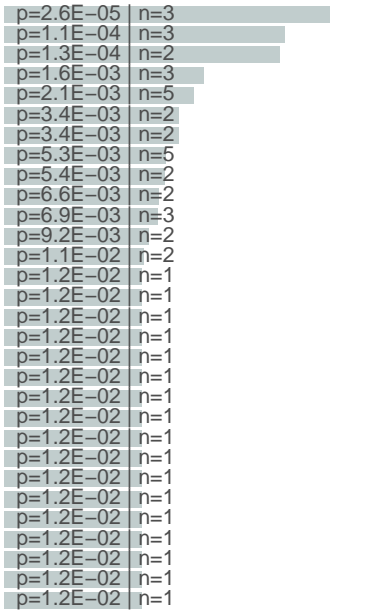


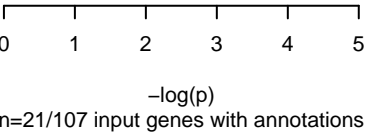
GO:BP  
Elav\_Nvec\_vc1.1\_XM\_048722445.1

fraction genes in fg and expected value

response to amphetamine	GO:0001975
response to bronchodilator	GO:0097366
axon target recognition	GO:0007412
positive regulation of dendrite developm...	GO:1900006
positive regulation of neuron differenti...	GO:0045666
positive regulation of potassium ion tra...	GO:1901381
positive regulation of protein localizat...	GO:1903078
positive regulation of neurogenesis	GO:0050769
long-term synaptic depression	GO:0060292
positive regulation of dendritic spine d...	GO:0060999
positive regulation of synaptic transmis...	GO:0050806
negative regulation of endocytosis	GO:0045806
regulation of synaptic vesicle cycle	GO:0098693
pancreatic juice secretion	GO:0030157
negative regulation of SNARE complex ass...	GO:0035544
regulation of pancreatic juice secretion	GO:0090186
negative regulation of pancreatic juice ...	GO:0090188
lipid tube assembly	GO:0060988
regulation of protein localization to ac...	GO:1904370
positive regulation of protein localizat...	GO:1904372
response to kainic acid	GO:1904373
negative regulation of digestive system ...	GO:0060457
cellular response to morphine	GO:0071315
cellular response to isoquinoline alkalo...	GO:0071317
negative regulation of presynapse assemb...	GO:1905607
protein localization to actin cytoskelet...	GO:1903119
membrane tubulation	GO:0097749
regulation of glutamate secretion, neuro...	GO:1903294
positive regulation of glutamate secreti...	GO:1903296
positive regulation of potassium ion imp...	GO:1903288



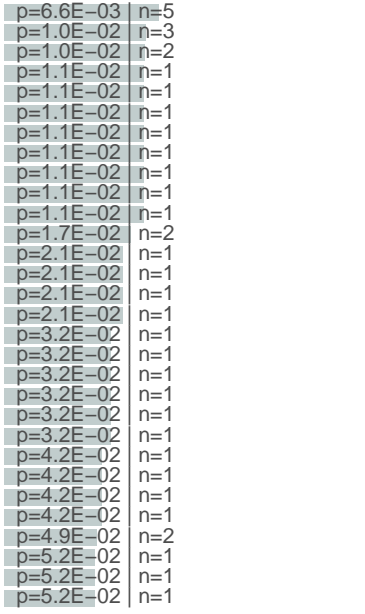
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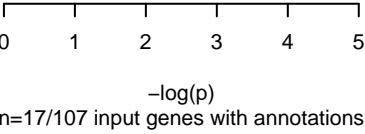
GO:MF  
Elav\_Nvec\_vc1.1\_XM\_048722445.1

fraction genes in fg and expected value

protein kinase binding	GO:0019901
protein C-terminus binding	GO:0008022
transmembrane transporter binding	GO:0044325
K48-linked polyubiquitin modification-de...	GO:0036435
methylmalonyl-CoA mutase activity	GO:0004494
cobalamin binding	GO:0031419
beta-1 adrenergic receptor binding	GO:0031697
U2 snRNA binding	GO:0030620
U5 snRNA binding	GO:0030623
U6 snRNA binding	GO:0017070
G-protein beta-subunit binding	GO:0031681
modification-dependent protein binding	GO:0140030
chloride channel inhibitor activity	GO:0019869
U1 snRNA binding	GO:0030619
second spliceosomal transesterification ...	GO:0000386
syntaxin-3 binding	GO:0030348
calcium-dependent protein kinase activit...	GO:0010857
pre-mRNA intronic binding	GO:0097157
intramolecular transferase activity	GO:0016866
calcium-dependent protein serine/threoni...	GO:0009931
D2 dopamine receptor binding	GO:0031749
neurexin family protein binding	GO:0042043
snRNA binding	GO:0017069
chloride channel regulator activity	GO:0017081
syntaxin-1 binding	GO:0017075
protein phosphatase inhibitor activity	GO:0004864
G protein-coupled receptor binding	GO:0001664
lysophospholipase activity	GO:0004622
lysophospholipid acyltransferase activit...	GO:0071617
phospholipase binding	GO:0043274



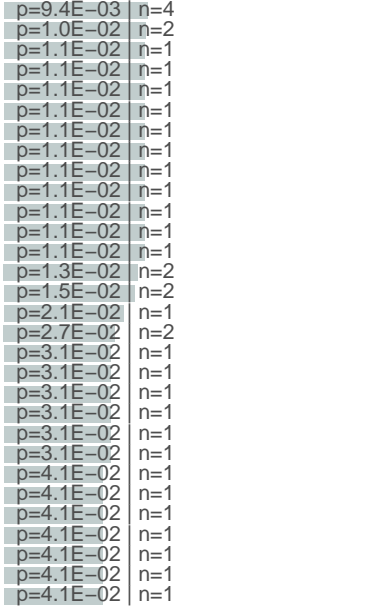
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GO:CC  
Elav\_Nvec\_vc1.1\_XM\_048722445.1

fraction genes in fg and expected value

distal axon	GO:0150034
protein serine/threonine phosphatase com...	GO:0008287
dosage compensation complex	GO:0046536
MSL complex	GO:0072487
X chromosome located dosage compensation...	GO:0016456
cytoplasmic side of dendritic spine plas...	GO:1990780
extrinsic component of presynaptic membr...	GO:0098888
protein phosphatase 4 complex	GO:0030289
dendritic spine neck	GO:0044326
dendritic spine head	GO:0044327
extrinsic component of presynaptic activ...	GO:0098891
hippocampal mossy fiber to CA3 synapse	GO:0098686
extrinsic component of synaptic membrane	GO:0099243
phosphatase complex	GO:1903293
presynaptic active zone	GO:0048786
U5 snRNP	GO:0005682
presynaptic membrane	GO:0042734
zymogen granule membrane	GO:0042589
growth cone lamellipodium	GO:1990761
dendritic spine membrane	GO:0032591
H4/H2A histone acetyltransferase complex	GO:0043189
cytolytic granule	GO:0044194
NuA4 histone acetyltransferase complex	GO:0035267
dendrite terminus	GO:0044292
X chromosome	GO:0000805
fungus-type vacuole	GO:0000324
fungus-type vacuole membrane	GO:0000329
presynaptic periaxonal zone	GO:0036062
chromocenter	GO:0010369
basal dendrite	GO:0097441



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