

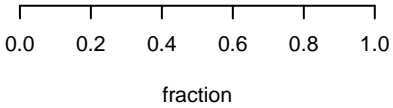
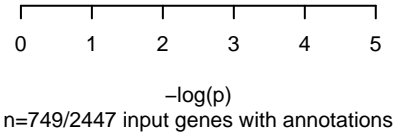
GO:BP
Elav_Nvec_vc1.1_XM_048723945.1

fraction genes in fg and expected value

regulation of oocyte development	GO:0060281
blood vessel remodeling	GO:0001974
macropinocytosis	GO:0044351
pole cell development	GO:0007277
response to platelet aggregation inhibit...	GO:0061478
positive regulation of oogenesis	GO:1905881
cellular response to glucagon stimulus	GO:0071377
positive regulation of gluconeogenesis	GO:0045722
mRNA polyadenylation	GO:0006378
positive regulation of cell maturation	GO:1903431
regulation of nitric-oxide synthase acti...	GO:0050999
photoreceptor cell maintenance	GO:0045494
immune response-regulating signaling pat...	GO:0002764
cellularization	GO:0007349
glutamine family amino acid metabolic pr...	GO:0009064
leukocyte activation involved in immune ...	GO:0002366
response to ketone	GO:1901654
response to antipsychotic drug	GO:0097332
response to clozapine	GO:0097338
modulation by symbiont of host cellular ...	GO:0044068
cellular response to forskolin	GO:1904322
regulation of spontaneous synaptic trans...	GO:0150003
regulation of protein localization to ce...	GO:1904776
negative regulation of histone acetylati...	GO:0035067
glomerulus morphogenesis	GO:0072102
lymphocyte differentiation	GO:0030098
cell differentiation involved in kidney ...	GO:0061005
mushroom body development	GO:0016319
regulation of embryonic pattern specific...	GO:1902875
B cell proliferation	GO:0042100

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p=8.8E-04	n=9
p=1.6E-03	n=5
p=1.6E-03	n=5
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p=2.4E-03	n=6
p=2.4E-03	n=6
p=2.4E-03	n=12
p=2.4E-03	n=8
p=2.6E-03	n=7
p=3.0E-03	n=11
p=4.0E-03	n=28
p=4.2E-03	n=14
p=4.2E-03	n=12
p=5.1E-03	n=50
p=5.5E-03	n=28
p=5.9E-03	n=4
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p=6.1E-03	n=25
p=6.9E-03	n=10
p=7.0E-03	n=12
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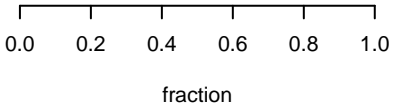
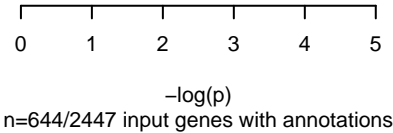
GO:MF
Elav_Nvec_vc1.1_XM_048723945.1

fraction genes in fg and expected value

mRNA binding	GO:0003729
translation repressor activity	GO:0030371
regulatory RNA binding	GO:0061980
ATPase binding	GO:0051117
mRNA 3'-UTR binding	GO:0003730
ubiquitin-like protein ligase binding	GO:0044389
solute:anion antiporter activity	GO:0140323
ligand-activated transcription factor ac...	GO:0098531
inorganic anion exchanger activity	GO:0005452
anion:anion antiporter activity	GO:0015301
nuclear receptor activity	GO:0004879
miRNA binding	GO:0035198
mRNA regulatory element binding translat...	GO:0000900
Wnt-activated receptor activity	GO:0042813
oxalate transmembrane transporter activi...	GO:0019531
formate transmembrane transporter activi...	GO:0015499
sulfate transmembrane transporter activi...	GO:0015116
tRNA (cytosine) methyltransferase activi...	GO:0016427
cytoskeletal protein binding	GO:0008092
transcription corepressor activity	GO:0003714
protein kinase binding	GO:0019901
transmembrane transporter binding	GO:0044325
antiporter activity	GO:0015297
nucleotide binding	GO:0000166
nucleoside phosphate binding	GO:1901265
purine nucleotide binding	GO:0017076
manganese ion binding	GO:0030145
ubiquitin protein ligase binding	GO:0031625
microtubule motor activity	GO:0003777
oxidoreductase activity, acting on CH-OH...	GO:0016614

p=2.4E-03	n=32
p=5.4E-03	n=4
p=5.4E-03	n=4
p=8.1E-03	n=14
p=1.3E-02	n=9
p=1.7E-02	n=34
p=1.8E-02	n=5
p=1.8E-02	n=5
p=1.8E-02	n=5
p=1.8E-02	n=5
p=2.0E-02	n=3
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p=2.0E-02	n=3
p=2.1E-02	n=64
p=2.5E-02	n=16
p=2.5E-02	n=56
p=3.2E-02	n=9
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p=3.3E-02	n=64
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p=3.6E-02	n=31
p=3.7E-02	n=7
p=3.8E-02	n=11

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GO:CC
Elav_Nvec_vc1.1_XM_048723945.1

fraction genes in fg and expected value

protein-containing complex	GO:0032991
photoreceptor inner segment	GO:0001917
centrosome	GO:0005813
sarcolemma	GO:0042383
cilium	GO:0005929
mitotic spindle	GO:0072686
female germ cell nucleus	GO:0001674
non-membrane-bounded organelle	GO:0043228
intracellular non-membrane-bounded organ...	GO:0043232
mitotic spindle pole	GO:0097431
motile cilium	GO:0031514
ciliary plasm	GO:0097014
germinal vesicle	GO:0042585
fibrillar center	GO:0001650
integral component of cytoplasmic side o...	GO:0071458
cAMP-dependent protein kinase complex	GO:0005952
polytene chromosome chromocenter	GO:0005701
ribonucleoprotein complex	GO:1990904
plasma membrane bounded cell projection ...	GO:0032838
axoneme	GO:0005930
nuclear lumen	GO:0031981
photoreceptor connecting cilium	GO:0032391
stress fiber	GO:0001725
contractile actin filament bundle	GO:0097517
90S preribosome	GO:0030686
secretory granule	GO:0030141
large ribosomal subunit	GO:0015934
membrane-enclosed lumen	GO:0031974
intracellular organelle lumen	GO:0070013
organelle lumen	GO:0043233

p=5.6E-05	n=361
p=2.1E-03	n=9
p=2.4E-03	n=54
p=6.2E-03	n=14
p=6.6E-03	n=61
p=6.8E-03	n=18
p=7.1E-03	n=6
p=1.3E-02	n=283
p=1.3E-02	n=283
p=1.3E-02	n=7
p=1.5E-02	n=23
p=1.6E-02	n=16
p=2.0E-02	n=5
p=2.0E-02	n=15
p=2.1E-02	n=3
p=2.1E-02	n=3
p=2.2E-02	n=4
p=2.3E-02	n=80
p=2.3E-02	n=19
p=2.7E-02	n=15
p=2.9E-02	n=276
p=3.0E-02	n=8
p=3.1E-02	n=6
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p=3.1E-02	n=6
p=3.3E-02	n=49
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