

GO_Biological_Process_2021

Term

negative regulation of metallopeptidase
activity (GO:1905049)

nucleotide-sugar metabolic process
(GO:0009225)

membrane fission (GO:0090148)

embryo development ending in birth or egg
hatching (GO:0009792)

mitochondrial fragmentation involved in
apoptotic process (GO:0043653)

positive regulation of production of miRNAs
involved in gene silencing by miRNA
(GO:1903800)

cardiac muscle cell membrane repolarization
(GO:0099622)

regulation of membrane protein ectodomain
proteolysis (GO:0051043)

positive regulation of mitotic sister
chromatid separation (GO:1901970)

positive regulation of mitotic
metaphase/anaphase transition (GO:0045842)

positive regulation of metaphase/anaphase
transition of cell cycle (GO:1902101)

0

2

4

6

Enrichment log(combined score)

GO_Cellular_Component_2021

Term

kinetochore microtubule (GO:0005828)

anaphase-promoting complex (GO:0005680)

neurofibrillary tangle (GO:0097418)

NatA complex (GO:0031415)

postsynaptic recycling endosome (GO:0098837)

Cul2-RING ubiquitin ligase complex
(GO:0031462)

pinosome (GO:0044352)

macropinosome (GO:0044354)

NSL complex (GO:0044545)

extrinsic component of external side of
plasma membrane (GO:0031232)

dendritic spine membrane (GO:0032591)

MICOS complex (GO:0061617)

HFE-transferrin receptor complex
(GO:1990712)

0

2

4

6

Enrichment log(combined score)

GO_Molecular_Function_2021

Term

3',5'-cyclic-GMP phosphodiesterase activity
(GO:0047555)

microtubule plus-end binding (GO:0051010)

metalloendopeptidase inhibitor activity
(GO:0008191)

cytochrome-c oxidase activity (GO:0004129)

voltage-gated calcium channel activity
involved in cardiac muscle cell action
potential (GO:0086007)

phosphotransferase activity, nitrogenous
group as acceptor (GO:0016775)

creatine kinase activity (GO:0004111)

choline transmembrane transporter activity
(GO:0015220)

oxidoreductase activity, acting on the CH-CH
group of donors, NAD or NADP as acceptor
(GO:0016628)

primary miRNA binding (GO:0070878)

CoA carboxylase activity (GO:0016421)

0

2

4

6

Enrichment log(combined score)