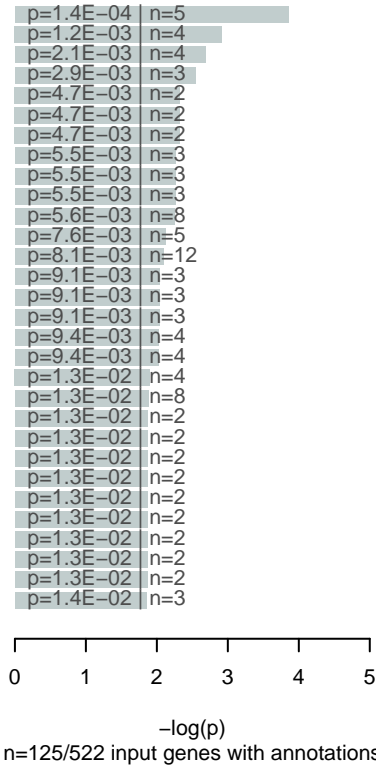


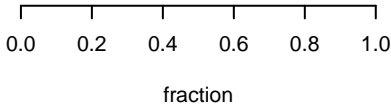
GO:BP
Fox_Nvec_vc1.1_XM_048732912.1

detection of chemical stimulus	GO:0009593
neuronal signal transduction	GO:0023041
sensory perception of taste	GO:0050909
positive regulation of cyclin-dependent ...	GO:1904031
detection of endogenous stimulus	GO:0009726
negative regulation of establishment of ...	GO:1903748
vulval location	GO:0034608
placenta blood vessel development	GO:0060674
regulation of lysosomal lumen pH	GO:0035751
eye pigment biosynthetic process	GO:0006726
regulation of Notch signaling pathway	GO:0008593
cranial nerve development	GO:0021545
Notch signaling pathway	GO:0007219
SA node cell action potential	GO:0086015
regulation of synaptic activity	GO:0060025
positive regulation of myoblast prolifer...	GO:2000288
positive regulation of potassium ion tra...	GO:0043268
positive regulation of cation channel ac...	GO:2001259
cranial nerve morphogenesis	GO:0021602
response to mechanical stimulus	GO:0009612
olfactory nerve development	GO:0021553
determination of liver left/right asymme...	GO:0071910
ocellus pigment biosynthetic process	GO:0008055
ommochrome metabolic process	GO:0046152
ocellus pigment metabolic process	GO:0046158
ommochrome biosynthetic process	GO:0006727
limb bud formation	GO:0060174
cellular metabolic compound salvage	GO:0043094
ocellus pigmentation	GO:0033060
cell differentiation in spinal cord	GO:0021515



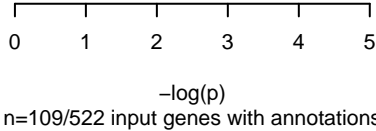
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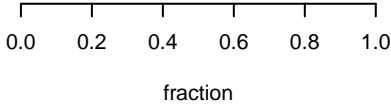
GO:MF
Fox_Nvec_vc1.1_XM_048732912.1

calcium-induced calcium release activity	GO:0048763
LIM domain binding	GO:0030274
outward rectifier potassium channel acti...	GO:0015271
adenyl-nucleotide exchange factor activi...	GO:0000774
alpha-actinin binding	GO:0051393
actinin binding	GO:0042805
aminoacyl-tRNA ligase activity	GO:0004812
ligase activity, forming carbon-oxygen b...	GO:0016875
ion channel inhibitor activity	GO:0008200
disordered domain specific binding	GO:0097718
voltage-gated sodium channel activity	GO:0005248
phospholipase A2 activity	GO:0004623
microtubule plus-end binding	GO:0051010
channel inhibitor activity	GO:0016248
ligase activity	GO:0016874
ATPase regulator activity	GO:0060590
delayed rectifier potassium channel acti...	GO:0005251
voltage-gated potassium channel activity...	GO:0086087
voltage-gated potassium channel activity...	GO:0086089
sodium channel activity	GO:0005272
intracellular cyclic nucleotide activate...	GO:0005221
voltage-gated potassium channel activity...	GO:0086008
cyclic nucleotide-gated ion channel acti...	GO:0043855
voltage-gated potassium channel activity...	GO:0086090
molecular function inhibitor activity	GO:0140678
hydrolase activity, acting on carbon-nit...	GO:0016810
potassium channel inhibitor activity	GO:0019870
ion gated channel activity	GO:0022839
kinesin binding	GO:0019894
voltage-gated calcium channel activity i...	GO:0086056



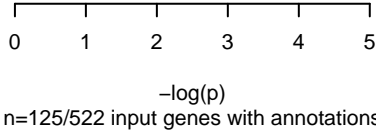
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GO:CC
Fox_Nvec_vc1.1_XM_048732912.1

GAIT complex	GO:0097452
aminoacyl-tRNA synthetase multienzyme co...	GO:0017101
integral component of cytoplasmic side o...	GO:0071458
mRNA editing complex	GO:0045293
RNA N6-methyladenosine methyltransferase...	GO:0036396
intrinsic component of endoplasmic reticu...	GO:0031227
integral component of endoplasmic reticu...	GO:0030176
plasma membrane bounded cell projection	GO:0120025
cell projection	GO:0042995
cell body	GO:0044297
luminal side of endoplasmic reticulum me...	GO:0098553
integral component of luminal side of en...	GO:0071556
dendrite	GO:0030425
dendritic tree	GO:0097447
late endosome	GO:0005770
integral component of organelle membrane	GO:0031301
integral component of Golgi membrane	GO:0030173
somatodendritic compartment	GO:0036477
intrinsic component of organelle membran...	GO:0031300
cytoplasmic side of endoplasmic reticulu...	GO:0098554
luminal side of membrane	GO:0098576
intracellular canalliculus	GO:0046691
sensory dendrite	GO:0071683
juxtaparanode region of axon	GO:0044224
intrinsic component of Golgi membrane	GO:0031228
cilium	GO:0005929
neuronal cell body	GO:0043025
cation channel complex	GO:0034703
neuron projection	GO:0043005
tertiary granule membrane	GO:0070821



fraction genes in fg and expected value

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