

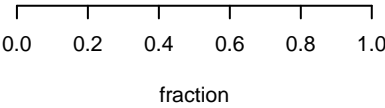
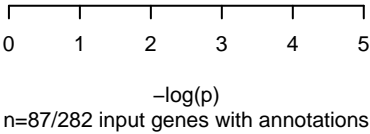
GO:BP
Elav_Nvec_vc1.1_XM_032363951.2

fraction genes in fg and expected value

detection of endogenous stimulus	GO:0009726
cellular amino acid biosynthetic process	GO:0008652
carbohydrate derivative transport	GO:1901264
glutamine family amino acid metabolic pr...	GO:0009064
protein hexamerization	GO:0034214
response to antipsychotic drug	GO:0097332
response to clozapine	GO:0097338
presynaptic active zone organization	GO:1990709
positive regulation of cellular response...	GO:1900409
negative regulation of stem cell prolife...	GO:2000647
positive regulation of cation transmembr...	GO:1904064
negative regulation of hydrogen peroxide...	GO:1903206
negative regulation of cellular response...	GO:2001039
cell cycle G1/S phase transition	GO:0044843
regulation of cell cycle G1/S phase tran...	GO:1902806
regulation of cell cycle phase transitio...	GO:1901987
protein ubiquitination	GO:0016567
cellular response to endogenous stimulus	GO:0071495
cellular response to organic substance	GO:0071310
positive regulation of cell cycle proces...	GO:0090068
positive regulation of cation channel ac...	GO:2001259
calcium ion import	GO:0070509
regulation of intrinsic apoptotic signal...	GO:2001242
response to hydrostatic pressure	GO:0051599
protein-lipid complex subunit organizati...	GO:0071825
gamma-aminobutyric acid signaling pathwa...	GO:0007214
export across plasma membrane	GO:0140115
pyrimidine-containing compound transmemb...	GO:0072531
2-oxoglutarate metabolic process	GO:0006103
positive regulation of necrotic cell dea...	GO:0010940

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p=4.0E-03	n=4
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p=5.9E-03	n=2
p=7.3E-03	n=3
p=8.7E-03	n=4
p=9.6E-03	n=2
p=9.6E-03	n=2
p=1.1E-02	n=7
p=1.1E-02	n=5
p=1.2E-02	n=10
p=1.2E-02	n=12
p=1.3E-02	n=17
p=1.3E-02	n=24
p=1.3E-02	n=7
p=1.3E-02	n=3
p=1.3E-02	n=3
p=1.4E-02	n=5
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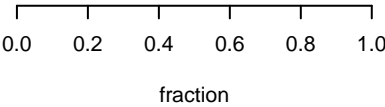
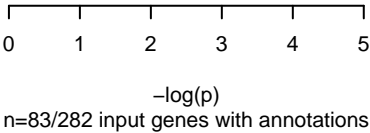
GO:MF
Elav_Nvec_vc1.1_XM_032363951.2

fraction genes in fg and expected value

transaminase activity	GO:0008483
BH domain binding	GO:0051400
mitogen-activated protein kinase p38 bin...	GO:0048273
outward rectifier potassium channel acti...	GO:0015271
ankyrin binding	GO:0030506
steroid binding	GO:0005496
nucleobase-containing compound transmemb...	GO:0015932
ubiquitin protein ligase binding	GO:0031625
JUN kinase binding	GO:0008432
calcium-activated potassium channel acti...	GO:0015269
ion gated channel activity	GO:0022839
ADP binding	GO:0043531
anion binding	GO:0043168
mitogen-activated protein kinase kinase ...	GO:0031434
amino acid binding	GO:0016597
solute:cation symporter activity	GO:0015294
protein kinase binding	GO:0019901
cytoskeletal protein binding	GO:0008092
small molecule binding	GO:0036094
adrenergic receptor binding	GO:0031690
cholesterol binding	GO:0015485
transmembrane transporter binding	GO:0044325
symporter activity	GO:0015293
vitamin B6 binding	GO:0070279
pyridoxal phosphate binding	GO:0030170
sterol binding	GO:0032934
carbohydrate derivative transmembrane tr...	GO:1901505
secondary active transmembrane transport...	GO:0015291
kinase binding	GO:0019900
phospholipid binding	GO:0005543

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p=7.7E-04	n=3
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p=2.1E-03	n=3
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p=7.3E-03	n=3
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p=7.3E-03	n=3
p=8.6E-03	n=8
p=1.1E-02	n=2
p=1.1E-02	n=2
p=1.2E-02	n=6
p=1.4E-02	n=3
p=1.4E-02	n=18
p=1.7E-02	n=2
p=1.7E-02	n=3
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p=2.0E-02	n=13
p=2.1E-02	n=14
p=2.3E-02	n=15
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p=3.2E-02	n=13
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GO:CC
Elav_Nvec_vc1.1_XM_032363951.2

fraction genes in fg and expected value

cytoplasmic side of endoplasmic reticulu...	GO:0098554
polytene chromosome puff	GO:0005703
dendritic spine membrane	GO:0032591
growth cone lamellipodium	GO:1990761
T-tubule	GO:0030315
cation channel complex	GO:0034703
intrinsic component of endoplasmic retic...	GO:0031227
methylosome	GO:0034709
cell cortex	GO:0005938
neuromuscular junction	GO:0031594
neuronal cell body membrane	GO:0032809
actomyosin	GO:0042641
cell body membrane	GO:0044298
sarcomere	GO:0030017
main axon	GO:0044304
contractile fiber	GO:0043292
cytoplasmic side of membrane	GO:0098562
unconventional myosin complex	GO:0016461
myosin complex	GO:0016459
integral component of endoplasmic reticu...	GO:0030176
cytoplasmic side of plasma membrane	GO:0009898
calcium channel complex	GO:0034704
neuronal cell body	GO:0043025
mitochondrial proton-transporting ATP sy...	GO:0005756
spectrosome	GO:0045170
mitochondrial proton-transporting ATP sy...	GO:0000276
L-type voltage-gated calcium channel com...	GO:1990454
dendritic spine head	GO:0044327
spectrin	GO:0008091
anchored component of the cytoplasmic si...	GO:0098753

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p=5.9E-03	n=2
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p=9.1E-03	n=6
p=9.6E-03	n=2
p=9.7E-03	n=8
p=1.1E-02	n=5
p=1.6E-02	n=3
p=1.6E-02	n=3
p=1.9E-02	n=3
p=1.9E-02	n=5
p=2.2E-02	n=3
p=2.2E-02	n=6
p=2.5E-02	n=8
p=2.5E-02	n=2
p=2.5E-02	n=2
p=2.8E-02	n=5
p=2.8E-02	n=5
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