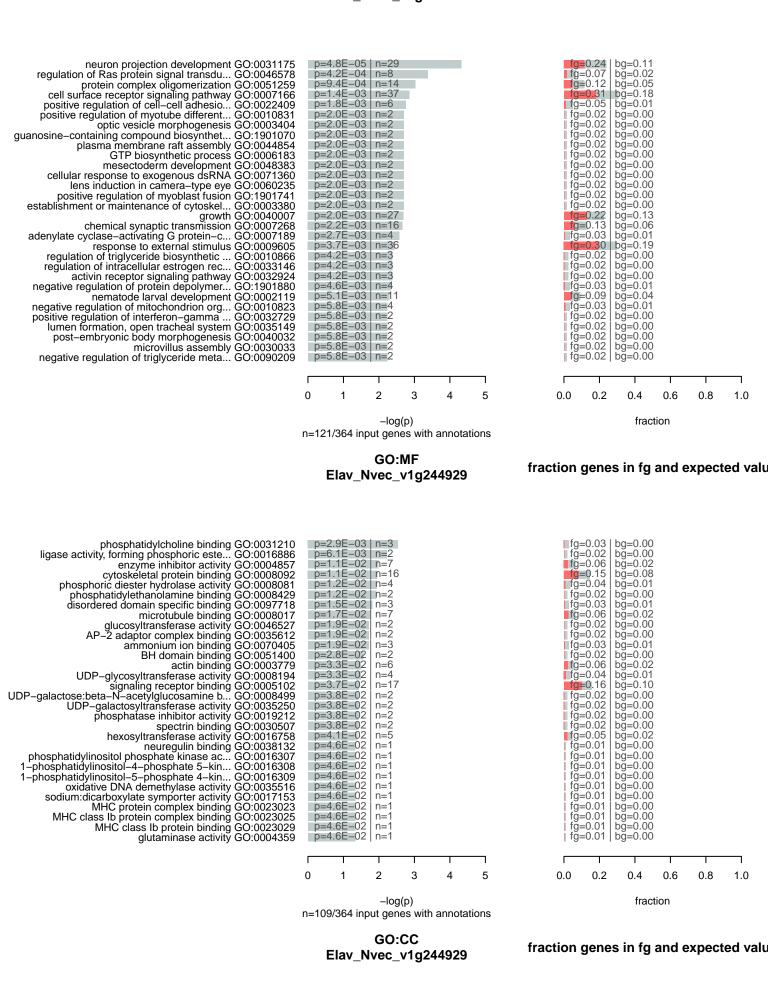
fraction



lamellipodium GO:0030027 synaptic membrane GO:0097060 p=3.5E-04 | n=9 p=7.0E-04 | n=9 bg=0.02 bg=0.02 fg=0.08 fg=0.08 p=1.9E-03 n=2 p=1.9E-03 n=2 fg=0.02 photoreceptor outer segment membrane GO:0042622 bg=0.00 collagen and cuticulin–based cuticle ext... GO:00031253
transcription elongation factor complex GO:008023 fg=0.02 bg=0.00 fg=0.02 bg=0.00 fg=0.10 bg=0.03 fg=0.03 bg=0.01 plasma membrane protein complex GO:0008025
plasma membrane protein complex GO:0098797
clathrin–coated pit GO:0005905
presynapse GO:0098793
growth cone GO:0030426
cytoplasmic side of plasma membrane GO:0004853
plasma membrane raft GO:0044853 bq=0.03 p=5.5E-03 n=3 p=5.6E-03 n=12 p=5.7E-03 n=7 fg=0.03 fg=0.10 bg=0.00 bg=0.05 bg=0.02 bg=0.02 bg=0.01 fg=0.06 p=7.8E-03 | n=7 fa=0.06 p=8.4E-03 n=5 fg=0.04 bg=0.09 bg=0.01 somatodendritic compartment GO:0036477 extrinsic component of plasma resultante GO:0018897 p=9.2E-03 | n=18 q = 0.15p=9.7E-03 | n=5 fg=0.04 filamentous actin GO:0031941 growth cone lamellipodium GO:1990761 p=1.1E-02 | n=2 g=0.02 bg=0.00 p=1.1E-02 | n=2 fg=0.02 bg=0.00 vacuolar lumen GO:0005775 extrinsic component of cytoplasmic side ... GO:0031234 p=1.1E-02 n=5 p=1.1E-02 n=4 fg=0.04 fg=0.03 bg=0.01 bg=0.01 fg=0.13 fg=0.03 secretory vesicle GO:0099503 caveola GO:0005901 p=1.4E-02 | n=15 bg=0.07 p=1.6E-02 n=4 bg=0.01 p=1.6E-02 n=5 p=1.6E-02 n=7 brush border GO:0005903 basolateral plasma membrane GO:0016323 fg=0.04 bg=0.01 fg=0.06 bg=0.02 p=1.6E-02 p=1.7E-02 p=1.7E-02 basal plasma membrane GO:0009925 n=7fg=0.06 bg=0.02 n=12 fg=0.10 bg=0.03 side of membrane GO:0098552 sensory dendrite GO:0071683 n=2 g=0.02 bg=0.00 fg=0.23 bg=0.15 cytoplasmic vesicle GO:0031410 intracellular vesicle GO:0097708 p=1.8E-02 n=27 p=1.8E-02 p=2.0E-02 n=27 n=10 fg=0.23 fg=0.09 bg=0.15 anchoring junction GO:0070161 cell-cell junction GO:0005911 bq=0.04 p=2.0E-02 fg=0.09 bg=0.04 0 2 3 5 0.0 1 4 0.2 0.4 0.6 0.8 1.0

-log(p)
n=117/364 input genes with annotations