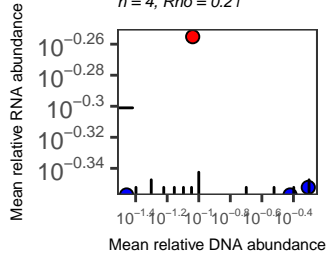
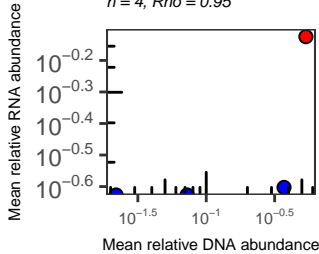


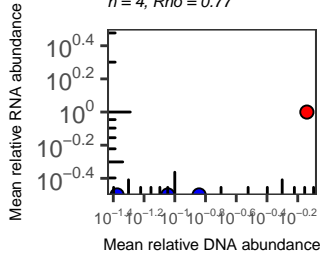
FUCCAT-PWY
fucose
degradation
 $n = 4, Rho = 0.21$



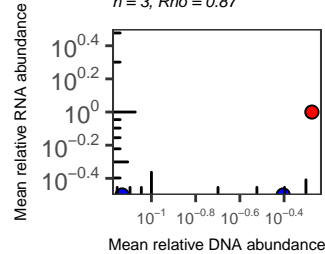
PWY-5384
sucrose
degradation IV (sucrose phospho)
 $n = 4, Rho = 0.95$



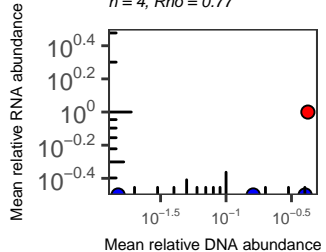
TCA-GLYOX-BYPASS
superpathway
of glyoxylate bypass and TCA
 $n = 4, Rho = 0.77$



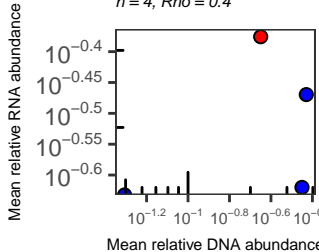
PWY-6270
isoprene
biosynthesis I
 $n = 3, Rho = 0.87$



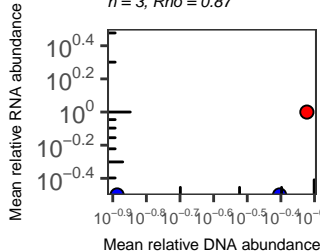
GLUCOSE1PMETAB-PWY
glucose
and glucose-1-phosphate degi
 $n = 4, Rho = 0.77$



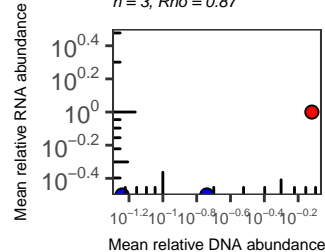
PWY-7456
mannan
degradation
 $n = 4, Rho = 0.4$



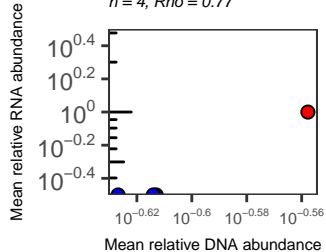
HCAMHPDEG-PWY
3-phenylpropanoate
and 3-(3-hydroxyphenyl)propa
 $n = 3, Rho = 0.87$



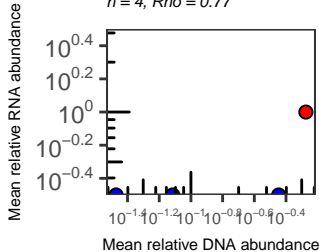
PWY-6612
superpathway
of tetrahydrofolate biosynthesis
 $n = 3, Rho = 0.87$



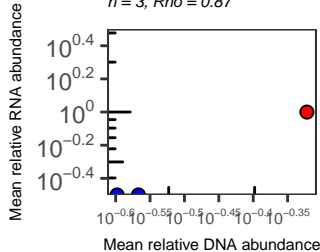
PWY-4984
urea
cycle
 $n = 4, Rho = 0.77$



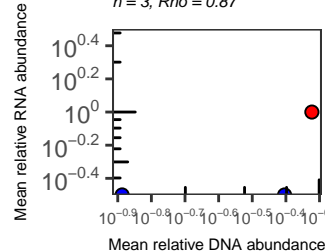
PWY-7560
methylerythritol
phosphate pathway II
 $n = 4, Rho = 0.77$



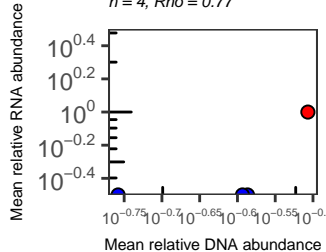
PWY-3781
aerobic
respiration I (cytochrome c)
 $n = 3, Rho = 0.87$



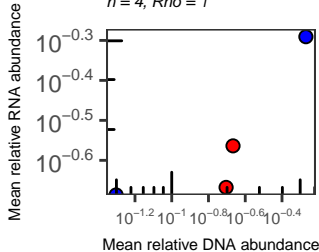
PWY-6690
cinnamate
and 3-hydroxycinnamate degra
 $n = 3, Rho = 0.87$



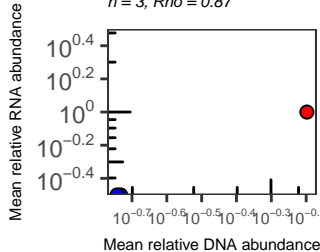
PWY-5138
unsaturated,
even numbered fatty acid & beta
 $n = 4, Rho = 0.77$



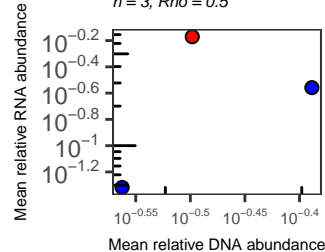
PWY0-1479
tRNA
processing
 $n = 4, Rho = 1$



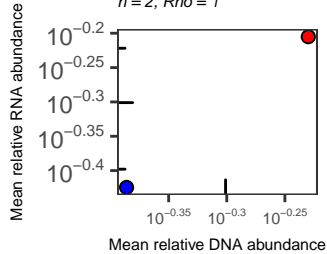
PWY-5136
fatty
acid & beta;-oxidation II (peroxi
 $n = 3, Rho = 0.87$



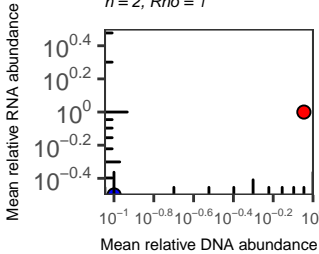
SALVADEHYPOX-PWY
adenosine
nucleotides degradation II
 $n = 3, Rho = 0.5$



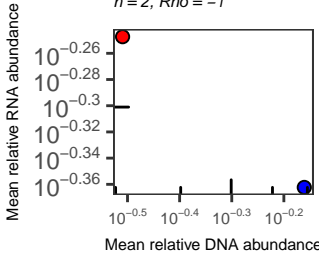
7ALPHADEHYDROX-PWY
cholate degradation (bacteria, anaerol
n = 2, Rho = 1



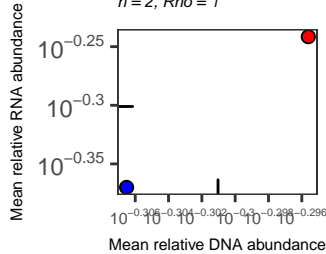
GLYCOL-GLYOXDEG-PWY
superpathway of glycol metabolism and degra
n = 2, Rho = 1



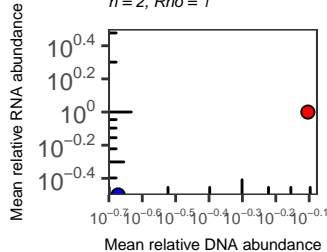
PWY-4702
phytate degradation I
n = 2, Rho = -1



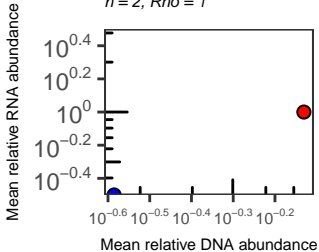
PWY-6588
pyruvate fermentation to acetone
n = 2, Rho = 1



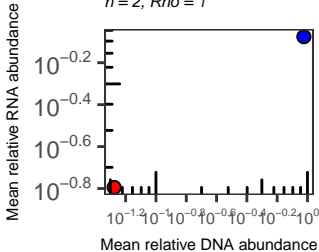
FOLSYN-PWY
superpathway of tetrahydrofolate biosynthesis
n = 2, Rho = 1



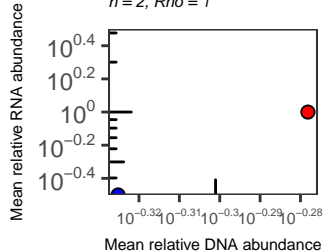
GLYCOLYSIS
glycolysis I (from glucose 6-phosphate)
n = 2, Rho = 1



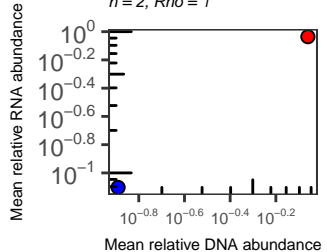
PWY-5101
L-isoleucine biosynthesis II
n = 2, Rho = 1



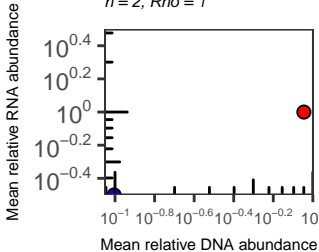
PWY-6630
superpathway of L-tyrosine biosynthesis
n = 2, Rho = 1



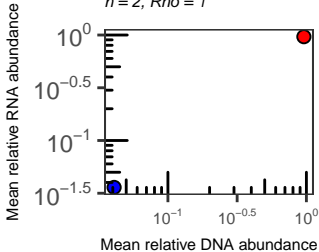
GALACT-GLUCUROCAT-PWY
superpathway of hexuronide and hexuronate c
n = 2, Rho = 1



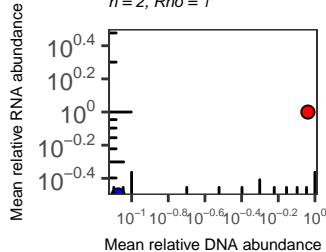
METHANOGENESIS-PWY
methanogenesis from H2 and CO2
n = 2, Rho = 1



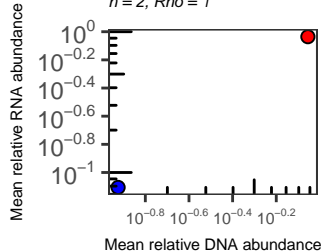
PWY-5177
glutaryl-CoA degradation
n = 2, Rho = 1



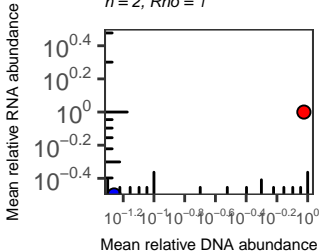
PWY-7269
NAD/NADP-NADH/NADPH mitochondrial interconversion (y
n = 2, Rho = 1



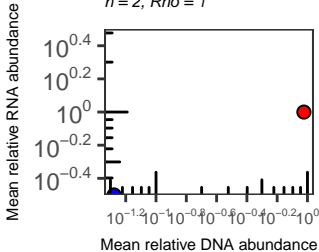
GLUCUROCAT-PWY
superpathway of β-D-glucuronide and D
n = 2, Rho = 1



PWY-2723
trehalose degradation V
n = 2, Rho = 1



PWY-5198
factor 420 biosynthesis
n = 2, Rho = 1



PWY-7315
dTDP-N-acetylthomosamine biosynthesis
n = 2, Rho = 1

