intracellular sequestering of iron ion (GO:0006880)

glutamate catabolic process (GO:0006538)

negative regulation of amyloid precursor protein biosynthetic process (GO:0042985)

SRP-dependent cotranslational protein targeting to membrane (GO:0006614)

cotranslational protein targeting to membrane (GO:0006613)

0 2 4
Enrichment log(combined score)

6

astrocyte projection (GO:0097449)

sodium:potassium-exchanging ATPase complex (GO:0005890)

autolysosome (GO:0044754)

cytosolic large ribosomal subunit (GO:0022625)

cytosolic small ribosomal subunit (GO:0022627)

0 2 4
Enrichment log(combined score)

6

ferric iron binding (GO:0008199) aldehyde-lyase activity (GO:0016832) sequence-specific single stranded DNA binding (GO:0098847) phosphotransferase activity, nitrogenous group as acceptor (GO:0016775) oxo-acid-lyase activity (GO:0016833) L-leucine transmembrane transporter activity (GO:0015190) L-lactate dehydrogenase activity (GO:0004459) ganglioside binding (GO:0035594) creatine kinase activity (GO:0004111) calcitonin family receptor activity (GO:0097642) cadmium ion binding (GO:0046870) benzodiazepine receptor binding (GO:0030156) 0 6

Enrichment log(combined score)