

GO_Biological_Process_2021

Term

zymogen inhibition (GO:0097341)

inhibition of cysteine-type endopeptidase
activity (GO:0097340)

positive regulation of endoplasmic reticulum
unfolded protein response (GO:1900103)

pentose-phosphate shunt (GO:0006098)

negative regulation of binding (GO:0051100)

T cell extravasation (GO:0072683)

ribose phosphate metabolic process
(GO:0019693)

regulation of type B pancreatic cell
apoptotic process (GO:2000674)

regulation of adherens junction organization
(GO:1903391)

pyrimidine deoxyribonucleotide biosynthetic
process (GO:0009221)

positive regulation of neutrophil
extravasation (GO:2000391)

positive regulation of mast cell chemotaxis
(GO:0060754)

negative regulation of plasminogen
activation (GO:0010757)

0

2

4

6

Enrichment log(combined score)

GO_Cellular_Component_2021

Term

spermatoproteasome complex (GO:1990111)

phosphatidylinositol 3-kinase complex, class I (GO:0097651)

early endosome lumen (GO:0031905)

shelterin complex (GO:0070187)

cis-Golgi network (GO:0005801)

cis-Golgi network membrane (GO:0033106)

cortical actin cytoskeleton (GO:0030864)

insulin-responsive compartment (GO:0032593)

sodium:potassium-exchanging ATPase complex (GO:0005890)

cortical cytoskeleton (GO:0030863)

0

2

4

Enrichment log(combined score)

GO_Molecular_Function_2021

Term

calcium-dependent cysteine-type
endopeptidase activity (GO:0004198)

lipase binding (GO:0035473)

leukotriene-C4 synthase activity
(GO:0004464)

ErbB-3 class receptor binding (GO:0043125)

chondroitin sulfotransferase activity
(GO:0034481)

carbonyl reductase (NADPH) activity
(GO:0004090)

oligosaccharyl transferase activity
(GO:0004576)

neurotrophin TRKA receptor binding
(GO:0005168)

neurotrophin TRK receptor binding
(GO:0005167)

N-acetylglucosamine 6-O-sulfotransferase
activity (GO:0001517)

interleukin-6 receptor binding (GO:0005138)

acid-amino acid ligase activity (GO:0016881)

0

2

4

Enrichment log(combined score)