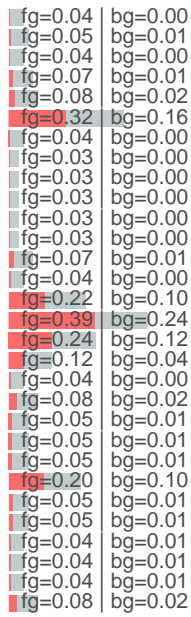
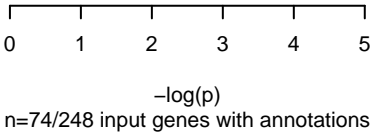
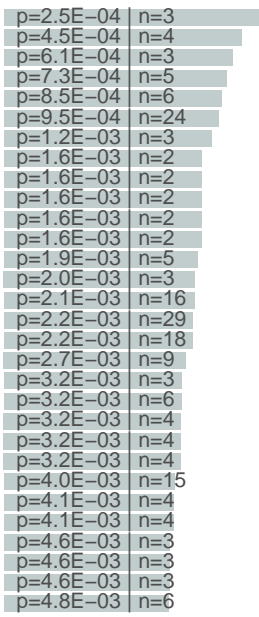


GO:BP  
Elav\_Nvec\_vc1.1\_XM\_048733498.1

fraction genes in fg and expected value

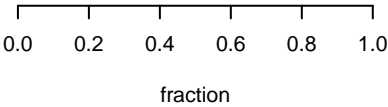
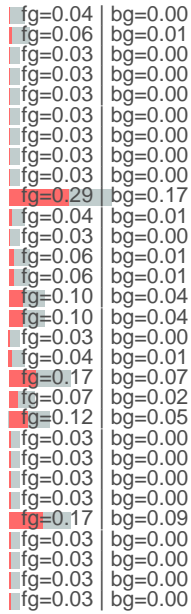
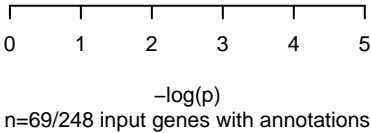
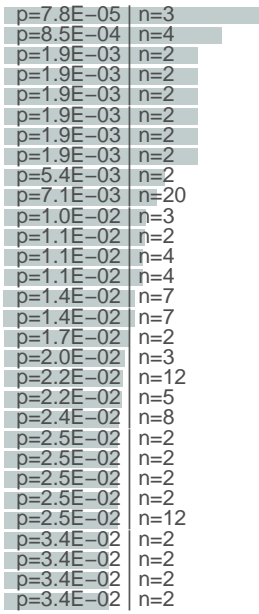
negative regulation of peptidyl-threonin... GO:0010801  
circadian regulation of gene expression GO:0032922  
G protein-coupled receptor internalizati... GO:0002031  
locomotor rhythm GO:0045475  
regulation of circadian rhythm GO:0042752  
negative regulation of gene expression GO:0010629  
positive 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\_048733498.1

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\_048733498.1

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

