

GO_Biological_Process_2021

Term

regulation of mRNA processing (GO:0050684)

regulation of RNA splicing (GO:0043484)

regulation of mRNA splicing, via spliceosome
(GO:0048024)

regulation of DNA methylation (GO:0044030)

regulation of cell–cell adhesion involved in
gastrulation (GO:0070587)

negative regulation of heterotypic cell–cell
adhesion (GO:0034115)

spliceosomal complex assembly (GO:0000245)

positive regulation of transcription from
RNA polymerase II promoter in response to
endoplasmic reticulum stress (GO:1990440)

positive regulation of non–canonical Wnt
signaling pathway (GO:2000052)

RNA processing (GO:0006396)

0

2

4

6

Enrichment log(combined score)

GO_Cellular_Component_2021

Term

TORC1 complex (GO:0031931)

MLL1/2 complex (GO:0044665)

MLL1 complex (GO:0071339)

TORC2 complex (GO:0031932)

Golgi stack (GO:0005795)

nucleus (GO:0005634)

U2-type catalytic step 1 spliceosome
(GO:0071006)

intracellular membrane-bounded organelle
(GO:0043231)

transcription factor TFTC complex
(GO:0033276)

nBAF complex (GO:0071565)

Enrichment log(combined score)

0

1

2

3

4

5

GO_Molecular_Function_2021

Term

RNA binding (GO:0003723)

double-stranded RNA adenosine deaminase activity (GO:0003726)

cytoskeleton-nuclear membrane anchor activity (GO:0140444)

ATPase inhibitor activity (GO:0042030)

transcription regulatory region nucleic acid binding (GO:0001067)

U4 snRNA binding (GO:0030621)

glucuronosyl-N-acetylgalactosaminyl-proteoglycan 4-beta-N-acetylgalactosaminyltransferase activity (GO:0047238)

double-stranded methylated DNA binding (GO:0010385)

S-methyltransferase activity (GO:0008172)

minor groove of adenine-thymine-rich DNA binding (GO:0003680)

interleukin-6 receptor binding (GO:0005138)

0

2

4

Enrichment log(combined score)