
rank breakdown
temperature homeostasis
cardiac conduction
negative regulation of cold-induced thermogenesis
cardiac muscle cell contraction
regulation of myelination
regulation of myeloid leukocyte differentiation
vasoconstriction
ensheathment of neurons
cardiac chamber development
mesenchymal cell differentiation
apoptotic process involved in development
positive regulation of myeloid cell differentiation
stem cell development
neuroblast proliferation
regulation of heart rate
digestive system process
T cell costimulation
energy homeostasis
positive regulation of axonogenesis
neuron maturation
regulation of vasoconstriction
regulation of glial cell differentiation
glycosylation
protein polyubiquitination
SCF-dependent proteasomal ubiquitin-dependent protein catabolic process
positive regulation of protein modification by small protein conjugation or removal
regulation of peptidyl-serine phosphorylation
protein monoubiquitination
peptidyl-serine modification
peptidyl-lysine trimethylation
signaling receptor ligand precursor processing
phospholipid catabolic process
positive regulation of MAP kinase activity
tRNA methylation
regulation of histone methylation
negative regulation of peptidyl-tyrosine phosphorylation
RNA-templated DNA biosynthetic process
monosaccharide metabolic process
carbohydrate biosynthetic process
carbohydrate catabolic process
carbohydrate homeostasis
energy homeostasis
dicarboxylic acid metabolic process
regulation of nervous system process
regulation of striated muscle contraction
muscle organ development
ventricular cardiac muscle tissue development
cardiac muscle tissue morphogenesis
energy homeostasis
anatomical structure homeostasis
regulation of digestive system process
logpval
0.0
−0.5
−1.0
−1.5
−2.0
−2.5
−3.0
−3.5