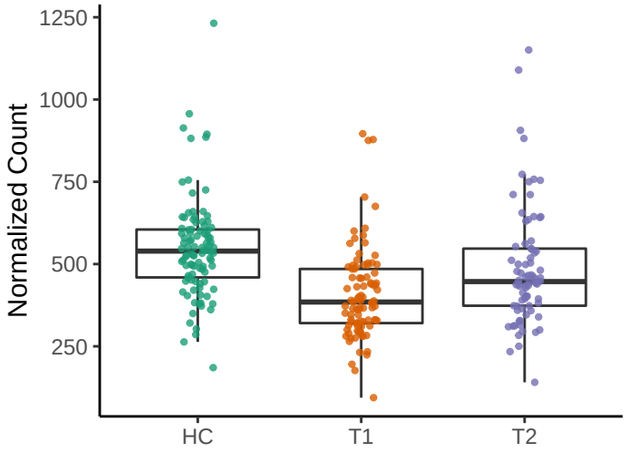
PYRIDNUCSAL−PWY: NAD salvage pa
HC vs. T1 adjusted p = 0.0067
HC vs. T2 adjusted p = 0.068
T1 vs. T2 adjusted p = 0.5



PWY−6737: starch degradation V
HC vs. T1 adjusted p = 3.7e−06
HC vs. T2 adjusted p = 0.073
T1 vs. T2 adjusted p = 0.019

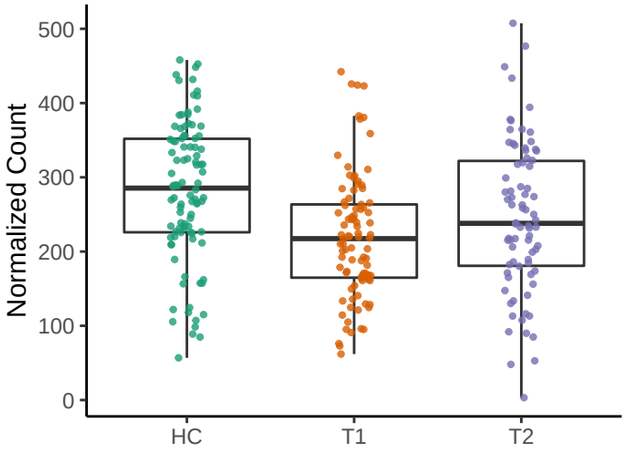


BRANCHED−CHAIN−AA−SYN−PWY:
HC vs. T1 adjusted p = 9.2e−09
HC vs. T2 adjusted p = 0.075
T1 vs. T2 adjusted p = 0.0045



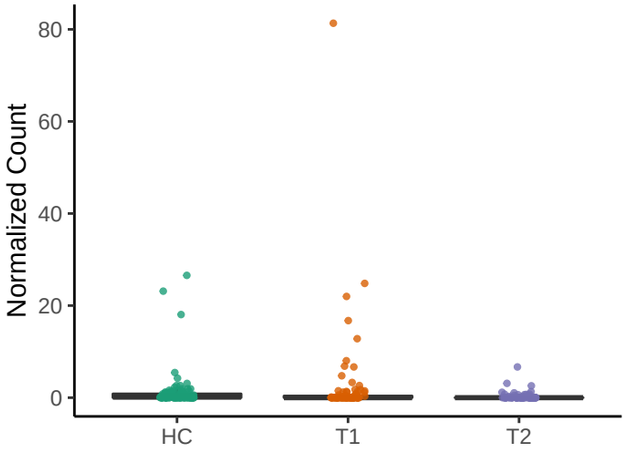
PWY−6527: stachyose degradation

HC vs. T1 adjusted p = 2.8e−05
HC vs. T2 adjusted p = 0.076
T1 vs. T2 adjusted p = 0.13



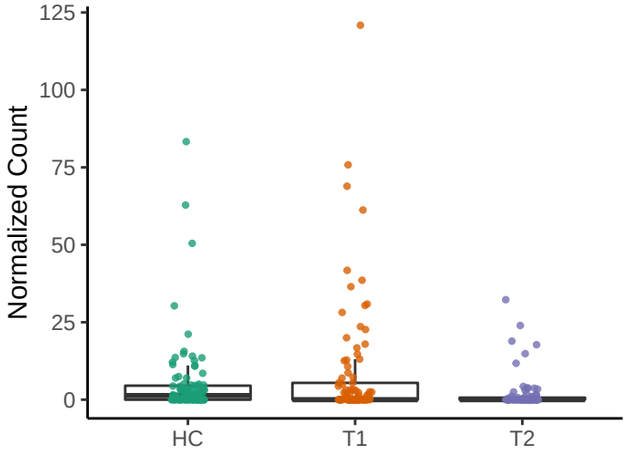
METHGLYUT−PWY: superpathway of m

HC vs. T1 adjusted p = 0.47
HC vs. T2 adjusted p = 0.079
T1 vs. T2 adjusted p = 0.05



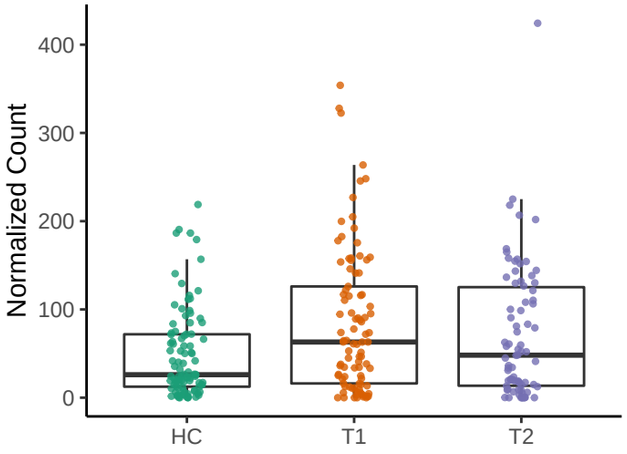
PWY−5861: superpathway of demethyl

HC vs. T1 adjusted p = 0.29
HC vs. T2 adjusted p = 0.086
T1 vs. T2 adjusted p = 0.018



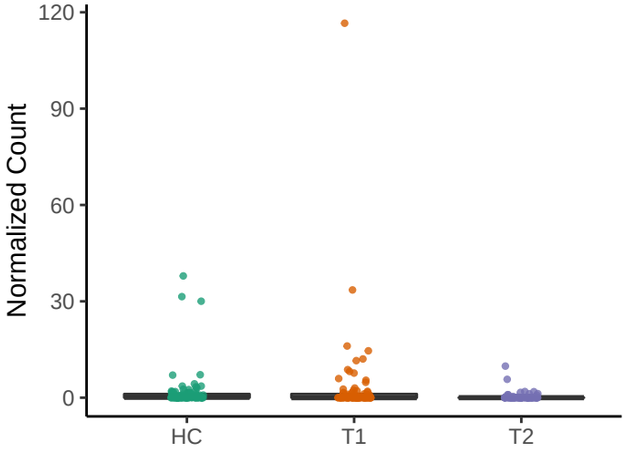
ARGININE−SYN4−PWY: L−ornithine d

HC vs. T1 adjusted p = 0.0015
HC vs. T2 adjusted p = 0.089
T1 vs. T2 adjusted p = 0.12



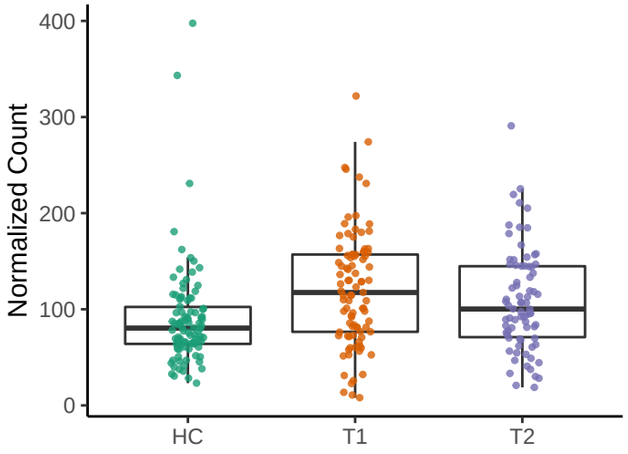
GLYCOL−GLYOXDEG−PWY: superpath

HC vs. T1 adjusted p = 0.53
HC vs. T2 adjusted p = 0.089
T1 vs. T2 adjusted p = 0.03



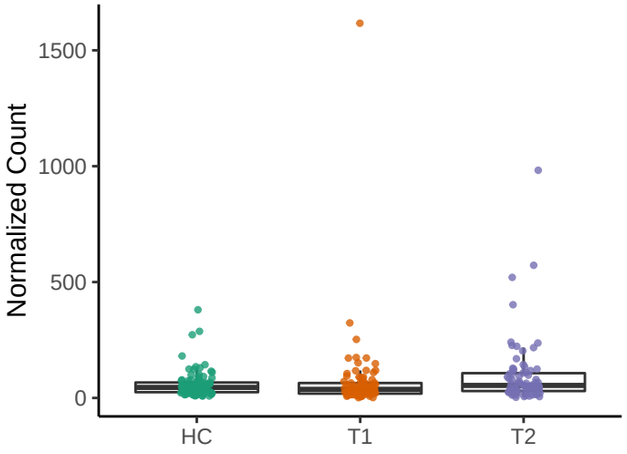
PWY−7197: pyrimidine deoxyribonucle

HC vs. T1 adjusted p = 0.0011
HC vs. T2 adjusted p = 0.089
T1 vs. T2 adjusted p = 0.15



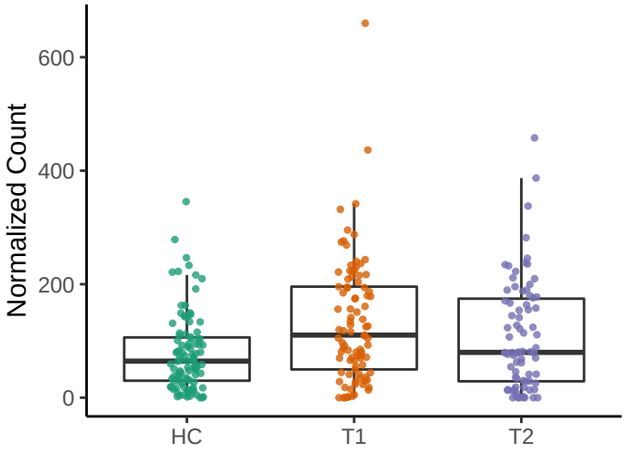
PWY−7237: myo−, chiro− and scillo−ir

HC vs. T1 adjusted p = 0.68
HC vs. T2 adjusted p = 0.089
T1 vs. T2 adjusted p = 0.38



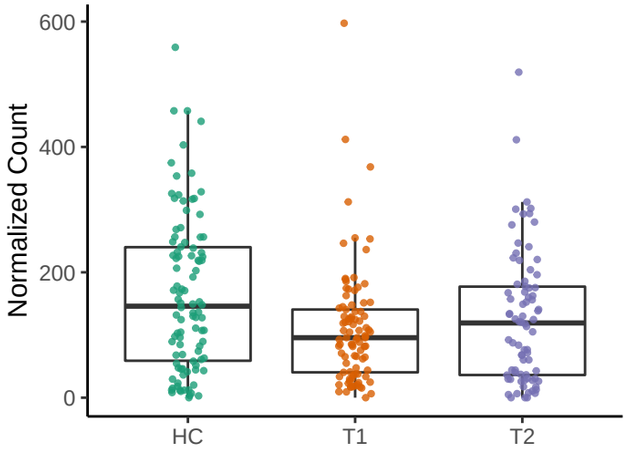
PWY0−845: superpathway of pyridoxal

HC vs. T1 adjusted p = 0.00046
HC vs. T2 adjusted p = 0.09
T1 vs. T2 adjusted p = 0.063



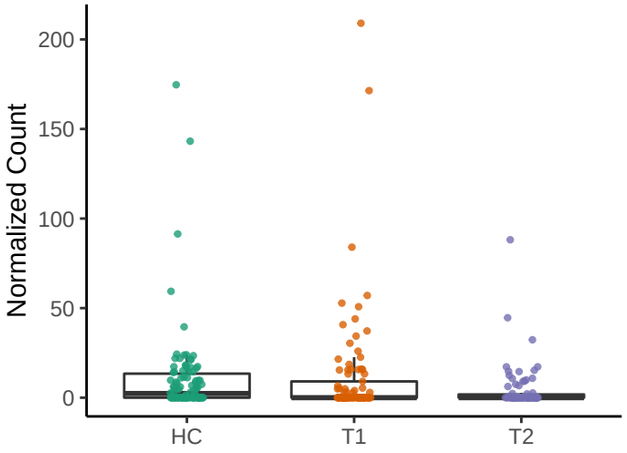
OANTIGEN−PWY: O−antigen building

HC vs. T1 adjusted p = 0.0026
HC vs. T2 adjusted p = 0.091
T1 vs. T2 adjusted p = 0.14



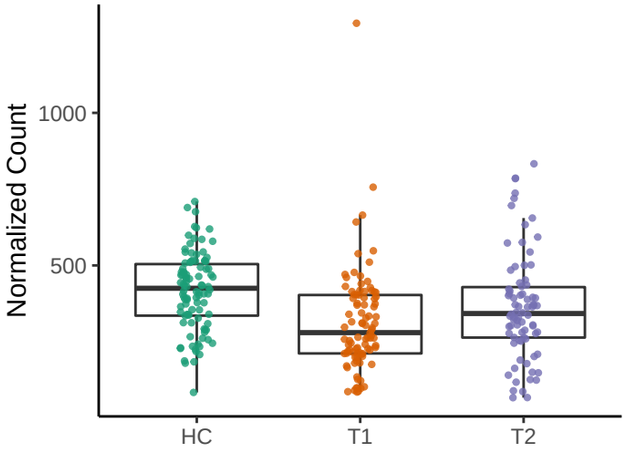
ENTBACSYN−PWY: enterobactin biosy

HC vs. T1 adjusted p = 0.89
HC vs. T2 adjusted p = 0.093
T1 vs. T2 adjusted p = 0.1



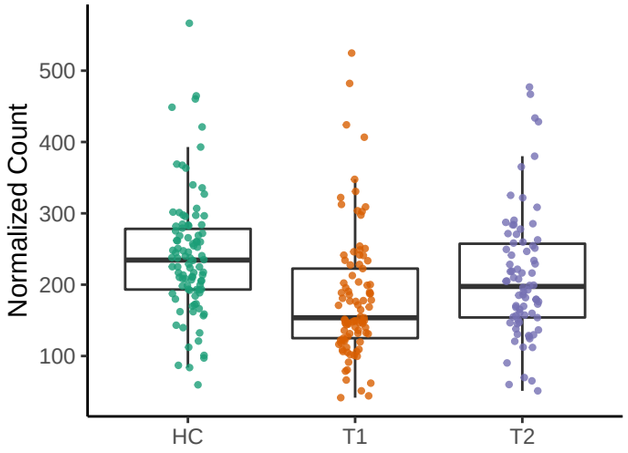
NONOXIPENT−PWY: pentose phosph

HC vs. T1 adjusted p = 3.5e−05
HC vs. T2 adjusted p = 0.093
T1 vs. T2 adjusted p = 0.022



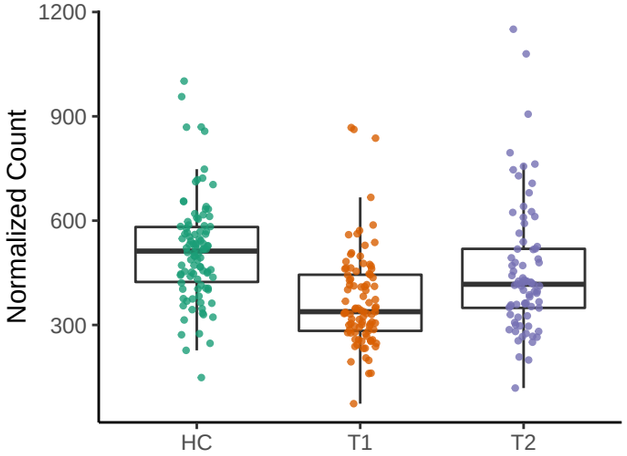
PWY−5100: pyruvate fermentation to ac

HC vs. T1 adjusted p = 4.7e−05
HC vs. T2 adjusted p = 0.093
T1 vs. T2 adjusted p = 0.05



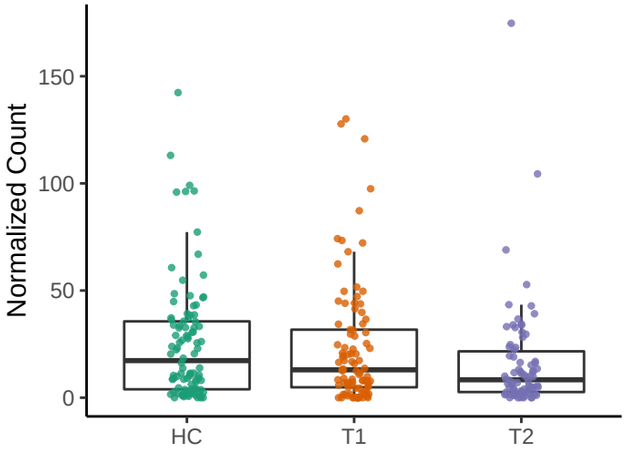
PWY−5103: L−isoleucine biosynthesis

HC vs. T1 adjusted p = 5.5e−09
HC vs. T2 adjusted p = 0.093
T1 vs. T2 adjusted p = 0.0045



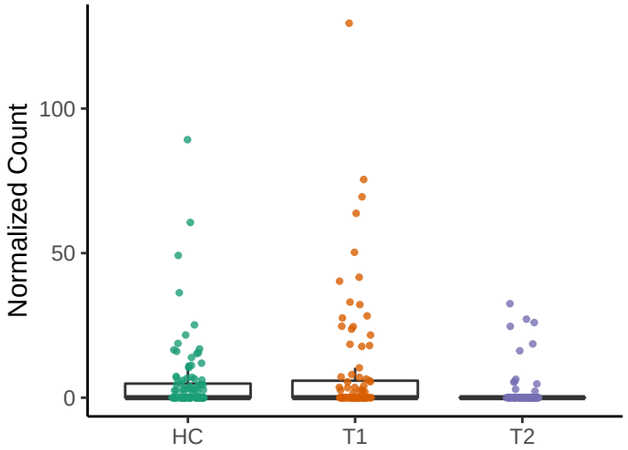
PWY−5345: superpathway of L−methio

HC vs. T1 adjusted p = 0.77
HC vs. T2 adjusted p = 0.093
T1 vs. T2 adjusted p = 0.17



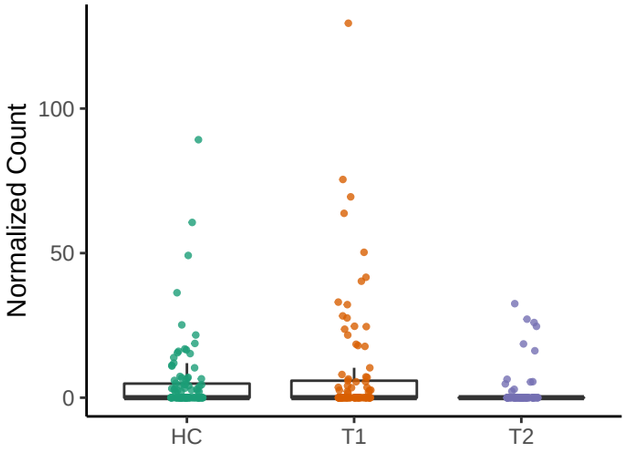
PWY−5845: superpathway of menaquin

HC vs. T1 adjusted p = 0.3
HC vs. T2 adjusted p = 0.093
T1 vs. T2 adjusted p = 0.018



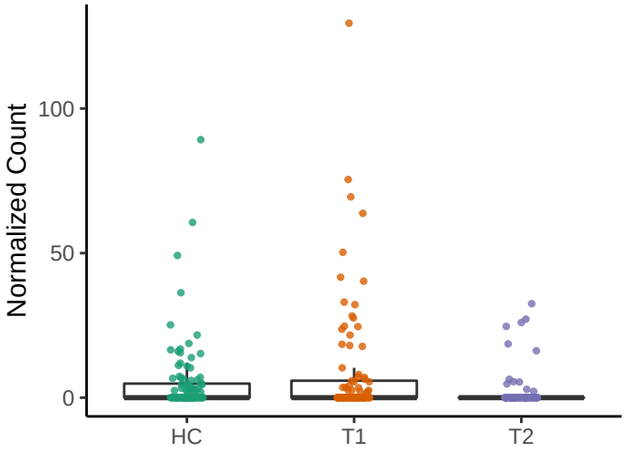
PWY−5850: superpathway of menaquir

HC vs. T1 adjusted p = 0.3
HC vs. T2 adjusted p = 0.093
T1 vs. T2 adjusted p = 0.018



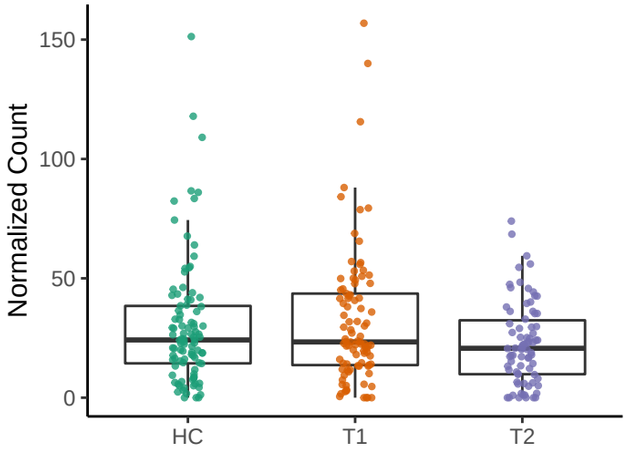
PWY−5896: superpathway of menaquir

HC vs. T1 adjusted p = 0.3
HC vs. T2 adjusted p = 0.093
T1 vs. T2 adjusted p = 0.018

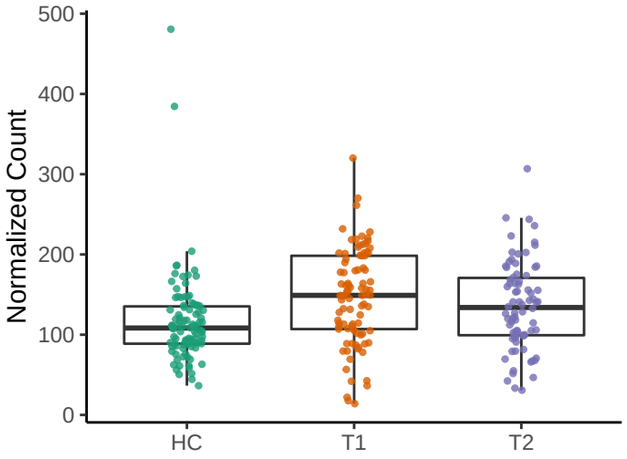


PWY−6901: superpathway of glucose a

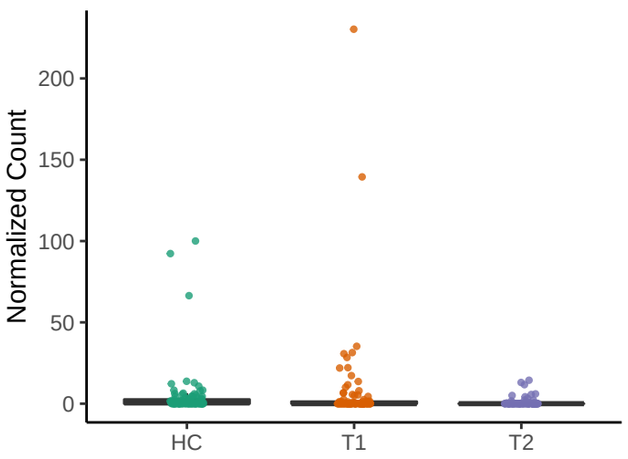
HC vs. T1 adjusted p = 0.68
HC vs. T2 adjusted p = 0.093
T1 vs. T2 adjusted p = 0.021



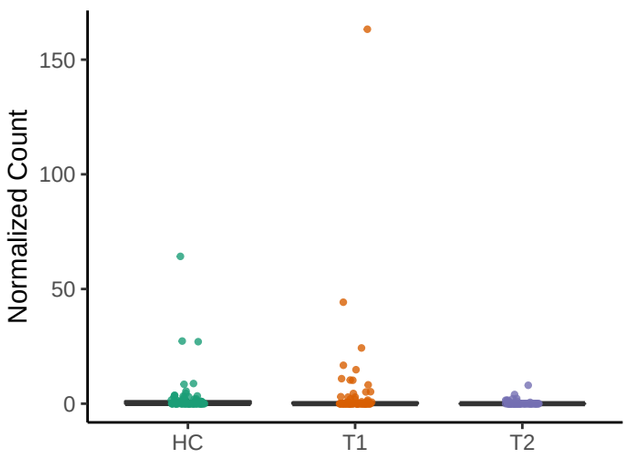
PWY−7184: pyrimidine deoxyribonucle(
HC vs. T1 adjusted p = 0.0017
HC vs. T2 adjusted p = 0.093
T1 vs. T2 adjusted p = 0.14



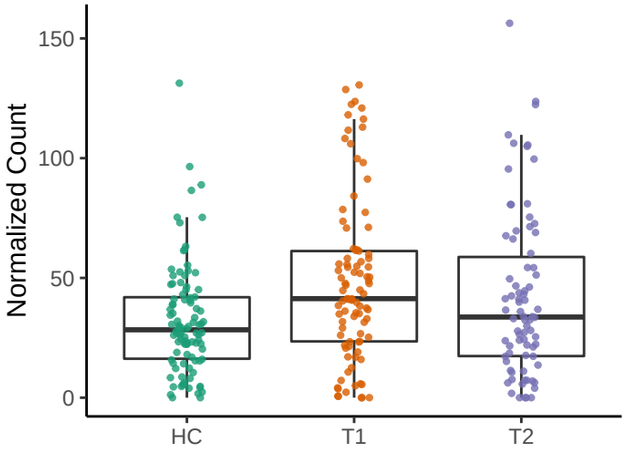
PWY−7315: dTDP−N−acetylthomosam
HC vs. T1 adjusted p = 0.54
HC vs. T2 adjusted p = 0.093
T1 vs. T2 adjusted p = 0.08



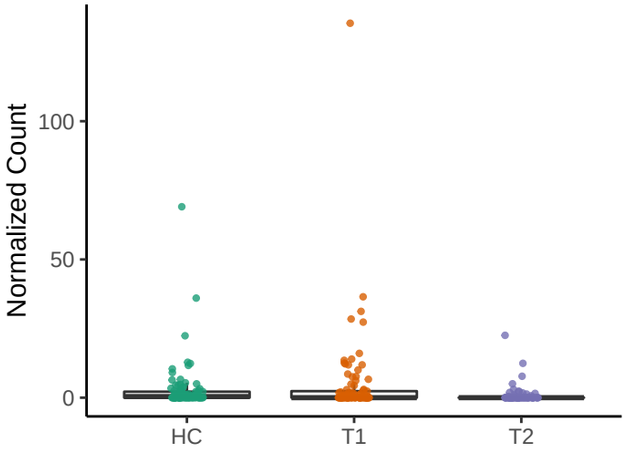
PWY−7409: phospholipid remodeling (p
HC vs. T1 adjusted p = 0.53
HC vs. T2 adjusted p = 0.093
T1 vs. T2 adjusted p = 0.044



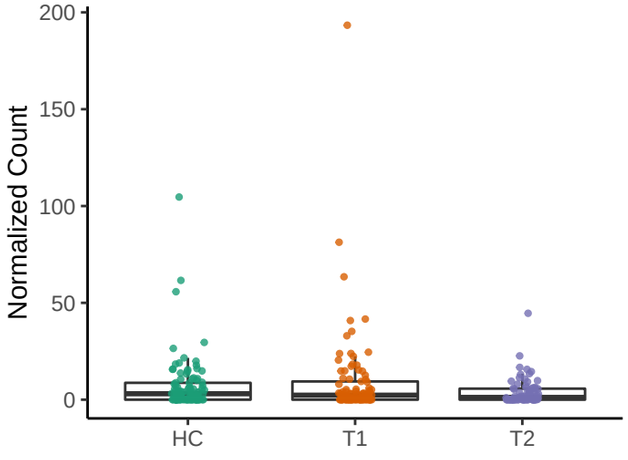
PWY66−399: gluconeogenesis III
HC vs. T1 adjusted p = 0.00074
HC vs. T2 adjusted p = 0.093
T1 vs. T2 adjusted p = 0.11



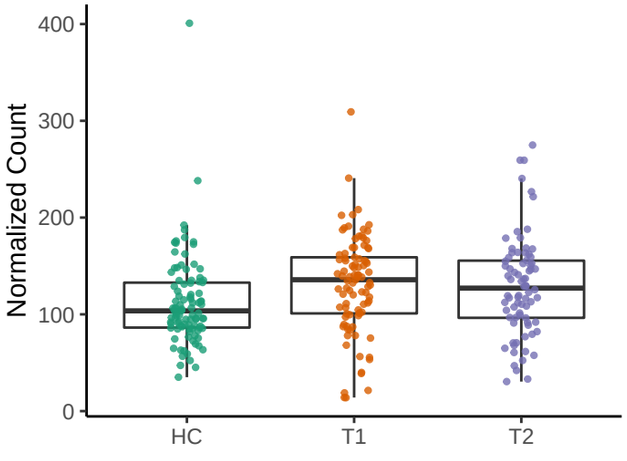
PWY−6803: phosphatidylcholine acyl e
HC vs. T1 adjusted p = 0.44
HC vs. T2 adjusted p = 0.095
T1 vs. T2 adjusted p = 0.019



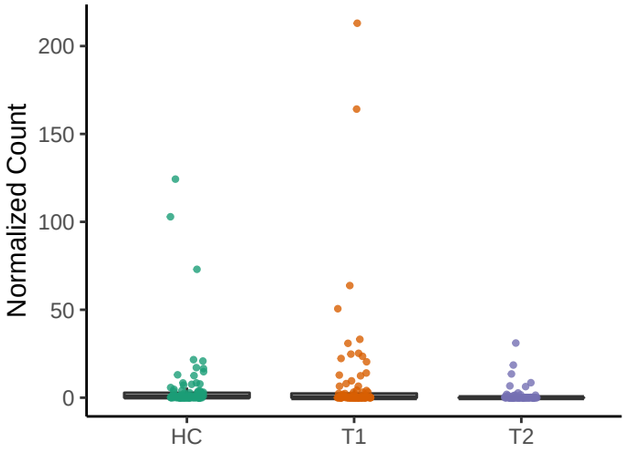
FUC−RHAMCAT−PWY: superpathway
HC vs. T1 adjusted p = 0.55
HC vs. T2 adjusted p = 0.096
T1 vs. T2 adjusted p = 0.042



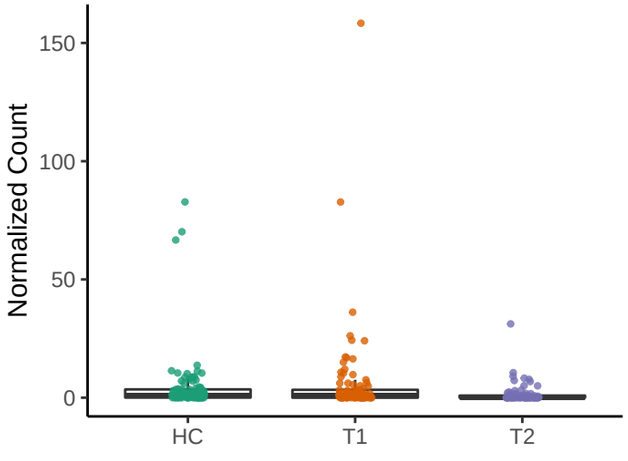
PWY−6545: pyrimidine deoxyribonucle(
HC vs. T1 adjusted p = 0.028
HC vs. T2 adjusted p = 0.096
T1 vs. T2 adjusted p = 0.61



PWY−6891: thiazole biosynthesis II (Ba
HC vs. T1 adjusted p = 0.51
HC vs. T2 adjusted p = 0.096
T1 vs. T2 adjusted p = 0.063

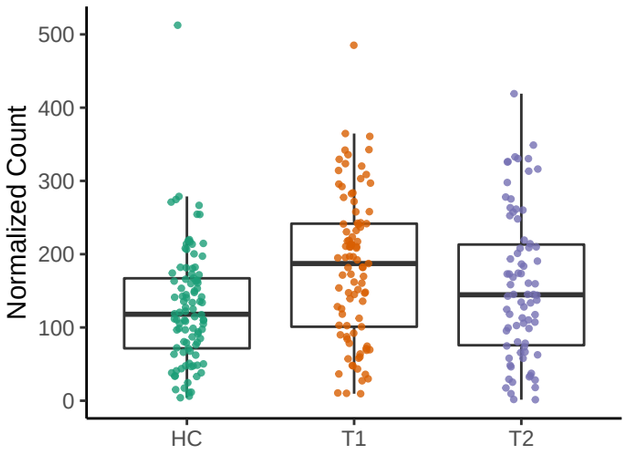


PWY−7254: TCA cycle VII (acetate−pro
HC vs. T1 adjusted p = 0.6
HC vs. T2 adjusted p = 0.096
T1 vs. T2 adjusted p = 0.05



PWY−7663: gondoate biosynthesis (an

HC vs. T1 adjusted p = 0.00022
HC vs. T2 adjusted p = 0.096
T1 vs. T2 adjusted p = 0.021



P161−PWY: acetylene degradation

HC vs. T1 adjusted p = 0.0098
HC vs. T2 adjusted p = 0.1
T1 vs. T2 adjusted p = 0.45