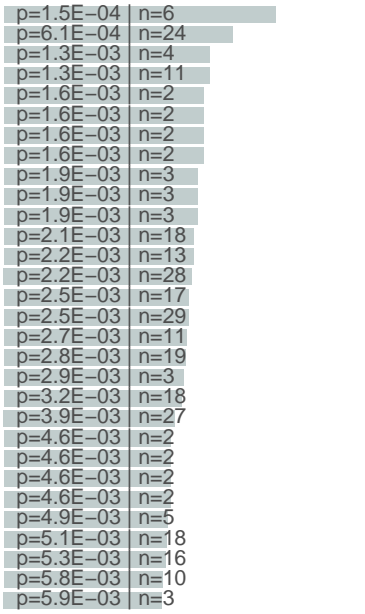


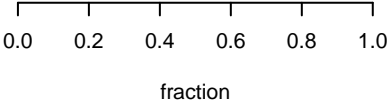
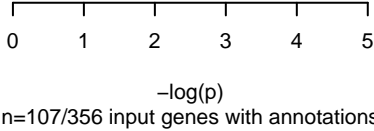
GO:BP
Elav_Nvec_vc1.1_XM_001627049.3

fraction genes in fg and expected value

regulation of mitochondrial membrane pot...	GO:0051881
negative regulation of cellular macromol...	GO:2000113
eye-antennal disc morphogenesis	GO:0007455
establishment or maintenance of cell pol...	GO:0007163
liver morphogenesis	GO:0072576
Golgi to transport vesicle transport	GO:0055108
negative regulation by host of viral gen...	GO:0044828
negative regulation of imaginal disc gro...	GO:0045571
negative regulation of cell fate specifi...	GO:0009996
pole cell migration	GO:0007280
regulation of R7 cell differentiation	GO:0045676
negative regulation of cellular protein ...	GO:0032269
positive regulation of neurogenesis	GO:0050769
negative regulation of gene expression	GO:0010629
oogenesis	GO:0048477
multi-organism reproductive process	GO:0044703
positive regulation of neuron differenti...	GO:0045666
regulation of anatomical structure morph...	GO:0022603
negative regulation of T cell proliferat...	GO:0042130
neuron projection morphogenesis	GO:0048812
positive regulation of developmental pro...	GO:0051094
phosphatidylglycerol acyl-chain remodeli...	GO:0036148
modulation by virus of host cellular pro...	GO:0019054
deoxynucleotide transport	GO:0030302
pyrimidine nucleotide import into mitoch...	GO:1990519
embryonic skeletal system morphogenesis	GO:0048704
negative regulation of multicellular org...	GO:0051241
brain development	GO:0007420
anatomical structure maturation	GO:0071695
response to gravity	GO:0009629



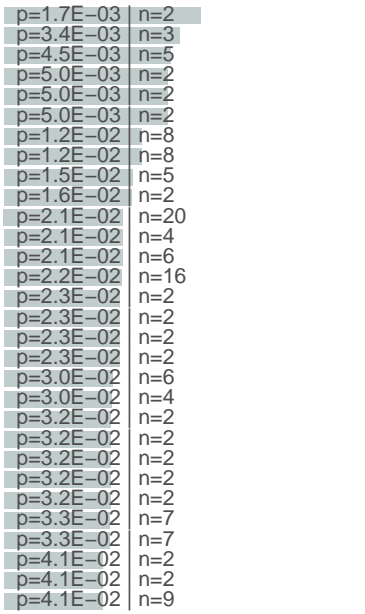
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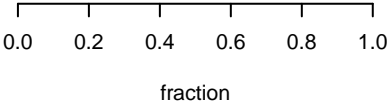
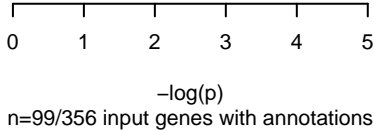
GO:MF
Elav_Nvec_vc1.1_XM_001627049.3

fraction genes in fg and expected value

DNA replication origin binding	GO:0003688
phosphotyrosine residue binding	GO:0001784
DNA-binding transcription repressor acti...	GO:0001227
pyrimidine nucleotide transmembrane tran...	GO:0015218
mRNA 3'-UTR AU-rich region binding	GO:0035925
mRNA regulatory element binding translat...	GO:0000900
cis-regulatory region sequence-specific ...	GO:0000987
RNA polymerase II cis-regulatory region ...	GO:0000978
channel regulator activity	GO:0016247
nucleoside monophosphate kinase activity	GO:0050145
transcription regulator activity	GO:0140110
myosin binding	GO:0017022
RNA polymerase II-specific DNA-binding t...	GO:0061629
DNA-binding transcription factor activit...	GO:0003700
mRNA 5'-UTR binding	GO:0048027
1-acylglycerol-3-phosphate O-acyltransfe...	GO:0003841
aryl hydrocarbon receptor binding	GO:0017162
cyclase activity	GO:0009975
DNA-binding transcription factor binding	GO:0140297
single-stranded DNA binding	GO:0003697
acylglycerol O-acyltransferase activity	GO:0016411
nuclear receptor activity	GO:0004879
adenylate cyclase binding	GO:0008179
ubiquitin-specific protease binding	GO:1990381
ligand-activated transcription factor ac...	GO:0098531
DNA-binding transcription activator acti...	GO:0001228
DNA-binding transcription activator acti...	GO:0001216
lysophosphatidic acid acyltransferase ac...	GO:0042171
lysophospholipid acyltransferase activit...	GO:0071617
transcription cis-regulatory region bind...	GO:0000976



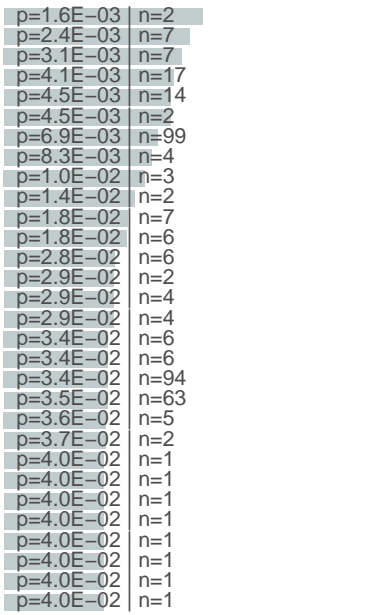
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GO:CC
Elav_Nvec_vc1.1_XM_001627049.3

fraction genes in fg and expected value

Lewy body	GO:0097413
lysosomal membrane	GO:0005765
postsynaptic density	GO:0014069
supramolecular complex	GO:0099080
dendrite	GO:0030425
myelin sheath abaxonal region	GO:0035748
intracellular organelle	GO:0043229
microvillus	GO:0005902
cytoplasmic stress granule	GO:0010494
intrinsic component of the cytoplasmic s...	GO:0031235
contractile fiber	GO:0043292
cytoplasmic side of plasma membrane	GO:0009898
cytoplasmic ribonucleoprotein granule	GO:0036464
excitatory synapse	GO:0060076
azurophil granule	GO:0042582
primary lysosome	GO:0005766
ribonucleoprotein granule	GO:0035770
cytoplasmic side of membrane	GO:0098562
membrane-bounded organelle	GO:0043227
nucleus	GO:0005634
chromosome, centromeric region	GO:0000775
immunological synapse	GO:0001772
microvesicle	GO:1990742
sperm connecting piece	GO:0097224
sperm mitochondrial sheath	GO:0097226
Ndc80 complex	GO:0031262
XPC complex	GO:0071942
apicomedial cortex	GO:0106037
Sec61 translocon complex	GO:0005784
alpha-heterochromatin	GO:0005723



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