Heatmap enrichment p−value (−log10)

oxoacid metabolic process
drug metabolic process
organonitrogen compound biosynthetic process
monocarboxylic acid metabolic process
ribose phosphate metabolic process
purine−containing compound metabolic process
carboxylic acid biosynthetic process
nucleoside phosphate biosynthetic process
cofactor metabolic process
nucleotide metabolic process
nucleobase−containing small molecule metabolic process
oxidation−reduction process
small molecule biosynthetic process
DNA integration
positive regulation of gene expression
regulation of primary metabolic process
positive regulation of macromolecule biosynthetic process
transcription, DNA−templated
regulation of cellular macromolecule biosynthetic process
regulation of biosynthetic process
RNA biosynthetic process
positive regulation of cellular biosynthetic process
DNA recombination
RNA−dependent DNA biosynthetic process
heterocycle biosynthetic process
regulation of biological process
aromatic compound biosynthetic process
regulation of cellular metabolic process
positive regulation of transcription by RNA polymerase II
proteolysis
cellular nitrogen compound biosynthetic process
positive regulation of nitrogen compound metabolic process
positive regulation of nucleic acid−templated transcription
organic cyclic compound biosynthetic process
positive regulation of RNA metabolic process
regulation of translational termination
cellular protein complex disassembly
maintenance of rDNA
lipid catabolic process
homologous chromosome segregation
meiotic nuclear division
response to stress
organelle disassembly
RNA metabolic process
negative regulation of RNA metabolic process
organic substance metabolic process
SCF−dependent proteasomal ubiquitin−dependent protein catabolic process
signal transduction
transposition, RNA−mediated
negative regulation of biosynthetic process
positive regulation of cellular process
cellular macromolecule metabolic process
nucleobase−containing compound metabolic process
negative regulation of cellular macromolecule biosynthetic process
nucleic acid phosphodiester bond hydrolysis
TOR signaling
mitotic cell cycle process
negative regulation of nucleic acid−templated transcription
RNA phosphodiester bond hydrolysis, endonucleolytic
cellular response to DNA damage stimulus
chromosome segregation

Causal   Affected