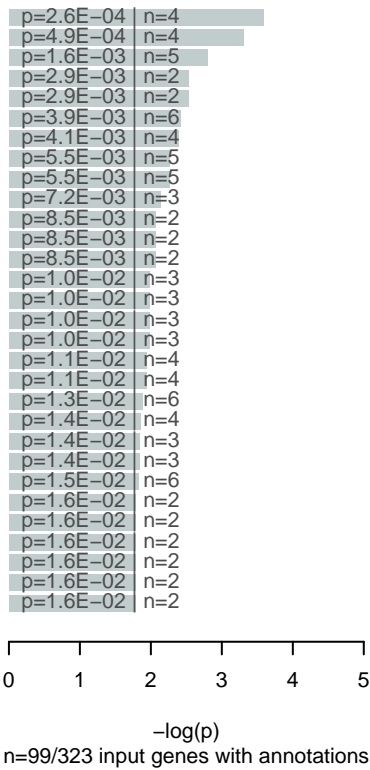


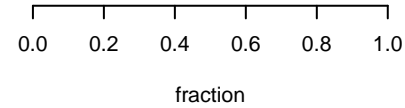
GO:BP
Elav_Nvec_vc1.1_XM_001629605.3

fraction genes in fg and expected value

positive regulation of morphogenesis of ...	GO:1905332
positive regulation of epithelial cell d...	GO:0030858
mRNA polyadenylation	GO:0006378
oviduct development	GO:0060066
positive regulation of inclusion body as...	GO:0090261
hindbrain development	GO:0030902
heterochromatin assembly	GO:0031507
positive regulation of chromatin organiz...	GO:1905269
molting cycle process	GO:0022404
histone monoubiquitination	GO:0010390
regulation of nephron tubule epithelial ...	GO:0072182
nephric duct morphogenesis	GO:0072178
regulation of histone phosphorylation	GO:0033127
intra-Golgi vesicle-mediated transport	GO:0006891
endoderm formation	GO:0001706
regulation of histone H3-K4 methylation	GO:0051569
regulation of morphogenesis of a branchi...	GO:0060688
histone H3-K4 methylation	GO:0051568
hair cycle process	GO:0022405
positive regulation of DNA metabolic pro...	GO:0051054
hair cycle	GO:0042633
regulation of epithelial cell apoptotic ...	GO:1904035
regulation of telomere maintenance	GO:0032204
retina development in camera-type eye	GO:0060041
regulation of heterochromatin organizati...	GO:0120261
metanephric renal vesicle morphogenesis	GO:0072283
retina layer formation	GO:0010842
regulation of heterochromatin assembly	GO:0031445
renal vesicle development	GO:0072087
renal vesicle morphogenesis	GO:0072077



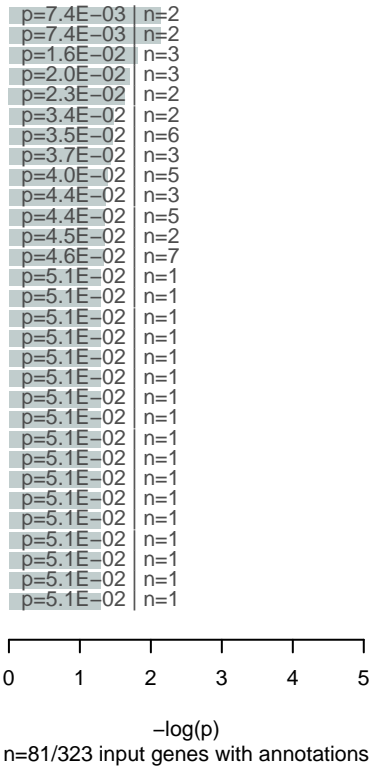
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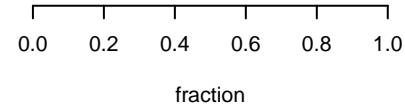
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Elav_Nvec_vc1.1_XM_001629605.3

fraction genes in fg and expected value

hexosaminidase activity	GO:0015929
NAD binding	GO:0051287
ubiquitin conjugating enzyme activity	GO:0061631
ubiquitin-like protein conjugating enzym...	GO:0061650
14-3-3 protein binding	GO:0071889
structural constituent of cytoskeleton	GO:0005200
mRNA binding	GO:0003729
transmembrane transporter binding	GO:0044325
RNA polymerase II-specific DNA-binding t...	GO:0061629
core promoter sequence-specific DNA bind...	GO:0001046
GTPase activator activity	GO:0005096
nuclear estrogen receptor binding	GO:0030331
phospholipid binding	GO:0005543
opsin binding	GO:0002046
NAD-dependent histone deacetylase activi...	GO:0046969
sphingolipid transfer activity	GO:0120016
ceramide transfer activity	GO:0120017
1-alkenylglycerophosphocholine O-acyltra...	GO:0047159
ferrochelataase activity	GO:0004325
steroid dehydrogenase activity	GO:0016229
histone methyltransferase binding	GO:1990226
protein phosphatase 5 binding	GO:1990634
mannosyl-oligosaccharide mannosidase act...	GO:0015924
estradiol 17-beta-dehydrogenase activity	GO:0004303
alpha-N-acetylglucosaminidase activity	GO:0004561
NAD+ binding	GO:0070403
NAD-dependent protein deacetylase activi...	GO:0034979
alpha-galactosidase activity	GO:0004557
V2 vasopressin receptor binding	GO:0031896
NAD-dependent histone deacetylase activi...	GO:0017136



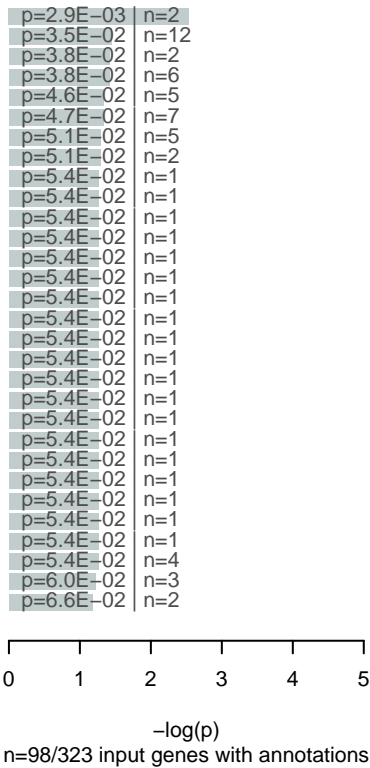
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GO:CC
Elav_Nvec_vc1.1_XM_001629605.3

fraction genes in fg and expected value

mRNA cleavage and polyadenylation specif...	GO:0005847
vacuole	GO:0005773
ficolin-1-rich granule membrane	GO:0101003
lytic vacuole membrane	GO:0098852
nuclear membrane	GO:0031965
vacuolar membrane	GO:0005774
secretory granule membrane	GO:0030667
PML body	GO:0016605
Cdc73/Paf1 complex	GO:0016593
piP-body	GO:0071547
beta-catenin-TCF7L2 complex	GO:0070369
platelet dense granule	GO:0042827
luminal side of Golgi membrane	GO:0098547
clathrin-sculpted gamma-aminobutyric aci...	GO:0061200
clathrin-sculpted gamma-aminobutyric aci...	GO:0061202
collagen and cuticulin-based cuticle ext...	GO:0060102
beta-catenin destruction complex	GO:0030877
nuclear outer membrane	GO:0005640
catenin-TCF7L2 complex	GO:0071664
protein kinase CK2 complex	GO:0005956
BLOC-3 complex	GO:0031085
fascia adherens	GO:0005916
Scrib-APC-beta-catenin complex	GO:0034750
spectrin-associated cytoskeleton	GO:0014731
nuclear pore inner ring	GO:0044611
clathrin-sculpted vesicle	GO:0060198
catenin complex	GO:0016342
ciliary basal body	GO:0036064
heterochromatin	GO:0000792
MKS complex	GO:0036038



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