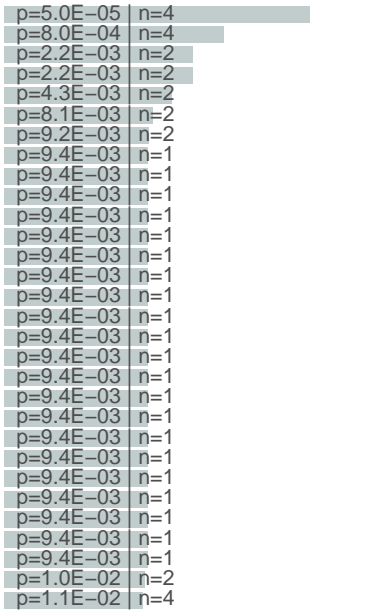


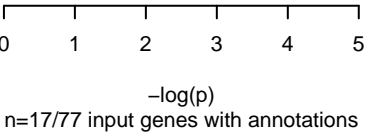
GO:BP
Fox_Nvec_vc1.1_XM_032378442.2

fraction genes in fg and expected value

potassium ion transmembrane transport	GO:0071805
positive regulation of cell cycle proces...	GO:0090068
pronephros development	GO:0048793
protein homotrimerization	GO:0070207
SRP-dependent cotranslational protein ta...	GO:0006614
embryonic placenta development	GO:0001892
receptor signaling pathway via JAK-STAT	GO:0007259
regulation of transcription involved in ...	GO:0000083
negative regulation of double-strand bre...	GO:2001033
cellular response to oscillatory fluid s...	GO:0097704
metanephric cortex development	GO:0072214
metanephric cortical collecting duct dev...	GO:0072219
cytoskeletal matrix organization at acti...	GO:0048789
positive regulation of microtubule bindi...	GO:1904528
detection of chemical stimulus involved ...	GO:0001581
cellular response to sodium arsenite	GO:1903936
metanephric smooth muscle tissue develop...	GO:0072208
detection of nodal flow	GO:0003127
positive regulation of inositol 1,4,5-tr...	GO:0031587
positive regulation of mediator complex ...	GO:2001178
cellular response to diamide	GO:0072738
positive regulation of viral entry into ...	GO:0046598
cellular response to hydrostatic pressur...	GO:0071464
embryonic process involved in female pre...	GO:0060136
positive regulation of mRNA polyadenylat...	GO:1900365
formation of cytoplasmic translation ini...	GO:0001732
viral translational termination-reinitia...	GO:0075525
positive regulation of iron ion transmem...	GO:0034761
response to pH	GO:0009268
negative regulation of cell population p...	GO:0008285



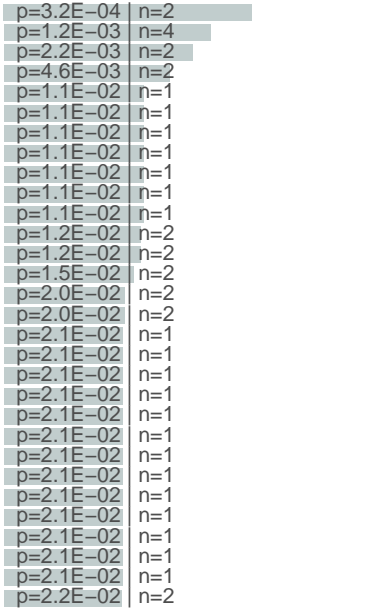
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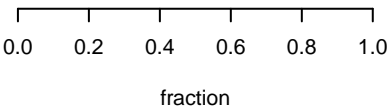
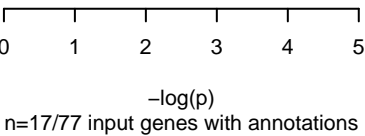
GO:MF
Fox_Nvec_vc1.1_XM_032378442.2

fraction genes in fg and expected value

calcium-activated potassium channel acti...	GO:0015269
potassium channel activity	GO:0005267
outward rectifier potassium channel acti...	GO:0015271
ribosome binding	GO:0043022
cytosine C-5 DNA demethylase activity	GO:0051747
leucine zipper domain binding	GO:0043522
HLH domain binding	GO:0043398
DNA demethylase activity	GO:0035514
translation elongation factor binding	GO:0061770
muscle alpha-actinin binding	GO:0051371
intracellular sodium activated potassium...	GO:0005228
core promoter sequence-specific DNA bind...	GO:0001046
translation factor activity, RNA binding	GO:0008135
translation regulator activity, nucleic ...	GO:0090079
DNA-binding transcription repressor acti...	GO:0001227
DNA-binding transcription repressor acti...	GO:0001217
LRR domain binding	GO:0030275
cAMP-dependent protein kinase inhibitor ...	GO:0004862
DNA-N1-methyladenine dioxygenase activit...	GO:0043734
sulfuric ester hydrolase activity	GO:0008484
cAMP-dependent protein kinase regulator ...	GO:0008603
nuclear retinoid X receptor binding	GO:0046965
calcium-induced calcium release activity	GO:0048763
N-acetylgalactosamine-4-sulfatase activi...	GO:0003943
protein kinase A catalytic subunit bindi...	GO:0034236
arylsulfatase activity	GO:0004065
sequence-specific single stranded DNA bi...	GO:0098847
STAT family protein binding	GO:0097677
beta-2 adrenergic receptor binding	GO:0031698
translation regulator activity	GO:0045182



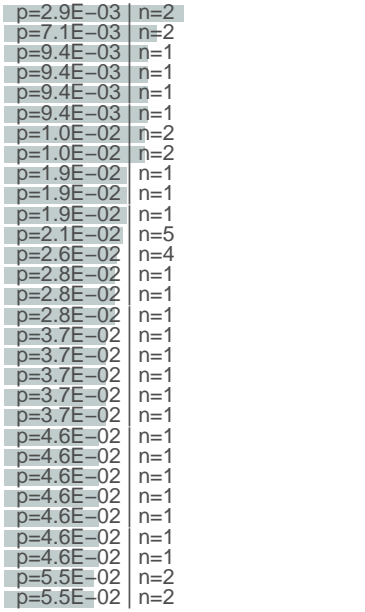
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GO:CC
Fox_Nvec_vc1.1_XM_032378442.2

fraction genes in fg and expected value

voltage-gated potassium channel complex	GO:0008076
protein kinase complex	GO:1902911
cytoskeleton of presynaptic active zone	GO:0048788
eukaryotic translation initiation factor...	GO:0071540
eukaryotic translation initiation factor...	GO:0071541
polycystin complex	GO:0002133
mitotic spindle	GO:0072686
rough endoplasmic reticulum	GO:0005791
multi-eIF complex	GO:0043614
IkkappaB kinase complex	GO:0008385
nuclear stress granule	GO:0097165
microtubule cytoskeleton	GO:0015630
ribonucleoprotein complex	GO:1990904
nucleotide-activated protein kinase comp...	GO:0031588
cAMP-dependent protein kinase complex	GO:0005952
integral component of cytoplasmic side o...	GO:0071458
basal cortex	GO:0045180
integral component of luminal side of en...	GO:0071556
germinal vesicle	GO:0042585
luminal side of endoplasmic reticulum me...	GO:0098553
cytosolic large ribosomal subunit	GO:0022625
female germ cell nucleus	GO:0001674
sensory dendrite	GO:0071683
basement membrane	GO:0005604
cytoplasmic side of endoplasmic reticul...	GO:0098554
luminal side of membrane	GO:0098576
messenger ribonucleoprotein complex	GO:1990124
mitotic spindle pole	GO:0097431
integral component of endoplasmic reticu...	GO:0030176
intrinsic component of endoplasmic retic...	GO:0031227



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