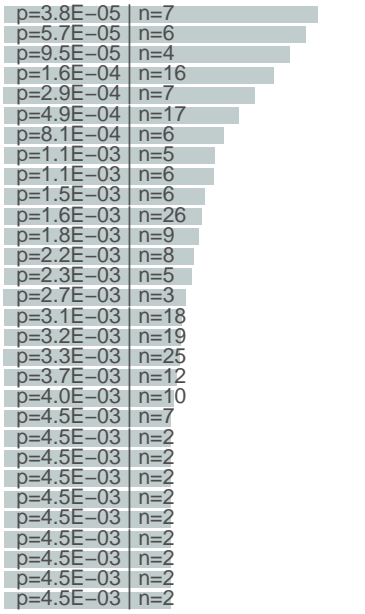


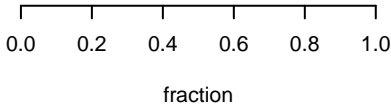
GO:BP
Fox_Nvec_vc1.1_XM_001636352.3

fraction genes in fg and expected value

integrin-mediated signaling pathway	GO:0007229
ephrin receptor signaling pathway	GO:0048013
response to muscle stretch	GO:0035994
Ras protein signal transduction	GO:0007265
actin cytoskeleton reorganization	GO:0031532
axon guidance	GO:0007411
negative regulation of synaptic assembly...	GO:0045886
sprouting angiogenesis	GO:0002040
positive regulation of JUN kinase activi...	GO:0043507
vascular endothelial growth factor recep...	GO:0048010
positive regulation of transport	GO:0051050
epidermal growth factor receptor signali...	GO:0007173
regulation of actin filament polymerizat...	GO:0030833
Fc-gamma receptor signaling pathway invo...	GO:0038096
peptide hormone processing	GO:0016486
import into cell	GO:0098657
positive regulation of cellular componen...	GO:0044089
regulation of hydrolase activity	GO:0051336
cytokine-mediated signaling pathway	GO:0019221
lymphocyte differentiation	GO:0030098
positive regulation of JNK cascade	GO:0046330
L-aspartate transmembrane transport	GO:0070778
diaphragm development	GO:0060539
cardiac conduction system development	GO:0003161
glycerol-3-phosphate metabolic process	GO:0006072
keratinization	GO:0031424
positive regulation of defense response ...	GO:0002230
pseudopodium organization	GO:0031268
regulation of cardioblast differentiatio...	GO:0051890
glycerophosphate shuttle	GO:0006127



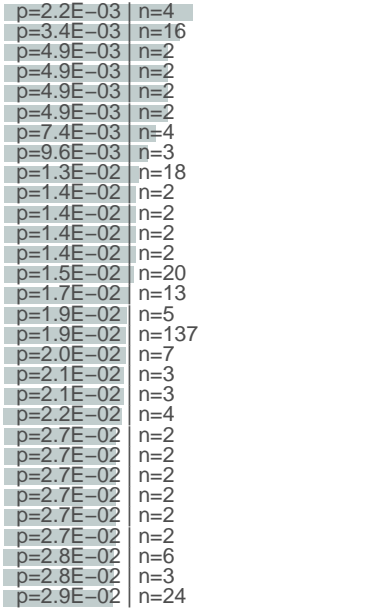
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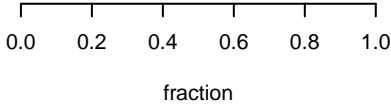
GO:MF
Fox_Nvec_vc1.1_XM_001636352.3

fraction genes in fg and expected value

signaling receptor complex adaptor activ...	GO:0030159
enzyme activator activity	GO:0008047
L-aspartate transmembrane transporter ac...	GO:0015183
glycerol-3-phosphate dehydrogenase [NAD+...	GO:0004367
glycerol-3-phosphate dehydrogenase (quin...	GO:0004368
L-glutamate transmembrane transporter ac...	GO:0005313
amide transmembrane transporter activity	GO:0042887
non-membrane spanning protein tyrosine k...	GO:0004715
protein kinase activity	GO:0004672
pyrimidine nucleotide-sugar transmembran...	GO:0015165
miRNA binding	GO:0035198
mRNA regulatory element binding translat...	GO:0000900
pre-mRNA 3'-splice site binding	GO:0030628
phosphotransferase activity, alcohol gro...	GO:0016773
protein serine/threonine kinase activity	GO:0004674
sodium ion transmembrane transporter act...	GO:0015081
binding	GO:0005488
signaling adaptor activity	GO:0035591
pre-mRNA binding	GO:0036002
peptide transmembrane transporter activi...	GO:1904680
protein tyrosine kinase binding	GO:1990782
regulatory RNA binding	GO:0061980
RNA stem-loop binding	GO:0035613
sequence-specific mRNA binding	GO:1990825
cyclase regulator activity	GO:0010851
translation repressor activity	GO:0030371
nucleotide-sugar transmembrane transport...	GO:0005338
ion channel regulator activity	GO:0099106
calmodulin-dependent protein kinase acti...	GO:0004683
transferase activity, transferring phosph...	GO:0016772



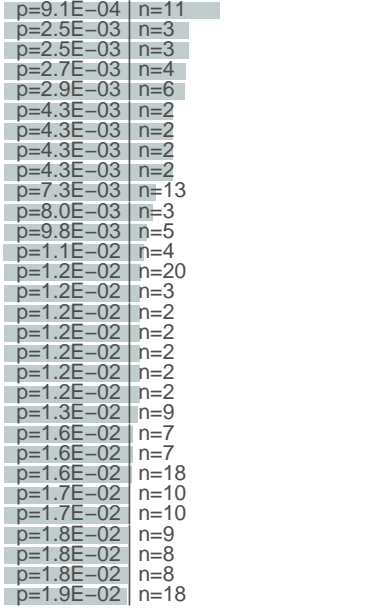
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GO:CC
Fox_Nvec_vc1.1_XM_001636352.3

fraction genes in fg and expected value

axon terminus	GO:0043679
septin cytoskeleton	GO:0032156
podosome	GO:0002102
stress fiber	GO:0001725
focal adhesion	GO:0005925
Arp2/3 protein complex	GO:0005885
U2AF complex	GO:0089701
glycerol-3-phosphate dehydrogenase compl...	GO:0009331
peripheral region of growth cone	GO:0090725
cell cortex	GO:0005938
axolemma	GO:0030673
extrinsic component of cytoplasmic side ...	GO:0031234
cleavage furrow	GO:0032154
cell body	GO:0044297
specific granule lumen	GO:0035580
septin ring	GO:0005940
sperm annulus	GO:0097227
septin complex	GO:0031105
type Ib terminal bouton	GO:0061176
type Is terminal bouton	GO:0061177
cytoplasmic ribonucleoprotein granule	GO:0036464
Golgi-associated vesicle	GO:0005798
terminal bouton	GO:0043195
neuronal cell body	GO:0043025
site of polarized growth	GO:0030427
trans-Golgi network	GO:0005802
ribonucleoprotein granule	GO:0035770
neuron spine	GO:0044309
dendritic spine	GO:0043197
dendrite	GO:0030425



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