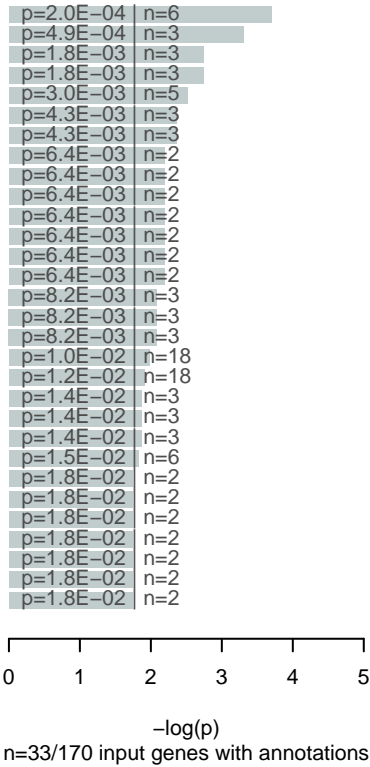


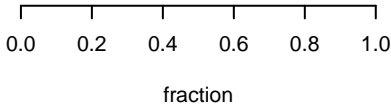
GO:BP
Elav_Nvec_vc1.1_XM_032370984.2

fraction genes in fg and expected value

regulation of synapse organization	GO:0050807
midgut development	GO:0007494
cardiac muscle hypertrophy in response t...	GO:0014898
heart valve morphogenesis	GO:0003179
mononuclear cell differentiation	GO:1903131
ventricular septum development	GO:0003281
regulation of skeletal muscle tissue dev...	GO:0048641
regulation of skeletal muscle cell diffe...	GO:2001014
negative regulation of cAMP-mediated sig...	GO:0043951
response to vitamin	GO:0033273
negative regulation of endothelial cell ...	GO:0001937
cardiac ventricle formation	GO:0003211
negative regulation of cell migration in...	GO:0090051
negative regulation of sprouting angioge...	GO:1903671
positive regulation of dephosphorylation	GO:0035306
negative regulation of small molecule me...	GO:0062014
endothelium development	GO:0003158
cellular macromolecule biosynthetic proc...	GO:0034645
macromolecule biosynthetic process	GO:0009059
regulation of neurotransmitter transport	GO:0051588
regulation of neurotransmitter secretion	GO:0046928
regulation of synapse assembly	GO:0051963
regulation of developmental growth	GO:0048638
imaginal disc-derived wing vein morphoge...	GO:0008586
response to parathyroid hormone	GO:0071107
regulation of cAMP-dependent protein kin...	GO:2000479
embryonic heart tube morphogenesis	GO:0003143
ventricular septum morphogenesis	GO:0060412
heart looping	GO:0001947
chromosome separation	GO:0051304



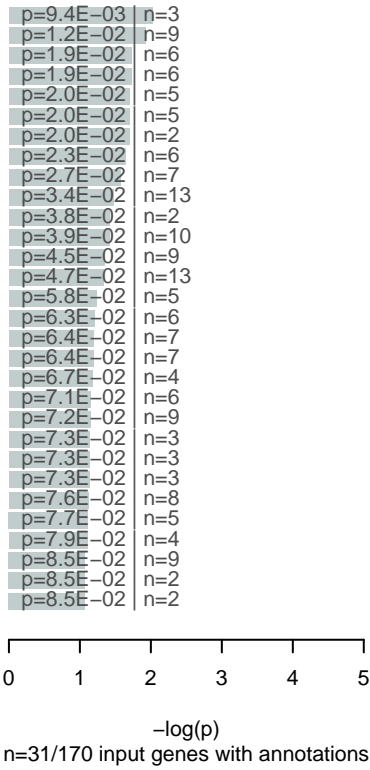
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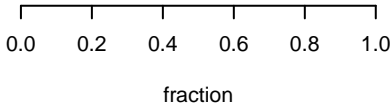
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Elav_Nvec_vc1.1_XM_032370984.2

fraction genes in fg and expected value

protein C-terminus binding	GO:0008022
DNA-binding transcription factor activit...	GO:0000981
DNA-binding transcription activator acti...	GO:0001228
DNA-binding transcription activator acti...	GO:0001216
cis-regulatory region sequence-specific ...	GO:0000987
RNA polymerase II cis-regulatory region ...	GO:0000978
cAMP binding	GO:0030552
metal ion transmembrane transporter acti...	GO:0046873
sequence-specific DNA binding	GO:0043565
heterocyclic compound binding	GO:1901363
cyclic nucleotide binding	GO:0030551
transcription regulator activity	GO:0140110
DNA-binding transcription factor activit...	GO:0003700
organic cyclic compound binding	GO:0097159
cation channel activity	GO:0005261
inorganic cation transmembrane transport...	GO:0022890
kinase binding	GO:0019900
protein kinase binding	GO:0019901
adenyl ribonucleotide binding	GO:0032559
cation transmembrane transporter activit...	GO:0008324
protein dimerization activity	GO:0046983
potassium channel activity	GO:0005267
calcium ion transmembrane transporter ac...	GO:0015085
potassium ion transmembrane transporter ...	GO:0015079
anion binding	GO:0043168
carbohydrate derivative binding	GO:0097367
adenyl nucleotide binding	GO:0030554
nucleic acid binding	GO:0003676
histone deacetylase binding	GO:0042826
adrenergic receptor binding	GO:0031690



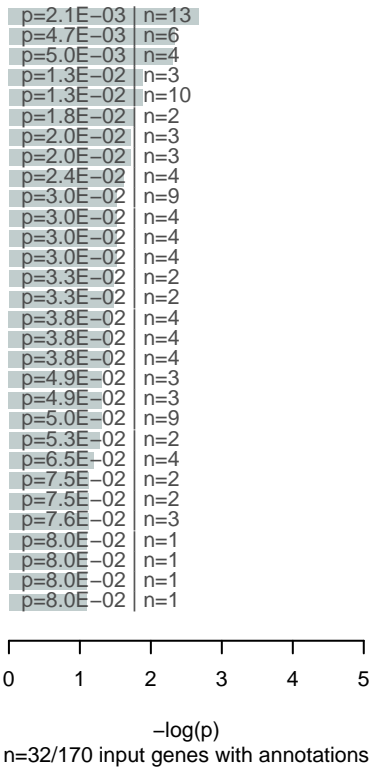
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GO:CC
Elav_Nvec_vc1.1_XM_032370984.2

fraction genes in fg and expected value

cell junction	GO:0030054
plasma membrane protein complex	GO:0098797
plasma membrane raft	GO:0044853
caveola	GO:0005901
synapse	GO:0045202
germ cell nucleus	GO:0043073
voltage-gated potassium channel complex	GO:0008076
potassium channel complex	GO:0034705
cation channel complex	GO:0034703
integral component of plasma membrane	GO:0005887
transporter complex	GO:1990351
transmembrane transporter complex	GO:1902495
ion channel complex	GO:0034702
extrinsic component of cytoplasmic side ...	GO:0031234
T-tubule	GO:0030315
basal plasma membrane	GO:0009925
nuclear chromosome	GO:0000228
basolateral plasma membrane	GO:0016323
neuromuscular junction	GO:0031594
cytoplasmic side of plasma membrane	GO:0009898
intrinsic component of plasma membrane	GO:0031226
extrinsic component of plasma membrane	GO:0019897
basal part of cell	GO:0045178
lipid droplet	GO:0005811
sarcolemma	GO:0042383
cytoplasmic side of membrane	GO:0098562
sperm principal piece	GO:0097228
integrin complex	GO:0008305
RNA polymerase II transcription represso...	GO:0090571
lateral element	GO:0000800



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