

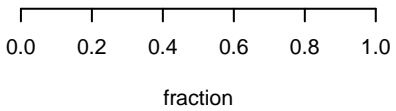
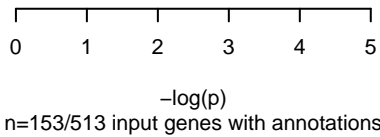
GO:BP
Elav_Nvec_vc1.1_XM_048723524.1

fraction genes in fg and expected value

signal complex assembly	GO:0007172
cellular protein metabolic process	GO:0044267
proximal/distal pattern formation	GO:0009954
negative regulation of Rho protein signa...	GO:0035024
retinoic acid receptor signaling pathway	GO:0048384
maintenance of epithelial integrity, ope...	GO:0035160
positive regulation of gastrulation	GO:2000543
determination of adult lifespan	GO:0008340
positive regulation of macromolecule met...	GO:0010604
cell adhesion mediated by integrin	GO:0033627
nematode larval development	GO:0002119
positive regulation of nervous system de...	GO:0051962
multicellular organism aging	GO:0010259
endoderm formation	GO:0001706
positive regulation of transforming grow...	GO:0030511
B cell differentiation	GO:0030183
epithelial tube morphogenesis	GO:0060562
regulation of execution phase of apoptos...	GO:1900117
insulin receptor signaling pathway	GO:0008286
positive regulation of nitrogen compound...	GO:0051173
endothelial cell migration	GO:0043542
lymphocyte activation involved in immune...	GO:0002285
axon regeneration	GO:0031103
PERK-mediated unfolded protein response	GO:0036499
tumor necrosis factor-mediated signaling...	GO:0033209
cellular catabolic process	GO:0044248
response to herbicide	GO:0009635
negative regulation of mitochondrial out...	GO:1901029
negative regulation of mitochondrial dep...	GO:0051902
compartment pattern specification	GO:0007386

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p=3.3E-03	n=13
p=3.3E-03	n=54
p=3.4E-03	n=4
p=4.3E-03	n=13
p=4.6E-03	n=17
p=4.6E-03	n=13
p=4.7E-03	n=4
p=5.3E-03	n=3
p=5.3E-03	n=5
p=5.5E-03	n=20
p=6.3E-03	n=4
p=7.2E-03	n=7
p=7.6E-03	n=50
p=8.0E-03	n=6
p=8.0E-03	n=5
p=8.0E-03	n=5
p=8.1E-03	n=3
p=8.3E-03	n=4
p=8.7E-03	n=42
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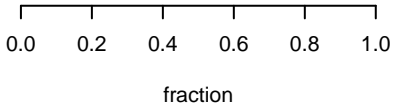
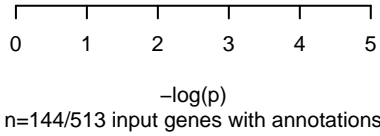
GO:MF
Elav_Nvec_vc1.1_XM_048723524.1

fraction genes in fg and expected value

protease binding	GO:0002020
integrin binding	GO:0005178
Notch binding	GO:0005112
vinculin binding	GO:0017166
nuclear retinoic acid receptor binding	GO:0042974
ubiquitin protein ligase binding	GO:0031625
metalloendopeptidase activity	GO:0004222
transcription coregulator binding	GO:0001221
ubiquitin-like protein ligase binding	GO:0044389
beta-catenin binding	GO:0008013
iodide transmembrane transporter activit...	GO:0015111
temperature-gated ion channel activity	GO:0097603
WW domain binding	GO:0050699
protein methyltransferase activity	GO:0008276
DNA-binding transcription repressor acti...	GO:0001227
DNA-binding transcription repressor acti...	GO:0001217
endopeptidase activity	GO:0004175
protein domain specific binding	GO:0019904
protein-lysine N-methyltransferase activ...	GO:0016279
transcription coregulator activity	GO:0003712
lysine N-methyltransferase activity	GO:0016278
hydro-lyase activity	GO:0016836
SH2 domain binding	GO:0042169
eukaryotic initiation factor eIF2 bindin...	GO:0071074
phospholipid scramblase activity	GO:0017128
JUN kinase binding	GO:0008432
transcription coactivator binding	GO:0001223
arrestin family protein binding	GO:1990763
Hsp70 protein binding	GO:0030544
nuclear androgen receptor binding	GO:0050681

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p=1.4E-02	n=3
p=1.4E-02	n=3
p=1.8E-02	n=11
p=1.9E-02	n=3
p=2.0E-02	n=3
p=2.0E-02	n=2
p=2.0E-02	n=2
p=2.1E-02	n=4
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p=2.1E-02	n=5
p=2.2E-02	n=6
p=2.5E-02	n=19
p=2.5E-02	n=3
p=2.6E-02	n=13
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p=3.2E-02	n=3
p=3.2E-02	n=3
p=3.3E-02	n=2
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p=3.5E-02	n=4
p=4.0E-02	n=3

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GO:CC
Elav_Nvec_vc1.1_XM_048723524.1

fraction genes in fg and expected value

endoplasmic reticulum quality control co...	GO:0044322
ribosome	GO:0005840
integral component of mitochondrial memb...	GO:0032592
integral component of Golgi membrane	GO:0030173
ribonucleoprotein complex	GO:1990904
focal adhesion	GO:0005925
Cul3-RING ubiquitin ligase complex	GO:0031463
stress fiber	GO:0001725
contractile actin filament bundle	GO:0097517
basal part of cell	GO:0045178
luminal side of endoplasmic reticulum me...	GO:0098553
integral component of luminal side of en...	GO:0071556
cortical endoplasmic reticulum	GO:0032541
organellar large ribosomal subunit	GO:0000315
mitochondrial large ribosomal subunit	GO:0005762
actin filament bundle	GO:0032432
basolateral plasma membrane	GO:0016323
basal plasma membrane	GO:0009925
intercellular canalliculus	GO:0046581
luminal side of membrane	GO:0098576
receptor complex	GO:0043235
intracellular membrane-bounded organelle	GO:0043231
intrinsic component of mitochondrial inn...	GO:0031304
integral component of mitochondrial inne...	GO:0031305
filopodium membrane	GO:0031527
cell surface	GO:0009986
cytosolic ribosome	GO:0022626
membrane-bounded organelle	GO:0043227
ribosomal subunit	GO:0044391
cytoplasmic ubiquitin ligase complex	GO:0000153

p=6.8E-04	n=3
p=1.6E-03	n=11
p=3.4E-03	n=4
p=4.7E-03	n=4
p=7.2E-03	n=23
p=8.0E-03	n=5
p=1.2E-02	n=3
p=1.6E-02	n=3
p=1.6E-02	n=3
p=1.7E-02	n=9
p=1.8E-02	n=2
p=1.8E-02	n=2
p=1.8E-02	n=2
p=2.1E-02	n=3
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p=2.1E-02	n=3
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p=3.2E-02	n=6
p=3.7E-02	n=129
p=4.1E-02	n=2
p=4.1E-02	n=2
p=4.1E-02	n=2
p=4.3E-02	n=10
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p=5.4E-02	n=7
p=5.6E-02	n=2

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