

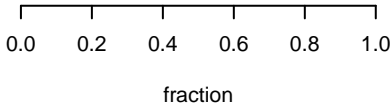
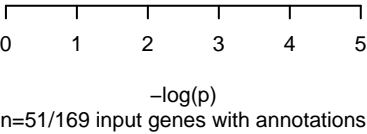
GO:BP
Elav_Nvec_vc1.1_XM_032380190.2

fraction genes in fg and expected value

salivary gland morphogenesis	GO:0007435
response to thyroid hormone	GO:0097066
cellular response to staurosporine	GO:0072734
ovarian follicle cell stalk formation	GO:0030713
negative regulation of erythrocyte diffe...	GO:0045647
melanotic encapsulation of foreign targe...	GO:0035011
ciliary basal body-plasma membrane docki...	GO:0097711
hemocyte migration	GO:0035099
ectodermal placode formation	GO:0060788
cytoplasmic transport, nurse cell to ooc...	GO:0007303
maintenance of cell polarity	GO:0030011
wound healing	GO:0042060
negative regulation of microtubule depol...	GO:0007026
regulation of clathrin-dependent endocyt...	GO:2000369
IRE1-mediated unfolded protein response	GO:0036498
regulation of microtubule depolymerizati...	GO:0031114
positive regulation of anion transport	GO:1903793
pigment cell differentiation	GO:0050931
calcium ion regulated lysosome exocytosi...	GO:1990927
second mitotic wave involved in compound...	GO:0016330
deoxyribonucleoside monophosphate metabo...	GO:0009162
deoxyribonucleoside monophosphate biosyn...	GO:0009157
pyrimidine deoxyribonucleoside monophosp...	GO:0009176
pyrimidine deoxyribonucleoside monophosp...	GO:0009177
negative regulation of synaptic vesicle ...	GO:1900243
intracellular sequestering of iron ion	GO:0006880
indole metabolic process	GO:0042431
hindbrain tangential cell migration	GO:0021934
cerebellar granule cell precursor tangen...	GO:0021935
2'-deoxyribonucleotide biosynthetic proc...	GO:0009265

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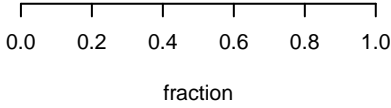
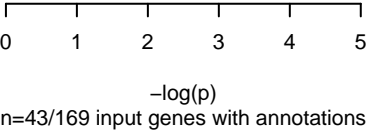
GO:MF
Elav_Nvec_vc1.1_XM_032380190.2

fraction genes in fg and expected value

protein binding	GO:0005515
aspartic-type endopeptidase activity	GO:0004190
hydrolase activity, acting on carbon-nit...	GO:0016814
CCR chemokine receptor binding	GO:0048020
hydroxymethyl-, formyl- and related tran...	GO:0016742
phosphoribosylaminoimidazolecarboxamide ...	GO:0004643
JUN kinase kinase kinase activity	GO:0004706
phosphopantothenate--cysteine ligase act...	GO:0004632
IMP cyclohydrolase activity	GO:0003937
apolipoprotein receptor binding	GO:0034190
GBD domain binding	GO:0032427
apolipoprotein A-I receptor binding	GO:0034191
protein tyrosine phosphatase activity, m...	GO:0030946
RNA polymerase I activity	GO:0001054
Rho GDP-dissociation inhibitor binding	GO:0051022
CCR5 chemokine receptor binding	GO:0031730
BIR domain binding	GO:1990525
dCMP deaminase activity	GO:0004132
clathrin binding	GO:0030276
identical protein binding	GO:0042802
cysteine-type endopeptidase activity	GO:0004197
chaperone binding	GO:0051087
unfolded protein binding	GO:0051082
phosphate ion binding	GO:0042301
cyclohydrolase activity	GO:0019238
beta-2 adrenergic receptor binding	GO:0031698
sodium:phosphate symporter activity	GO:0005436
cysteine-type endopeptidase activity inv...	GO:0097200
GDP-dissociation inhibitor binding	GO:0051021
aspartic endopeptidase activity, intrame...	GO:0042500

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GO:CC
Elav_Nvec_vc1.1_XM_032380190.2

fraction genes in fg and expected value

dendritic spine	GO:0043197
proteasome regulatory particle, base sub...	GO:0008540
melanosome	GO:0042470
pigment granule	GO:0048770
intermediate filament cytoskeleton	GO:0045111
centriole	GO:0005814
dense core granule membrane	GO:0032127
BLOC-3 complex	GO:0031085
microvesicle	GO:1990742
platelet dense granule	GO:0042827
precatalytic spliceosome	GO:0071011
vacuole	GO:0005773
smooth endoplasmic reticulum	GO:0005790
T-tubule	GO:0030315
actin filament	GO:0005884
PAN complex	GO:0031251
dendritic spine neck	GO:0044326
secretory granule	GO:0030141
cytoskeleton	GO:0005856
spliceosomal complex	GO:0005681
myelin sheath	GO:0043209
microtubule cytoskeleton	GO:0015630
integral component of cytoplasmic side o...	GO:0071458
inhibitory synapse	GO:0060077
dense core granule	GO:0031045
early phagosome	GO:0032009
Hrd1p ubiquitin ligase ERAD-L complex	GO:0000839
cAMP-dependent protein kinase complex	GO:0005952
late endosome membrane	GO:0031902
P-body	GO:0000932

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