

# GO\_Biological\_Process\_2021

Term

cellular response to X-ray (GO:0071481)

regulation of skeletal muscle cell  
differentiation (GO:2001014)

regulation of axon guidance (GO:1902667)

presynaptic membrane assembly (GO:0097105)

amino sugar metabolic process (GO:0006040)

B cell homeostasis (GO:0001782)

regulation of dendritic spine development  
(GO:0060998)

presynaptic membrane organization  
(GO:0097090)

regulation of vascular associated smooth  
muscle cell migration (GO:1904752)

mitochondrial genome maintenance  
(GO:0000002)

0

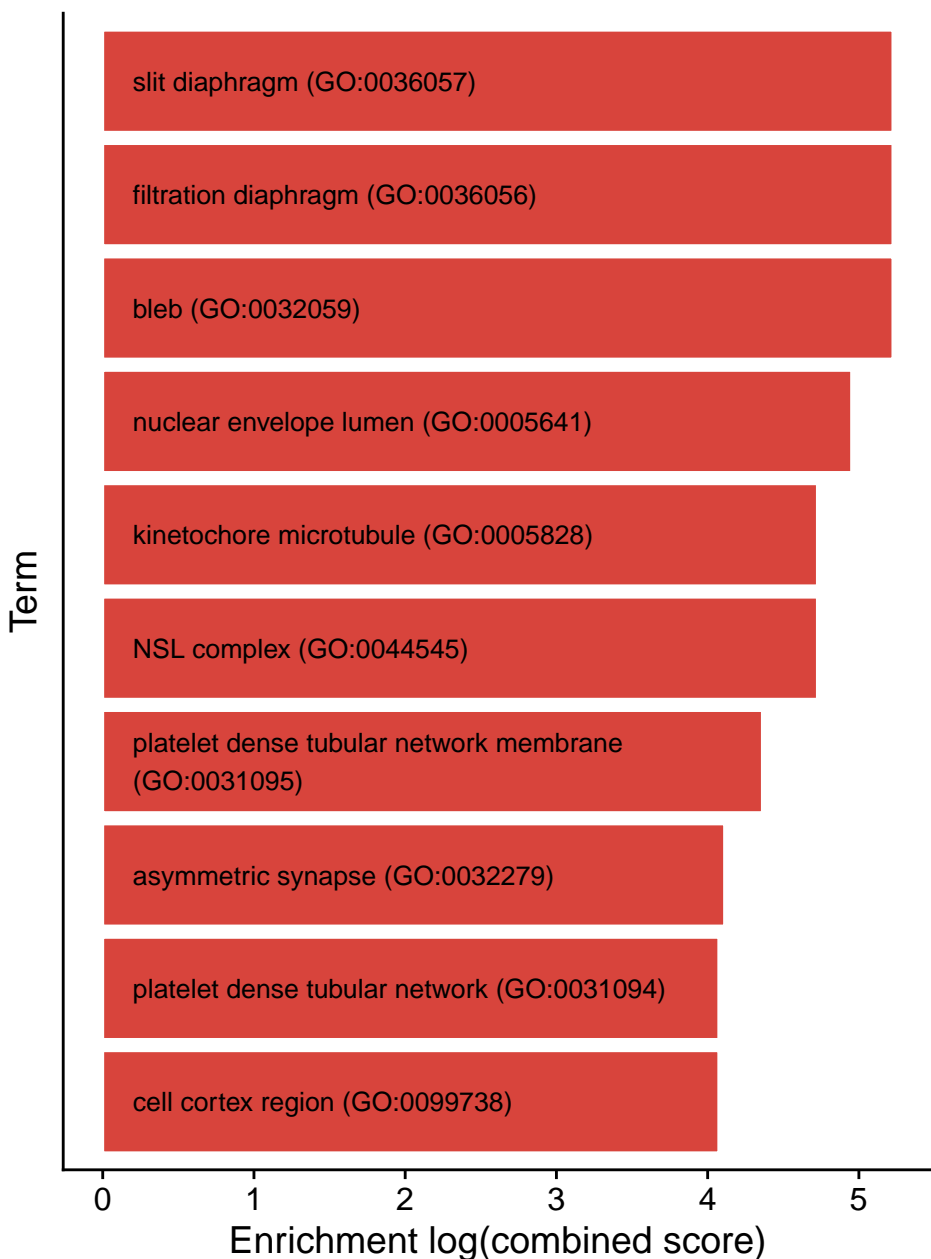
2

4

6

Enrichment log(combined score)

# GO\_Cellular\_Component\_2021



# GO\_Molecular\_Function\_2021

Term

GTPase activator activity (GO:0005096)

calmodulin-dependent protein phosphatase activity (GO:0033192)

histone deacetylase binding (GO:0042826)

primary miRNA binding (GO:0070878)

oncostatin-M receptor activity (GO:0004924)

leukemia inhibitory factor receptor activity (GO:0004923)

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

chromo shadow domain binding (GO:0070087)

cation:cation antiporter activity (GO:0015491)

calcium:cation antiporter activity (GO:0015368)

MHC class II protein binding (GO:0042289)

0

2

4

Enrichment log(combined score)