

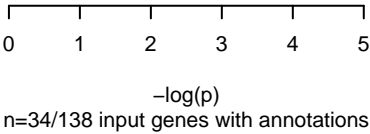
GO:BP
Elav_Nvec_vc1.1_XM_032378021.2

fraction genes in fg and expected value

negative regulation of neuroblast prolif... GO:0007406
negative regulation of stem cell prolife... GO:2000647
sensory perception of mechanical stimulu... GO:0050954
establishment or maintenance of polarity... GO:0016334
interleukin-6-mediated signaling pathway GO:0070102
gamma-aminobutyric acid secretion GO:0014051
positive regulation of histone H3-K4 met... GO:0051571
aorta morphogenesis GO:0035909
chitin-based larval cuticle pattern form... GO:0035293
histone phosphorylation GO:0016572
establishment of synaptic vesicle locali... GO:0097480
dicarboxylic acid transport GO:0006835
movement of cell or subcellular componen... GO:0006928
post-embryonic animal organ development GO:0048569
negative regulation of mRNA catabolic pr... GO:1902373
sympathetic nervous system development GO:0048485
positive regulation of neurotransmitter ... GO:0001956
cellular response to dexamethasone stimu... GO:0071549
cellular response to pH GO:0071467
regulation of neurotransmitter levels GO:0001505
synaptic vesicle transport GO:0048489
chordate embryonic development GO:0043009
regulation of response to drug GO:2001023
positive regulation of transcription elo... GO:0032968
triglyceride biosynthetic process GO:0019432
response to auditory stimulus GO:0010996
regulation of dopamine metabolic process GO:0042053
synaptic vesicle priming GO:0016082
negative regulation of protein kinase ac... GO:0006469
positive regulation of epithelial cell p... GO:0050679

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p=4.0E-03	n=2
p=4.0E-03	n=3
p=4.7E-03	n=13
p=4.9E-03	n=7
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p=5.4E-03	n=9
p=5.8E-03	n=4
p=5.9E-03	n=7
p=6.4E-03	n=3
p=6.7E-03	n=2
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p=7.0E-03	n=4
p=8.3E-03	n=3

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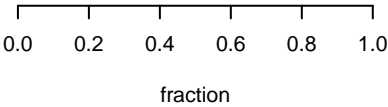
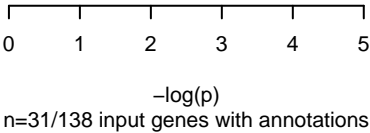
GO:MF
Elav_Nvec_vc1.1_XM_032378021.2

fraction genes in fg and expected value

outward rectifier potassium channel acti... GO:0015271
ion binding GO:0043167
histone kinase activity GO:0035173
iron-responsive element binding GO:0030350
protein tyrosine phosphatase activity, m... GO:0030946
acid-sensing ion channel activity GO:0044736
ferrochelatase activity GO:0004325
choline dehydrogenase activity GO:0008812
7SK snRNA binding GO:0097322
intracellular sodium activated potassium... GO:0005228
myosin II binding GO:0045159
short-branched-chain-acyl-CoA dehydrogen... GO:0016937
myosin binding GO:0017022
kinesin binding GO:0019894
kinase inhibitor activity GO:0019210
protein kinase inhibitor activity GO:0004860
purine ribonucleoside triphosphate bindi... GO:0035639
carbohydrate derivative binding GO:0097367
anion binding GO:0043168
oxidoreductase activity, acting on CH-OH... GO:0016614
ligand-gated cation channel activity GO:0099094
sodium ion transmembrane transporter act... GO:0015081
ligand-gated sodium channel activity GO:0015280
oxysterol binding GO:0008142
ATPase-coupled organic acid transmembran... GO:0033283
ATPase-coupled carboxylic acid transmemb... GO:0033284
ATPase-coupled monocarboxylic acid trans... GO:0033285
bile acid transmembrane transporter acti... GO:0015125
canalicular bile acid transmembrane tran... GO:0015126
substance P receptor activity GO:0016496

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p=1.9E-02	n=1
p=2.5E-02	n=2
p=2.9E-02	n=2
p=2.9E-02	n=2
p=2.9E-02	n=2
p=3.1E-02	n=5
p=3.2E-02	n=6
p=3.3E-02	n=8
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GO:CC
Elav_Nvec_vc1.1_XM_032378021.2

fraction genes in fg and expected value

transcription elongation factor complex GO:0008023
potassium channel complex GO:0034705
voltage-gated potassium channel complex GO:0008076
synaptic vesicle GO:0008021
egg chorion GO:0042600
micropyle GO:0070825
Dbf4-dependent protein kinase complex GO:0031431
Cdc73/Paf1 complex GO:0016593
female germline ring canal outer rim GO:0035182
Dsl1/NZR complex GO:0070939
smooth septate junction GO:0005920
exocytic vesicle GO:0070382
intercellular bridge GO:0045171
neuronal cell body membrane GO:0032809
cell body membrane GO:0044298
external encapsulating structure GO:0030312
apical part of cell GO:0045177
cell junction GO:0030054
gut granule GO:0044840
gut granule membrane GO:0044841
female germline ring canal GO:0035324
extrinsic component of endoplasmic retic... GO:0042406
NELF complex GO:0032021
presynapse GO:0098793
growth cone GO:0030426
site of polarized growth GO:0030427
axon GO:0030424
synapse GO:0045202
cell cortex GO:0005938
cell periphery GO:0071944

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p=1.2E-02	n=2
p=1.3E-02	n=4
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p=1.9E-02	n=1
p=1.9E-02	n=1
p=1.9E-02	n=1
p=1.9E-02	n=1
p=1.9E-02	n=1
p=2.0E-02	n=4
p=2.1E-02	n=2
p=2.9E-02	n=2
p=3.3E-02	n=2
p=3.3E-02	n=2
p=3.6E-02	n=5
p=3.7E-02	n=9
p=3.8E-02	n=1
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