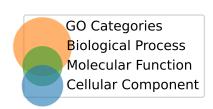
## GO Terms Network Node size represents significance (-log10 p-value)

clathrin-coated endocytic vesicle membrane (GO:0030669) p=1.72e-02



clathrin-coated endocytic vesicle (GO:0045334) p=2.24e-02 insulin-like growth factor binding (GO:0005520) p=2.08e-02

cell-substrate junction (GO:0030055) p=1.54e-02

riboflavin transmembrane transporter activity (GO:0032217) p=8.37e-02

positive regulation
of protein
modification
process
(GO:0031401)
p=3.60e-03

insulin-like growth factor I binding (GO:0031994) p=2.08e-02

vascular transport (GO:0010232) p=9.44e-03 positive regulation of phosphorylation (G0:0042327) p=1.05e-03

regulation of ERK1

and ERK2 cascade (GO:0070372)

p=2.52e-02

clathrin-coated vesicle membrane (GO:0030665) p=2.24e-02 GPI-linked ephrin receptor activity (GO:0005004) p=8.37e-02 transport across blood-brain barrier (G0:0150104) p=9.44e-03 regulation of protein phosphorylation (GO:0001932) p=2.01e-04

ABC-type xenobiotic transporter activity (GO:0008559) p=2.08e-02

focal adhesion (GO:0005925) p=1.54e-02

regulation of stress-activated MAPK cascade (GO:0032872) p=2.52e-02

urate transmembrane transporter activity (GO:0015143) p=8.37e-02 re regulation protein

positive regulation of protein phosphorylation (G0:0001934) p=2.52e-02

syndecan binding (GO:0045545) p=8.37e-02

vesicle (GO:0031982) p=3.01e-02 type 1 fibroblast growth factor receptor binding (GO:0005105) p=8.37e-02

peroxisome (GO:0005777) p=3.81e-02

brush border membrane (GO:0031526) p=3.81e-02