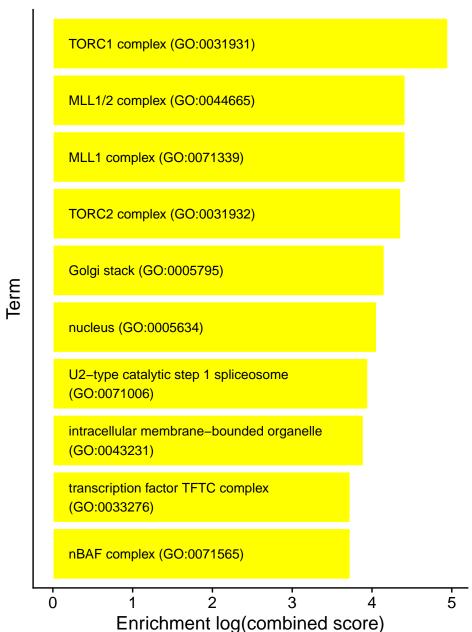
GO_Biological_Process_2021

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 6

Enrichment log(combined score)

GO_Cellular_Component_2021



GO_Molecular_Function_2021

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

Enrichment log(combined score)