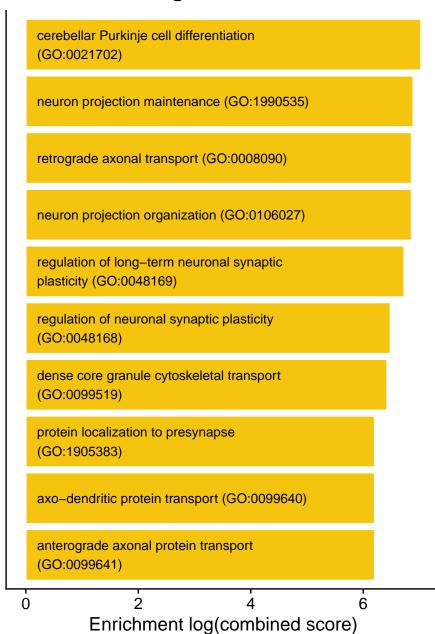
GO_Biological_Process_2021



axonal growth cone (GO:0044295) vacuolar proton-transporting V-type ATPase, V1 domain (GO:0000221) proton-transporting V-type ATPase, V1 domain (GO:0033180) intrinsic component of synaptic vesicle membrane (GO:0098563) synaptic vesicle membrane (GO:0030672) exocytic vesicle membrane (GO:0099501) neuron projection (GO:0043005) vacuolar proton-transporting V-type ATPase complex (GO:0016471) axon (GO:0030424) integral component of synaptic vesicle membrane (GO:0030285) 0 6

Enrichment log(combined score)

pyrophosphate hydrolysis-driven proton transmembrane transporter activity (GO:0009678) cuprous ion binding (GO:1903136) proton-transporting ATPase activity, rotational mechanism (GO:0046961) ATPase activity, coupled to transmembrane movement of ions, rotational mechanism (GO:0044769) MHC class II protein complex binding (GO:0023026) phosphoserine residue binding (GO:0050815) P-type calcium transporter activity (GO:0005388) disordered domain specific binding (GO:0097718) cell adhesion mediator activity (GO:0098631) protein kinase A catalytic subunit binding (GO:0034236) P-type ion transporter activity (GO:0015662) 0 6

Enrichment log(combined score)