GO_Biological_Process_2021

protein localization to axon (GO:0099612)

presynaptic membrane assembly (GO:0097105)

presynaptic membrane organization (GO:0097090)

presynapse assembly (GO:0099054)

astral microtubule organization (GO:0030953)

membrane assembly (GO:0071709)

negative regulation of locomotion (GO:0040013)

cellular potassium ion homeostasis (GO:0030007)

negative regulation of cellular component movement (GO:0051271)

ubiquitin-dependent SMAD protein catabolic process (GO:0030579)

regulation of retinal ganglion cell axon guidance (GO:0090259)

regulation of receptor catabolic process (GO:2000644)

regulation of histone H4-K16 acetylation (GO:2000618)

regulation of basement membrane assembly involved in embryonic body morphogenesis (GO:1904259)

regulation of Fc-gamma receptor signaling pathway involved in phagocytosis (GO:1905449)

receptor transactivation (GO:0035624)

protein processing involved in protein targeting to mitochondrion (GO:0006627)

positive regulation of vascular associated smooth muscle cell apoptotic process (GO:1905461)

positive regulation of basement membrane assembly involved in embryonic body morphogenesis (GO:1904261)

positive regulation of amino acid transport (GO:0051957)

monocyte extravasation (GO:0035696)

extracellular exosome biogenesis (GO:0097734)

cellular response to heparin (GO:0071504)

Roundabout signaling pathway (GO:0035385)

Term

Enrichment log(combined score)

0   2   4   6

GO_Cellular_Component_2021

kinetochore microtubule (GO:0005828)

synaptic membrane (GO:0097060)

MPP7−DLG1−LIN7 complex (GO:0097025)

postsynaptic recycling endosome (GO:0098837)

cyclin/CDK positive transcription elongation factor complex (GO:0008024)

neuromuscular junction (GO:0031594)

postsynaptic density membrane (GO:0098839)

postsynaptic specialization membrane (GO:0099634)

glutamatergic synapse (GO:0098978)

spectrin−associated cytoskeleton (GO:0014731)

NMDA selective glutamate receptor complex (GO:0017146)

Term

0   2   4   6

Enrichment log(combined score)

GO_Molecular_Function_2021

glutamate–gated calcium ion channel activity (GO:0022849)

GPI–linked ephrin receptor activity (GO:0005004)

1–phosphatidylinositol–4–phosphate 5–kinase activity (GO:0016308)

interleukin–6 receptor binding (GO:0005138)

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

mitogen–activated protein kinase kinase binding (GO:0031434)

Roundabout binding (GO:0048495)

NMDA glutamate receptor activity (GO:0004972)

cadherin binding (GO:0045296)

sodium:chloride symporter activity (GO:0015378)

potassium:chloride symporter activity (GO:0015379)

potassium channel inhibitor activity (GO:0019870)

interleukin–1 binding (GO:0019966)

guanylate kinase activity (GO:0004385)

bioactive lipid receptor activity (GO:0045125)

Term

Enrichment log(combined score)