

# GO\_Biological\_Process\_2021

Term

positive regulation of protein sumoylation  
(GO:0033235)

negative regulation of homotypic cell–cell  
adhesion (GO:0034111)

cellular response to thyroid hormone  
stimulus (GO:0097067)

attachment of mitotic spindle microtubules  
to kinetochore (GO:0051315)

response to thyroid hormone (GO:0097066)

regulation of homotypic cell–cell adhesion  
(GO:0034110)

kinetochore organization (GO:0051383)

maturation of LSU–rRNA from tricistronic  
rRNA transcript (SSU–rRNA, 5.8S rRNA,  
LSU–rRNA) (GO:0000463)

homologous chromosome pairing at meiosis  
(GO:0007129)

synaptonemal complex organization  
(GO:0070193)

synaptonemal complex assembly (GO:0007130)

macrophage migration (GO:1905517)

0

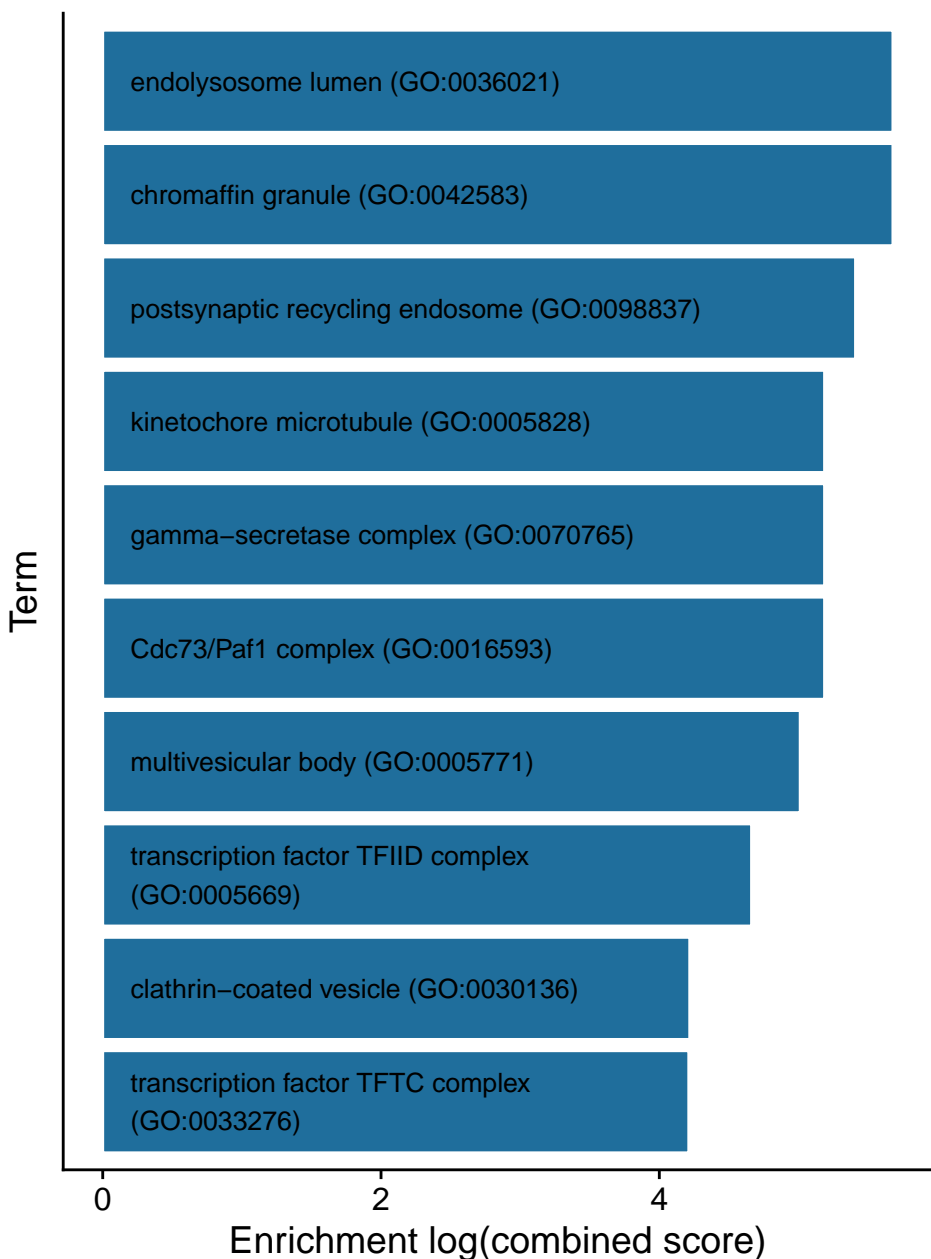
2

4

6

Enrichment log(combined score)

# GO\_Cellular\_Component\_2021



# GO\_Molecular\_Function\_2021

Term

calmodulin-dependent protein phosphatase  
activity (GO:0033192)

Y-form DNA binding (GO:0000403)

D-loop DNA binding (GO:0062037)

C-acetyltransferase activity (GO:0016453)

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

four-way junction helicase activity  
(GO:0009378)

acetyl-CoA C-acyltransferase activity  
(GO:0003988)

ion channel inhibitor activity (GO:0008200)

calcium-dependent protein serine/threonine  
phosphatase activity (GO:0004723)

MutLalpha complex binding (GO:0032405)

3',5'-cyclic-GMP phosphodiesterase activity  
(GO:0047555)

0

2

4

Enrichment log(combined score)