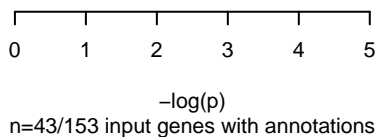
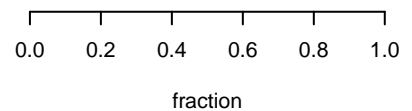


GO:BP
Fox_Nvec_vc1.1_XM_032372264.2

fraction genes in fg and expected value

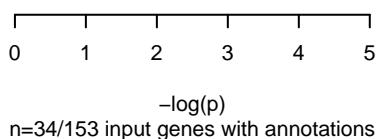
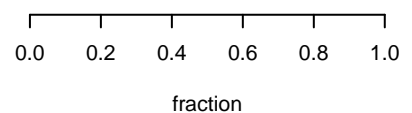
regulation of cilium beat frequency invo...	GO:0060296
phospholipid transport	GO:0015914
organic anion transport	GO:0015711
cellular response to glucagon stimulus	GO:0071377
feeding behavior	GO:0007663
glycerolipid metabolic process	GO:0046486
retina morphogenesis in camera-type eye	GO:0060042
regulation of G protein-coupled receptor...	GO:0008277
phospholipid metabolic process	GO:0006644
phospholipid biosynthetic process	GO:0008654
sensory perception of chemical stimulus	GO:0007606
phospholipid translocation	GO:0045332
lipid translocation	GO:0034204
glycerolipid biosynthetic process	GO:0045017
cell activation involved in immune respo...	GO:0002263
cellular response to cAMP	GO:0071320
regulation of membrane lipid distributio...	GO:0097035
bone mineralization involved in bone mat...	GO:0035630
cellular response to tunicamycin	GO:1904577
regulation of postsynaptic neurotransmit...	GO:0098962
positive regulation of phospholipid bios...	GO:0071073
neurofilament cytoskeleton organization	GO:0060052
negative regulation of interleukin-4 pro...	GO:0032713
collateral sprouting in absence of injur...	GO:0048669
positive regulation of N-terminal peptid...	GO:2000761
N-terminal protein amino acid modificati...	GO:0031365
regulation of N-terminal peptidyl-lysine...	GO:2000759
negative regulation of appetite by lepti...	GO:0038108
noradrenergic neuron differentiation	GO:0003357
regulation of response to calcium ion	GO:1905945

[illegible][illegible]

GO:MF
Fox_Nvec_vc1.1_XM_032372264.2

fraction genes in fg and expected value

phospholipid transporter activity	GO:0005548
mRNA binding	GO:0003729
protein domain specific binding	GO:0019904
peptidase activator activity	GO:0016504
deubiquitinase activator activity	GO:0035800
neurotransmitter receptor regulator acti...	GO:0099602
calcitonin family binding	GO:0097644
amylin binding	GO:0097645
mevalonate kinase activity	GO:0004496
phosphatidylcholine transporter activity	GO:0008525
heparan sulfate binding	GO:1904399
intracellular sodium activated potassium...	GO:0005228
acetylcholine receptor regulator activit...	GO:0030548
acetylcholine receptor activator activit...	GO:0030549
ATPase-coupled cation transmembrane tran...	GO:0019829
ATPase-coupled ion transmembrane transpo...	GO:0042625
active transmembrane transporter activit...	GO:0022804
primary active transmembrane transporter...	GO:0015399
active ion transmembrane transporter act...	GO:0022853
GDP-dissociation inhibitor activity	GO:0005092
CoA-ligase activity	GO:0016405
peroxisome proliferator activated recept...	GO:0042975
heparan sulfate proteoglycan binding	GO:0043395
PTB domain binding	GO:0051425
1-alkylglycerophosphocholine O-acetyltra...	GO:0047192
solute:bicarbonate symporter activity	GO:0140410
phosphatidylinositol transfer activity	GO:0008526
phosphatidic acid transfer activity	GO:1990050
apolipoprotein binding	GO:0034185
sodium:bicarbonate symporter activity	GO:0008510

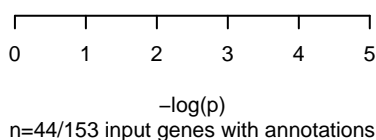
[illegible][illegible]

GO:CC
Fox_Nvec_vc1.1_XM_032372264.2

fraction genes in fg and expected value

secretory granule membrane	GO:0030667
cilium	GO:0005929
amyloid-beta complex	GO:0106003
delta DNA polymerase complex	GO:0043625
intermediate-density lipoprotein particl...	GO:0034363
high-density lipoprotein particle	GO:0034364
replisome	GO:0030894
lipoprotein particle	GO:1990777
plasma lipoprotein particle	GO:0034358
triglyceride-rich plasma lipoprotein par...	GO:0034385
endosome to plasma membrane transport ve...	GO:0070381
PR-DUB complex	GO:0035517
nuclear replisome	GO:0043601
secretory granule	GO:0030141
specific granule membrane	GO:0035579
endosome lumen	GO:0031904
ciliary rootlet	GO:0035253
platelet alpha granule membrane	GO:0031092
BBSome	GO:0034464
protein-lipid complex	GO:0032994
growth cone filopodium	GO:1990812
nuclear replication fork	GO:0043596
ciliary membrane	GO:0060170
Golgi apparatus subcompartment	GO:0098791
intrinsic component of membrane	GO:0031224
microtubule organizing center	GO:0005815
dendritic shaft	GO:0043198
zymogen granule membrane	GO:0042589
DNA polymerase complex	GO:0042575
astrocyte projection	GO:0097449

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p=6.3E-02	n=12
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p=7.1E-02	n=1
p=7.1E-02	n=1



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