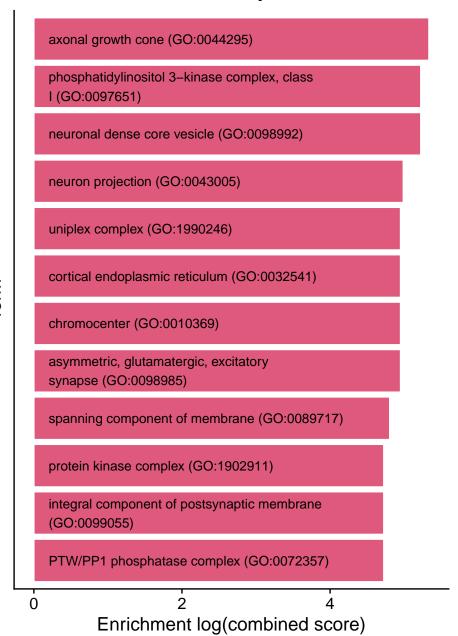
## GO\_Biological\_Process\_2021

negative regulation of anion transport (GO:1903792) negative regulation of amino acid transport (GO:0051956) regulation of transforming growth factor beta2 production (GO:0032909) regulation of oxidative stress-induced neuron intrinsic apoptotic signaling pathway (GO:1903376) regulation of myosin-light-chain-phosphatase activity (GO:0035507) positive regulation of aspartic-type endopeptidase activity involved in amyloid precursor protein catabolic process (GO:1902961) positive regulation of adenylate cyclase activity (GO:0045762) neurogenesis (GO:0022008) positive regulation of aspartic-type peptidase activity (GO:1905247) regulation of aspartic-type endopeptidase activity involved in amyloid precursor protein catabolic process (GO:1902959) 0 6 Enrichment log(combined score)

## GO\_Cellular\_Component\_2021



## GO\_Molecular\_Function\_2021

transmembrane receptor protein tyrosine kinase adaptor activity (GO:0005068) protein phosphorylated amino acid binding (GO:0045309) insulin-like growth factor receptor binding (GO:0005159) voltage-gated calcium channel activity involved in cardiac muscle cell action potential (GO:0086007) syndecan binding (GO:0045545) neuropeptide Y receptor activity (GO:0004983) nerve growth factor binding (GO:0048406) calcitonin family receptor activity (GO:0097642) beta-galactosidase activity (GO:0004565) GPI-linked ephrin receptor activity (GO:0005004) FATZ binding (GO:0051373) ErbB-3 class receptor binding (GO:0043125) 0

Enrichment log(combined score)