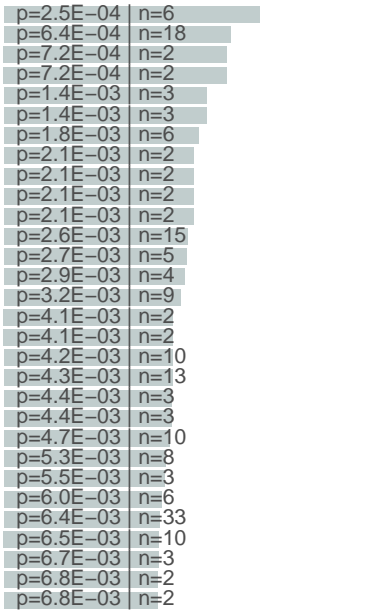


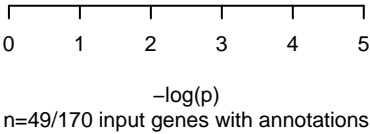
GO:BP  
Elav\_Nvec\_vc1.1\_XM\_001626384.3

fraction genes in fg and expected value

regeneration	GO:0031099
multi-organism reproductive process	GO:0044703
involuntary skeletal muscle contraction	GO:0003011
negative regulation by host of viral gen...	GO:0044828
negative regulation of vasculature devel...	GO:1901343
negative regulation of JNK cascade	GO:0046329
learning	GO:0007612
modulation by virus of host cellular pro...	GO:0019054
skeletal muscle tissue growth	GO:0048630
positive regulation of endothelial cell ...	GO:2001028
regulation of smooth muscle cell differe...	GO:0051150
gamete generation	GO:0007276
multi-multicellular organism process	GO:0044706
response to axon injury	GO:0048678
cellular response to organonitrogen comp...	GO:0071417
regulation of vascular endothelial growt...	GO:0030947
negative regulation of oxidoreductase ac...	GO:0051354
sensory system development	GO:0048880
cellular process involved in reproductio...	GO:0022412
response to cold	GO:0009409
regulation of lipid transport	GO:0032368
regulation of neuron differentiation	GO:0045664
positive regulation of neurogenesis	GO:0050769
glial cell migration	GO:0008347
response to xenobiotic stimulus	GO:0009410
protein metabolic process	GO:0019538
response to organic cyclic compound	GO:0014070
regulation of blood vessel endothelial c...	GO:0043535
positive regulation of ATP biosynthetic ...	GO:2001171
regulation of protein polyubiquitination	GO:1902914



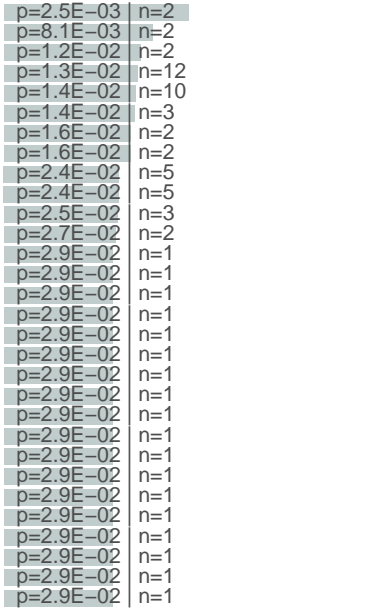
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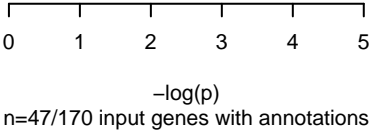
GO:MF  
Elav\_Nvec\_vc1.1\_XM\_001626384.3

fraction genes in fg and expected value

receptor-receptor interaction	GO:0090722
phospholipase binding	GO:0043274
G-protein alpha-subunit binding	GO:0001965
protein dimerization activity	GO:0046983
protein homodimerization activity	GO:0042803
histone deacetylase binding	GO:0042826
peptide transmembrane transporter activi...	GO:1904680
adrenergic receptor binding	GO:0031690
ubiquitin-like protein ligase activity	GO:0061659
ubiquitin protein ligase activity	GO:0061630
translation regulator activity	GO:0045182
amide transmembrane transporter activity	GO:0042887
G-protein beta-subunit binding	GO:0031681
[3-methyl-2-oxobutanoate dehydrogenase (...	GO:0047323
RNA-3'-phosphate cyclase activity	GO:0003963
dopamine binding	GO:0035240
ATP-gated ion channel activity	GO:0035381
1-alkylglycerophosphocholine O-acyltrans...	GO:0047191
bradykinin receptor binding	GO:0031711
B2 bradykinin receptor binding	GO:0031713
F-box domain binding	GO:1990444
nucleotide receptor activity	GO:0016502
mitochondrial ribosome binding	GO:0097177
1-alkenylglycerophosphocholine O-acyltra...	GO:0047159
mRNA 5'-UTR binding	GO:0048027
beta1-adrenergic receptor activity	GO:0004940
beta2-adrenergic receptor activity	GO:0004941
extracellularly ATP-gated cation channel...	GO:0004931
beta-adrenergic receptor activity	GO:0004939
purinergic nucleotide receptor activity	GO:0001614



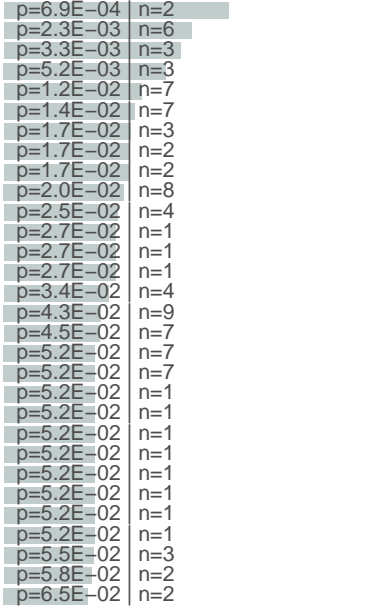
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GO:CC  
Elav\_Nvec\_vc1.1\_XM\_001626384.3

fraction genes in fg and expected value

Lewy body	GO:0097413
membrane raft	GO:0045121
polysome	GO:0005844
specific granule membrane	GO:0035579
lysosome	GO:0005764
lytic vacuole	GO:0000323
plasma membrane raft	GO:0044853
germ cell nucleus	GO:0043073
azurophil granule membrane	GO:0035577
perinuclear region of cytoplasm	GO:0048471
lysosomal membrane	GO:0005765
MCM8-MCM9 complex	GO:0097362
dihydrolipoyl dehydrogenase complex	GO:0045240
mitochondrial alpha-ketoglutarate dehydr...	GO:0005947
lytic vacuole membrane	GO:0098852
somatodendritic compartment	GO:0036477
vacuole	GO:0005773
dendritic tree	GO:0097447
dendrite	GO:0030425
mitochondrial tricarboxylic acid cycle e...	GO:0030062
tricarboxylic acid cycle enzyme complex	GO:0045239
Parkin-FBXW7-Cul1 ubiquitin ligase compl...	GO:1990452
glycosylphosphatidylinositol-N-acetylglu...	GO:0000506
intrinsic component of nuclear inner mem...	GO:0031229
sarcoplasmic reticulum membrane	GO:0033017
integral component of nuclear inner memb...	GO:0005639
MCM complex	GO:0042555
terminal bouton	GO:0043195
focal adhesion	GO:0005925
caveola	GO:0005901



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