

# GO\_Biological\_Process\_2021

Term

RNA processing (GO:0006396)

mRNA splice site selection (GO:0006376)

spliceosomal complex assembly (GO:0000245)

mRNA processing (GO:0006397)

mRNA splicing, via spliceosome (GO:0000398)

RNA splicing, via transesterification

reactions with bulged adenosine as

nucleophile (GO:0000377)

regulation of alternative mRNA splicing, via

spliceosome (GO:0000381)

regulation of RNA metabolic process

(GO:0051252)

regulation of maintenance of mitotic sister

chromatid cohesion (GO:0034182)

positive regulation of protein localization

to cilium (GO:1903566)

positive regulation of maintenance of sister

chromatid cohesion (GO:0034093)

positive regulation of maintenance of

mitotic sister chromatid cohesion

(GO:0034184)

negative regulation of protein

polyubiquitination (GO:1902915)

left/right axis specification (GO:0070986)

late endosome to Golgi transport

(GO:0034499)

0

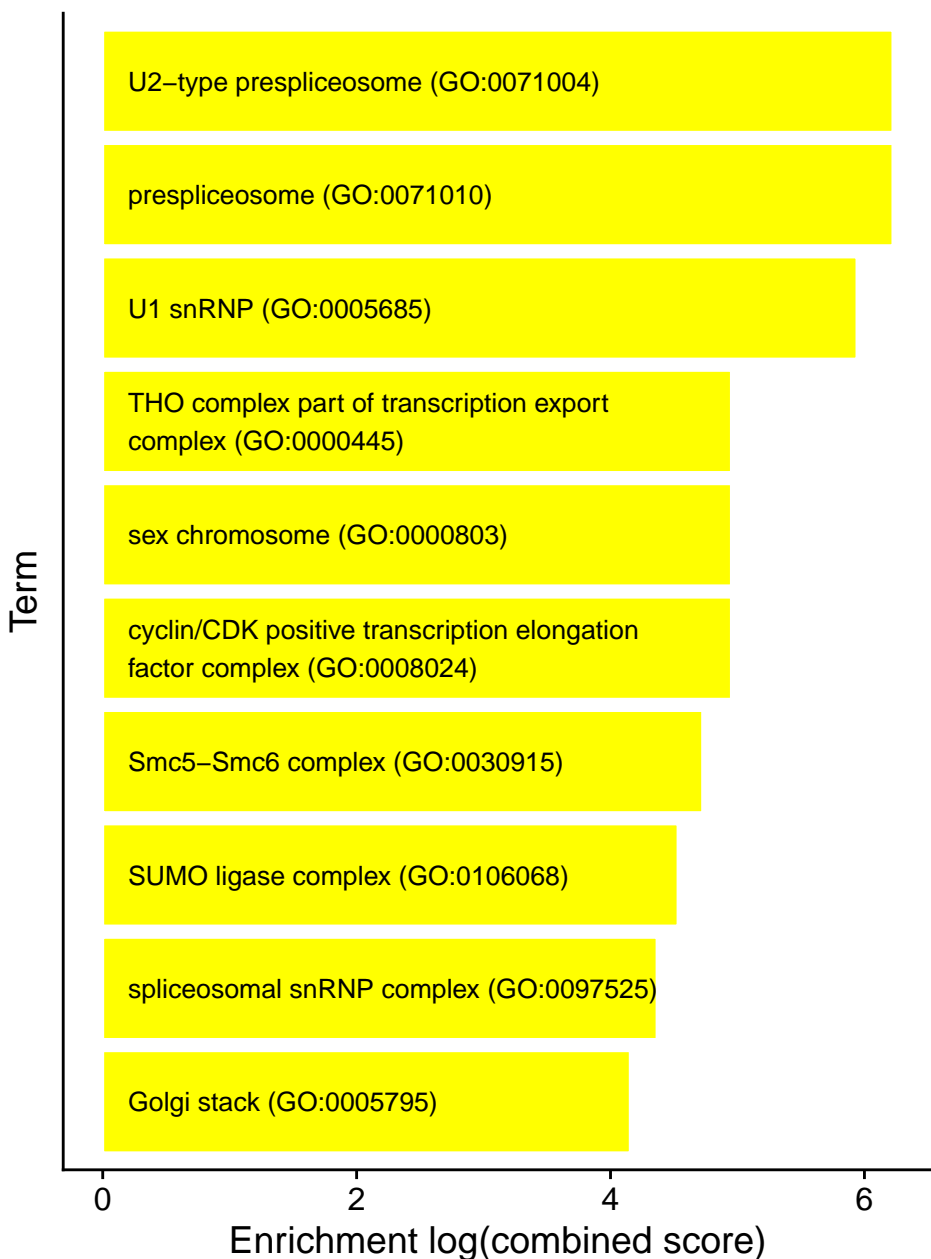
2

4

6

Enrichment log(combined score)

# GO\_Cellular\_Component\_2021



# GO\_Molecular\_Function\_2021

