



filopodium GO:0030175 pronucleus GO:0045120 cell leading edge GO:0031252 striated muscle dense body GO:0055120 synaptic membrane GO:0097060 presynaptic membrane GO:0042734 basal plasma membrane GO:0016232 p=3.1E-03 n=3 p=5.2E-03 n=2 p=7.2E-03 n=5 p=9.5E-03 n=2 fg=0.09 fg=0.06 bg=0.01 bg=0.00 fg=0.14 fg=0.06 bg=0.04 bg=0.00 p=1.0E-02 n=4 p=1.1E-02 n=3 p=1.2E-02 n=4 p=1.2E-02 n=4 fg=0.11 bg=0.02 fg=0.09 bg=0.01 fg=0.11 fg=0.11 bğ=0.03 basal plasma membrane GO:0009925
basolateral plasma membrane GO:0016323
melanosome GO:0042470
mediator complex GO:0016592
pigment granule GO:0048770
postsynaptic membrane GO:0045211
RNA polymerase III transcription regulat... GO:0090576
lamellipodium membrane GO:0031258
extrinsic component of mitochondrial inn... GO:0031314
clathrin-sculpted vesicle GO:0060198 ba = 0.03p=1.2E-02 n=2 p=1.2E-02 n=2 fg=0.06 fg=0.06 bg=0.00 bq=0.00 bg=0.01 bg=0.02 bg=0.00 fg=0.06 p=1.5E-02 | n=2 p=1.9F-02fa=0.09 n=3fg=0.03 fg=0.03 | bg=0.00 fg=0.03 | bg=0.00 p=1.9E-02n=1 p=1.9E-02 n=1 clathrin–sculpted vesicle GO:0060198 transcription factor TFIIK complex GO:0070985 transcription factor TFIIIC complex GO:0000127 fg=0.03 p=1.9E-02n=1bg=0.00 p=1.9E -02 n=1fg=0.03 bg=0.00 fg=0.03 fg=0.03 p=1.9E-02n=1bg=0.00 clathrin–sculpted gamma–aminobutyric aci... GO:0061200 clathrin–sculpted gamma–aminobutyric aci... GO:0061202 interphase microtubule organizing center GO:0031021 basal part of cell GO:0045178 lamellipodium GO:0030027 p=1.9E-02 bg=0.00 n=1fg=0.03 fg=0.03 bg=0.00 bg=0.00 p=1.9E-02n=1p=1.9E-02 n=1p=2.0E-02fg=0.11 fg=0.09 n=4bg=0.03 p=2.8E-02 bg=0.02 n=3vacuolar membrane GO:0005774 p=3.0E-02 n=4fg=0.11 bg=0.03 p=3.1E-02 p=3.3E-02 fg=0.17 vacuole GO:0005773 photoreceptor outer segment GO:0001750 lysosome GO:0005764 bg=0.07 n=6 fg=0.06 n=2bg=0.01 p=3.5E-02 bg=0.05 n=5 $f_{q=0.14}$ p=3.7E-02 p=3.8E-02 main axon GO:0044304 n=2 fg=0.06 bg=0.01 fg=0.03 carboxy-terminal domain protein kinase c... GO:0032806 type lb terminal bouton GO:0061176 bq = 0.00n=1 p=3.8E-02 fg=0.03 bg=0.00 0 2 3 5 0.0 1 0.2 0.4 0.6 0.8 1.0 fraction -log(p) n=35/127 input genes with annotations