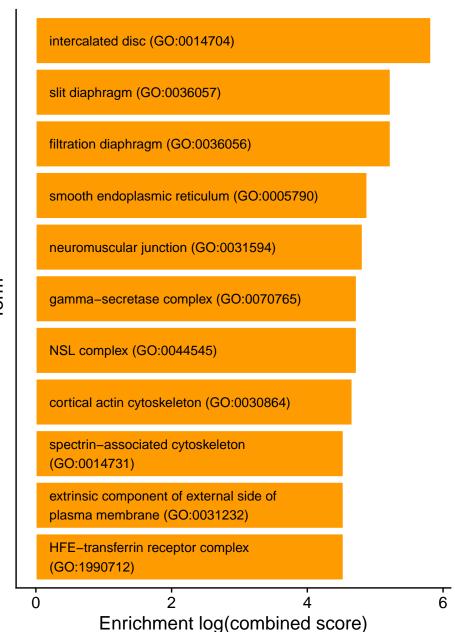
GO_Biological_Process_2021

protein localization to axon (GO:0099612) regulation of release of sequestered calcium ion into cytosol by sarcoplasmic reticulum (GO:0010880) negative regulation of keratinocyte proliferation (GO:0010839) positive regulation of cardiac muscle tissue growth (GO:0055023) regulation of actin filament-based movement (GO:1903115) positive regulation of sodium ion transmembrane transport (GO:1902307) digestive system development (GO:0055123) regulation of cell communication by electrical coupling (GO:0010649) regulation of ryanodine-sensitive calcium-release channel activity (GO:0060314) positive regulation of ion transmembrane transporter activity (GO:0032414) 0 6

Enrichment log(combined score)

GO_Cellular_Component_2021



GO_Molecular_Function_2021

```
3',5'-cyclic-GMP phosphodiesterase activity
  (GO:0047555)
  3',5'-cyclic-nucleotide phosphodiesterase
  activity (GO:0004114)
  cyclic-nucleotide phosphodiesterase activity
  (GO:0004112)
  3',5'-cyclic-AMP phosphodiesterase activity
  (GO:0004115)
  phosphotransferase activity, nitrogenous
  group as acceptor (GO:0016775)
  phosphodiesterase I activity (GO:0004528)
  phosphatidylserine flippase activity
  (GO:0140346)
  oxidoreductase activity, acting on a sulfur
  group of donors, disulfide as acceptor
  (GO:0016671)
  neuroligin family protein binding
   (GO:0097109)
  nerve growth factor binding (GO:0048406)
  fibroblast growth factor-activated receptor
  activity (GO:0005007)
  creatine kinase activity (GO:0004111)
  calcium, potassium:sodium antiporter
  activity (GO:0008273)
  beta-2 adrenergic receptor binding
   (GO:0031698)
                                                             6
0
```

Enrichment log(combined score)