

GO_Biological_Process_2021

Term

regulation of kidney development
(GO:0090183)
positive regulation of myoblast
proliferation (GO:2000288)
positive regulation of gamma–delta T cell
activation (GO:0046645)
nephron tubule formation (GO:0072079)
negative regulation of cell volume
(GO:0045794)
metanephric nephron tubule morphogenesis
(GO:0072282)
kidney morphogenesis (GO:0060993)
camera–type eye photoreceptor cell
differentiation (GO:0060219)
tRNA surveillance (GO:0106354)
spinal cord motor neuron differentiation
(GO:0021522)
positive regulation of osteoblast
proliferation (GO:0033690)
positive regulation of kidney development
(GO:0090184)
nuclear polyadenylation–dependent tRNA
catabolic process (GO:0071038)
nuclear polyadenylation–dependent rRNA
catabolic process (GO:0071035)
nuclear polyadenylation–dependent ncRNA
catabolic process (GO:0071046)
neuroepithelial cell differentiation
(GO:0060563)
negative regulation of transporter activity
(GO:0032410)
cellular response to cold (GO:0070417)
cardiac ventricle formation (GO:0003211)

0

2

4

6

Enrichment log(combined score)

GO_Cellular_Component_2021

Term

Cul5–RING ubiquitin ligase complex
(GO:0031466)

transcription factor TFIIH core complex
(GO:0000439)

post–mRNA release spliceosomal complex
(GO:0071014)

transcription factor TFIIH holo complex
(GO:0005675)

proton–transporting V–type ATPase complex
(GO:0033176)

U2–type precatalytic spliceosome
(GO:0071005)

platelet alpha granule membrane (GO:0031092)

precatalytic spliceosome (GO:0071011)

endoplasmic reticulum tubular network
(GO:0071782)

U2–type spliceosomal complex (GO:0005684)

0

2

4

Enrichment log(combined score)

GO_Molecular_Function_2021

Term

solute:proton antiporter activity
(GO:0015299)

JUN kinase binding (GO:0008432)

solute:cation antiporter activity
(GO:0015298)

7SK snRNA binding (GO:0097322)

volume-sensitive anion channel activity
(GO:0005225)

double-stranded telomeric DNA binding
(GO:0003691)

voltage-gated chloride channel activity
(GO:0005247)

leucine zipper domain binding (GO:0043522)

cyclin-dependent protein serine/threonine
kinase inhibitor activity (GO:0004861)

kinase inhibitor activity (GO:0019210)

0

2

4

6

Enrichment log(combined score)