

GObp in module 1.2.3.1.3.1.2

Term

positive regulation of protein deacetylation
mRNA modification
late endosome to vacuole transport
mitophagy
negative regulation of necroptotic process
regulation of histone deacetylation
kinetochore organization
protein mannosylation
acetyl-CoA biosynthetic process
intracellular protein transmembrane transport
regulation of TORC1 signaling
protein K48-linked ubiquitination
GPI anchor biosynthetic process
positive regulation of mRNA catabolic process
protein K63-linked ubiquitination
mitotic metaphase plate congression
regulation of centrosome duplication
regulation of macroautophagy
protein deubiquitination

0

1

2

log2Enrichment

GOcc in module 1.2.3.1.3.1.2

Term

endoribonuclease activity, producing 5'-phosphomonoesters
Rab guanyl-nucleotide exchange factor activity
4 iron, 4 sulfur cluster binding
protein kinase inhibitor activity
thiol-dependent ubiquitin-specific protease activity
S-acyltransferase activity

0.0

0.5

1.0

1.5

2.0

log2Enrichment

GOfm in module 1.2.3.1.3.1.2

Term

retromer complex
INO80-type complex
integrator complex
ciliary base
histone deacetylase complex

0

1

2

log2Enrichment