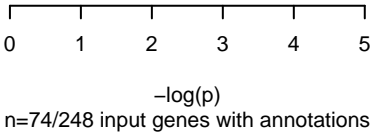
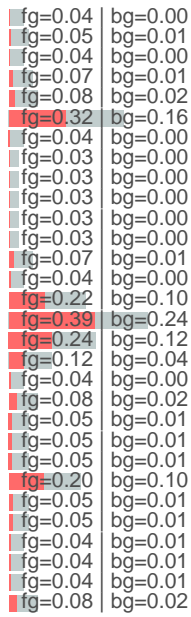
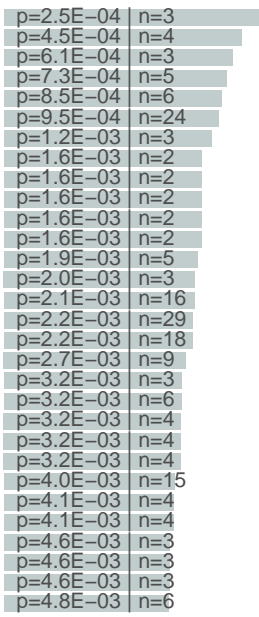


GO:BP
Elav_Nvec_vc1.1_XM_032361813.2

fraction genes in fg and expected value

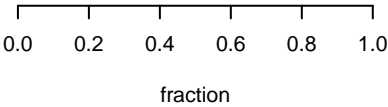
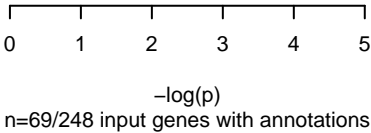
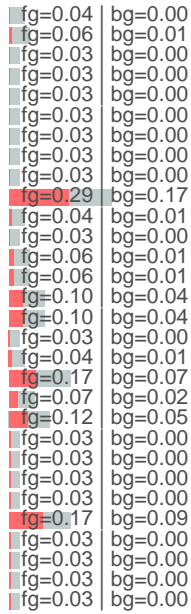
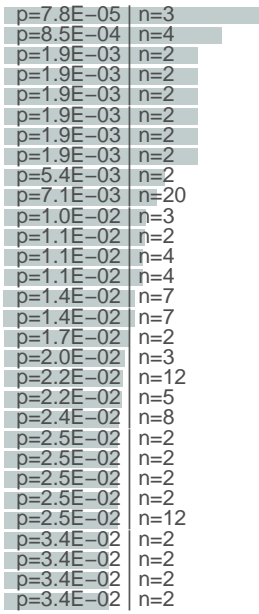
negative regulation of peptidyl-threonin... GO:0010801
circadian regulation of gene expression GO:0032922
G protein-coupled receptor internalizati... GO:0002031
locomotor rhythm GO:0045475
regulation of circadian rhythm GO:0042752
negative regulation of gene expression GO:0010629
positive regulation of cell cycle G2/M p... GO:1902751
follicle-stimulating hormone signaling p... GO:0042699
regulation of dendritic spine maintenanc... GO:1902950
neuroblast fate specification GO:0014018
positive regulation of transforming grow... GO:0071636
negative regulation of protein targeting... GO:0090315
positive regulation of behavior GO:0048520
negative regulation of peptidyl-serine p... GO:0033137
negative regulation of transcription, DN... GO:0045892
regulation of developmental process GO:0050793
negative regulation of cellular macromol... GO:2000113
response to light stimulus GO:0009416
regulation of toll-like receptor signali... GO:0034121
associative learning GO:0008306
positive regulation of mitotic cell cycl... GO:1901992
germarium-derived egg chamber formation GO:0007293
cell chemotaxis GO:0060326
positive regulation of transcription by ... GO:0045944
glycogen metabolic process GO:0005977
neuromuscular synaptic transmission GO:0007274
negative regulation of muscle cell apopt... GO:0010656
inositol phosphate metabolic process GO:0043647
Malpighian tubule development GO:0072002
protein homooligomerization GO:0051260



GO:MF
Elav_Nvec_vc1.1_XM_032361813.2

fraction genes in fg and expected value

cAMP response element binding GO:0035497
double-stranded RNA binding GO:0003725
primary miRNA binding GO:0070878
protein kinase B binding GO:0043422
phosphatidylinositol-3,4,5-trisphosphate... GO:0034485
cAMP response element binding protein bi... GO:0008140
D1 dopamine receptor binding GO:0031748
inositol-polyphosphate 5-phosphatase act... GO:0004445
inositol-1,3,4,5-tetrakisphosphate 5-pho... GO:0052659
identical protein binding GO:0042802
phosphatidylinositol-3,4,5-trisphosphate... GO:0005547
acetylcholine receptor binding GO:0033130
DNA-binding transcription repressor acti... GO:0001227
DNA-binding transcription repressor acti... GO:0001217
DNA-binding transcription activator acti... GO:0001228
DNA-binding transcription activator acti... GO:0001216
arrestin family protein binding GO:1990763
SMAD binding GO:0046332
transcription factor binding GO:0008134
enzyme inhibitor activity GO:0004857
lipid binding GO:0008289
endopeptidase inhibitor activity GO:0004866
cyclin-dependent protein serine/threonin... GO:0016538
potassium channel inhibitor activity GO:0019870
peptidase inhibitor activity GO:0030414
protein domain specific binding GO:0019904
adrenergic receptor binding GO:0031690
N-acetyltransferase activity GO:0008080
DEAD/H-box RNA helicase binding GO:0017151
endopeptidase regulator activity GO:0061135



GO:CC
Elav_Nvec_vc1.1_XM_032361813.2

fraction genes in fg and expected value

euchromatin GO:0000791
transcription regulator complex GO:0005667
cell leading edge GO:0031252
dendritic shaft GO:0043198
perinuclear region of cytoplasm GO:0048471
cytosol GO:0005829
nuclear envelope GO:0005635
axon GO:0030424
postsynapse GO:0098794
protein phosphatase type 1 complex GO:0000164
mitotic spindle pole GO:0097431
actin-based cell projection GO:0098858
distal axon GO:0150034
protein serine/threonine phosphatase com... GO:0008287
pronucleus GO:0045120
neuron spine GO:0044309
dendritic spine GO:0043197
growth cone GO:0030426
filopodium GO:0030175
cell-cell contact zone GO:0044291
site of polarized growth GO:0030427
phosphatase complex GO:1903293
PML body GO:0016605
ruffle GO:0001726
cell junction GO:0030054
endosome to plasma membrane transport ve... GO:0070381
bleb GO:0032059
lipoprotein particle GO:1990777
amyloid-beta complex GO:0106003
eukaryotic translation initiation factor... GO:0016281

