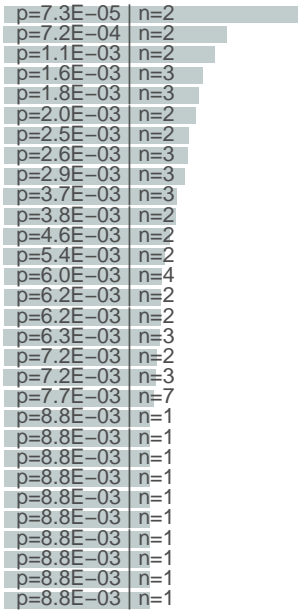


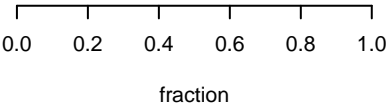
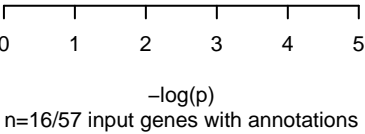
GO:BP
Elav_Nvec_vc1.1_XM_032385745.2

fraction genes in fg and expected value

positive regulation of protein kinase A ...	GO:0010739
cellular response to amyloid-beta	GO:1904646
regulation of urine volume	GO:0035809
positive regulation of cellular amide me...	GO:0034250
neural tube formation	GO:0001841
positive regulation of long-term synapti...	GO:1900273
positive regulation of bone mineralizati...	GO:0030501
mating	GO:0007618
adult locomotory behavior	GO:0008344
associative learning	GO:0008306
dorsal/ventral neural tube patterning	GO:0021904
male courtship behavior	GO:0008049
positive regulation of nervous system pr...	GO:0031646
carbohydrate metabolic process	GO:0005975
cellular response to nerve growth factor...	GO:1990090
negative regulation of endopeptidase act...	GO:0010951
central nervous system neuron developmen...	GO:0021954
midbrain development	GO:0030901
inner ear development	GO:0048839
positive regulation of gene expression	GO:0010628
positive regulation of odontogenesis	GO:0042482
oculomotor nerve formation	GO:0021623
negative regulation of oskar mRNA transl...	GO:0007319
positive regulation of mini excitatory p...	GO:0061885
positive regulation of heart rate by epi...	GO:0001996
positive regulation of nuclear-transcrib...	GO:0060213
positive regulation of AMPA receptor act...	GO:2000969
inhibitory postsynaptic potential	GO:0060080
collateral sprouting in absence of injur...	GO:0048669
positive regulation of tooth mineralizat...	GO:0070172



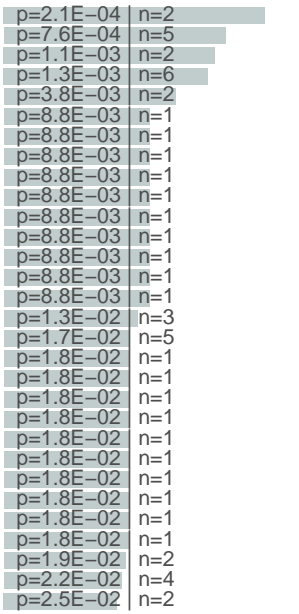
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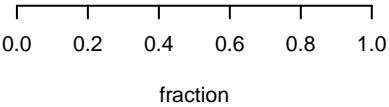
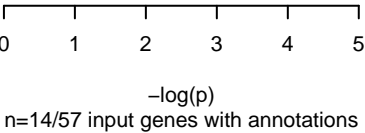
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Elav_Nvec_vc1.1_XM_032385745.2

fraction genes in fg and expected value

mRNA regulatory element binding translat...	GO:0000900
protein heterodimerization activity	GO:0046982
endopeptidase inhibitor activity	GO:0004866
protein homodimerization activity	GO:0042803
mRNA 3'-UTR binding	GO:0003730
acetylcholine receptor activator activit...	GO:0030549
B2 bradykinin receptor binding	GO:0031713
galactokinase activity	GO:0004335
amylin binding	GO:0097645
heparan sulfate binding	GO:1904399
beta1-adrenergic receptor activity	GO:0004940
beta2-adrenergic receptor activity	GO:0004941
oxygen sensor activity	GO:0019826
RNA polymerase II CTD heptapeptide repea...	GO:0008353
dopamine binding	GO:0035240
transcription coactivator activity	GO:0003713
transcription regulator activity	GO:0140110
serine-type endopeptidase inhibitor acti...	GO:0004867
heparan sulfate proteoglycan binding	GO:0043395
PTB domain binding	GO:0051425
cysteine-type endopeptidase inhibitor ac...	GO:0043027
carbon monoxide binding	GO:0070025
nitric oxide binding	GO:0070026
mRNA 3'-UTR AU-rich region binding	GO:0035925
norepinephrine binding	GO:0051380
apolipoprotein binding	GO:0034185
alpha-2A adrenergic receptor binding	GO:0031694
nuclear receptor binding	GO:0016922
DNA-binding transcription factor activit...	GO:0003700
hormone receptor binding	GO:0051427



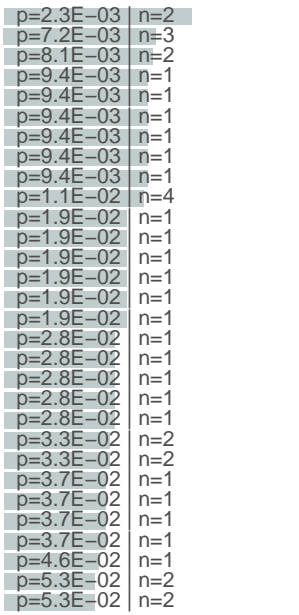
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GO:CC
Elav_Nvec_vc1.1_XM_032385745.2

fraction genes in fg and expected value

MKS complex	GO:0036038
cytoplasmic ribonucleoprotein granule	GO:0036464
clathrin-coated vesicle membrane	GO:0030665
ciliary cap	GO:0061822
transcription factor TFIIF complex	GO:0070985
amyloid-beta complex	GO:0106003
intermediate-density lipoprotein particl...	GO:0034363
high-density lipoprotein particle	GO:0034364
endosome to plasma membrane transport ve...	GO:0070381
extracellular space	GO:0005615
growth cone filopodium	GO:1990812
ciliary rootlet	GO:0035253
platelet alpha granule membrane	GO:0031092
endosome lumen	GO:0031904
cyclin-dependent protein kinase activati...	GO:0019907
nuclear cyclin-dependent protein kinase ...	GO:0019908
astrocyte projection	GO:0097449
glial cell projection	GO:0097386
growth cone lamellipodium	GO:1990761
enzyme activator complex	GO:0150005
polysomal ribosome	GO:0042788
trans-Golgi network membrane	GO:0032588
receptor complex	GO:0043235
cyclin-dependent protein kinase holoenzy...	GO:0000307
guanylate cyclase complex, soluble	GO:0008074
platelet alpha granule lumen	GO:0031093
spindle midzone	GO:0051233
transcription factor TFTC complex	GO:0033276
neuron spine	GO:0044309
dendritic spine	GO:0043197



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