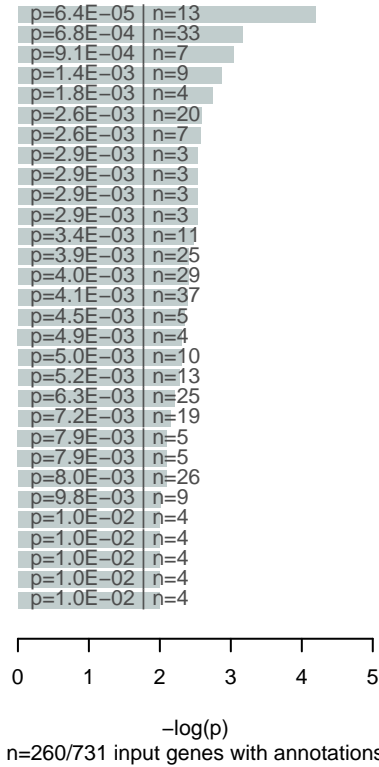


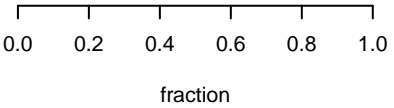
GO:BP
Fox_Nvec_vc1.1_XM_032376930.1

lysosomal transport	GO:0007041
regulation of response to external stimu...	GO:0032101
nuclear-transcribed mRNA catabolic proce...	GO:0000184
cytoplasmic translation	GO:0002181
negative regulation of stem cell differe...	GO:2000737
microtubule-based movement	GO:0007018
protein localization to cilium	GO:0061512
negative regulation of viral-induced cyt...	GO:0039532
natural killer cell degranulation	GO:0043320
response to herbicide	GO:0009635
transcription by RNA polymerase III	GO:0006383
post-Golgi vesicle-mediated transport	GO:0006892
import into cell	GO:0098657
endocytosis	GO:0006897
leukocyte activation	GO:0045321
positive regulation of NF-kappaB transcr...	GO:0051092
encapsulation of foreign target	GO:0035010
protein targeting to membrane	GO:0006612
negative regulation of hydrolase activit...	GO:0051346
cell activation involved in immune respo...	GO:0002263
determination of adult lifespan	GO:0008340
negative regulation of protein ubiquitin...	GO:0031397
protein targeting to vacuole	GO:0006623
regulation of vesicle-mediated transport	GO:0060627
ribosome biogenesis	GO:0042254
regulation of antigen receptor-mediated ...	GO:0050854
regulation of peptidyl-threonine phospho...	GO:0010799
pigment metabolic process involved in de...	GO:0043324
pigment metabolic process involved in pi...	GO:0043474
eye pigment metabolic process	GO:0042441



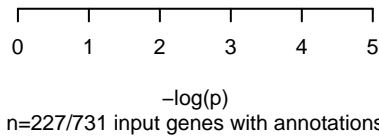
fraction genes in fg and expected value

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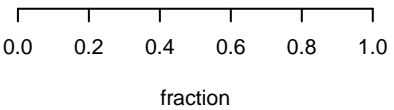
GO:MF
Fox_Nvec_vc1.1_XM_032376930.1

structural constituent of ribosome	GO:0003735
enzyme binding	GO:0019899
growth factor receptor binding	GO:0070851
peroxisome proliferator activated recept...	GO:0042975
dipeptidase activity	GO:0016805
annealing activity	GO:0140666
eukaryotic translation initiation factor...	GO:0004694
heparan-alpha-glucosaminide N-acetyltran...	GO:0015019
catalytic activity, acting on a protein	GO:0140096
ubiquitin-like protein transferase activ...	GO:0019787
ephrin receptor binding	GO:0046875
acetylglucosaminyltransferase activity	GO:0008375
ubiquitin-like protein ligase activity	GO:0061659
ubiquitin protein ligase activity	GO:0061630
ubiquitin-protein transferase activity	GO:0004842
ubiquitin protein ligase binding	GO:0031625
enzyme inhibitor activity	GO:0004857
protein tyrosine kinase binding	GO:1990782
beta-tubulin binding	GO:0048487
peptide hormone binding	GO:0017046
neurexin family protein binding	GO:0042043
glucosidase activity	GO:0015926
eukaryotic initiation factor eIF2 bindin...	GO:0071074
ubiquitin-like protein ligase binding	GO:0044389
protein kinase A binding	GO:0051018
N-acetyltransferase activity	GO:0008080
GTPase binding	GO:0051020
Hsp90 protein binding	GO:0051879
hydrolase activity, acting on glycosyl b...	GO:0016798
histone binding	GO:0042393



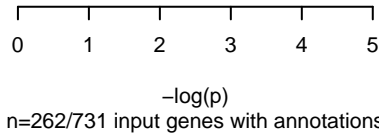
fraction genes in fg and expected value

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GO:CC
Fox_Nvec_vc1.1_XM_032376930.1

trans-Golgi network	GO:0005802
polysome	GO:0005844
cytosolic small ribosomal subunit	GO:0022627
nuclear membrane	GO:0031965
centrosome	GO:0005813
cytoplasmic vesicle	GO:0031410
cytosolic ribosome	GO:0022626
perinuclear region of cytoplasm	GO:0048471
rough endoplasmic reticulum membrane	GO:0030867
cytosolic large ribosomal subunit	GO:0022625
ribosomal subunit	GO:0044391
lytic vacuole membrane	GO:0098852
endosome	GO:0005768
polytene chromosome interband	GO:0005705
large ribosomal subunit	GO:0015934
preribosome, small subunit precursor	GO:0030688
cytoplasmic side of rough endoplasmic re...	GO:0098556
U2-type catalytic step 2 spliceosome	GO:0071007
oligosaccharyltransferase complex	GO:0008250
late endosome	GO:0005770
autolysosome	GO:0044754
microtubule cytoskeleton	GO:0015630
transcription repressor complex	GO:0017053
A band	GO:0031672
multivesicular body	GO:0005771
secretory granule	GO:0030141
early endosome	GO:0005769
lysosomal membrane	GO:0005765
cytoskeleton	GO:0005856
intracellular anatomical structure	GO:0005622



fraction genes in fg and expected value

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