
Heatmap enrichment p−value (−log10)
establishment or maintenance of cell polarity
response to copper ion
RNA import into nucleus
establishment or maintenance of actin cytoskeleton polarity
protein import into mitochondrial outer membrane
protein localization
protein targeting
double−strand break repair via nonhomologous end joining
tricarboxylic acid cycle
regulation of nucleocytoplasmic transport
regulation of G1/S transition of mitotic cell cycle
regulation of establishment of protein localization
mRNA destabilization
response to furfural
amide transport
biological regulation
protein transport
regulation of cell growth
NAD biosynthetic process
positive regulation of cell cycle G1/S phase transition
aging
cellular developmental process
positive regulation of mRNA metabolic process
regulation of mRNA catabolic process
replicative cell aging
small GTPase mediated signal transduction
regulation of biological process
regulation of RNA stability
Rab protein signal transduction
traversing start control point of mitotic cell cycle
tricarboxylic acid metabolic process
cell morphogenesis
iron ion transport
sequestering of metal ion
intracellular sequestering of iron ion
chromatin remodeling at centromere
chitin catabolic process
TORC2 signaling
amino sugar catabolic process
organelle organization
positive regulation of mitotic cell cycle phase transition
allantoin catabolic process
cardiolipin metabolic process
oligosaccharide catabolic process
carbohydrate metabolic process
maltose catabolic process
triglyceride mobilization
endoplasmic reticulum inheritance
drug metabolic process
reactive oxygen species metabolic process
ribonucleoside diphosphate metabolic process
ADP metabolic process
purine nucleoside diphosphate metabolic process
sucrose catabolic process
regulation of transcription regulatory region DNA binding
coflocculation
glycerol transport
cell wall organization
nitrogen utilization
regulation of intracellular protein transport
positive regulation of cellular response to amino acid starvation
nucleotide transmembrane transport
nucleoside monophosphate phosphorylation
nuclear pore complex assembly
modified amino acid transport
fungal−type cell wall beta−glucan biosynthetic process
flocculation
endoplasmic reticulum tubular network organization
cell wall beta−glucan metabolic process
ER−dependent peroxisome organization
cell aggregation
fungal−type cell wall biogenesis
nucleoside diphosphate phosphorylation
gluconeogenesis
fungal−type cell wall polysaccharide metabolic process
cellular carbohydrate catabolic process
cellular response to nitrogen compound
disaccharide metabolic process
glycerolipid metabolic process
organophosphate biosynthetic process
cellular response to organic substance
glycolytic process
regulation of protein acetylation
phosphatidylethanolamine biosynthetic process
regulation of histone acetylation
nucleoside phosphate catabolic process
regulation of lipid metabolic process
2
1.5
1
0.5
0
Causal
Affected