

Category	Term
GOTERM_BP_DIRECT	GO:0055114~oxidation-reduction process
GOTERM_BP_DIRECT	GO:0006730~one-carbon metabolic process
GOTERM_BP_DIRECT	GO:0005975~carbohydrate metabolic process
GOTERM_BP_DIRECT	GO:0006635~fatty acid beta-oxidation
GOTERM_BP_DIRECT	GO:0006189~'de novo' IMP biosynthetic process
GOTERM_BP_DIRECT	GO:0033539~fatty acid beta-oxidation using acyl-CoA dehydrogenase
GOTERM_BP_DIRECT	GO:0006098~pentose-phosphate shunt
GOTERM_BP_DIRECT	GO:0055088~lipid homeostasis
GOTERM_BP_DIRECT	GO:0035999~tetrahydrofolate interconversion
GOTERM_BP_DIRECT	GO:0055085~transmembrane transport
GOTERM_BP_DIRECT	GO:0006164~purine nucleotide biosynthetic process
GOTERM_BP_DIRECT	GO:0008152~metabolic process
GOTERM_BP_DIRECT	GO:0006090~pyruvate metabolic process
GOTERM_BP_DIRECT	GO:0006006~glucose metabolic process
GOTERM_BP_DIRECT	GO:0006552~leucine catabolic process
GOTERM_BP_DIRECT	GO:0009083~branched-chain amino acid catabolic process
GOTERM_BP_DIRECT	GO:0006574~valine catabolic process
GOTERM_BP_DIRECT	GO:0006568~tryptophan metabolic process
GOTERM_BP_DIRECT	GO:0046323~glucose import
GOTERM_BP_DIRECT	GO:0035428~hexose transmembrane transport
GOTERM_BP_DIRECT	GO:0006551~leucine metabolic process
GOTERM_BP_DIRECT	GO:0015718~monocarboxylic acid transport
GOTERM_BP_DIRECT	GO:0009396~folic acid-containing compound biosynthetic process
GOTERM_BP_DIRECT	GO:0006567~threonine catabolic process
GOTERM_BP_DIRECT	GO:0045454~cell redox homeostasis
GOTERM_BP_DIRECT	GO:0006206~pyrimidine nucleobase metabolic process
GOTERM_BP_DIRECT	GO:0001561~fatty acid alpha-oxidation
GOTERM_BP_DIRECT	GO:0030431~sleep
GOTERM_BP_DIRECT	GO:0006520~cellular amino acid metabolic process
GOTERM_BP_DIRECT	GO:0006032~chitin catabolic process
GOTERM_BP_DIRECT	GO:0033499~galactose catabolic process via UDP-galactose
GOTERM_BP_DIRECT	GO:0019318~hexose metabolic process
GOTERM_BP_DIRECT	GO:0006096~glycolytic process

Count	%	PValue
41	19.1588785	1.92E-18
7	3.27102804	1.38E-06
12	5.60747664	4.74E-06
7	3.27102804	1.47E-05
4	1.86915888	1.31E-04
5	2.3364486	1.49E-04
4	1.86915888	2.26E-04
6	2.80373832	2.72E-04
3	1.40186916	0.00107342
14	6.54205607	0.00297479
3	1.40186916	0.00516756
8	3.73831776	0.01959286
3	1.40186916	0.02461484
3	1.40186916	0.03648275
2	0.93457944	0.03781367
2	0.93457944	0.03781367
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3	1.40186916	0.05497774
2	0.93457944	0.05618379
2	0.93457944	0.05618379
2	0.93457944	0.05618379
2	0.93457944	0.05618379
4	1.86915888	0.07320319
2	0.93457944	0.07420511
2	0.93457944	0.07420511
6	2.80373832	0.07945898
3	1.40186916	0.08164158
3	1.40186916	0.08164158
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2	0.93457944	0.09188419
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Genes

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