

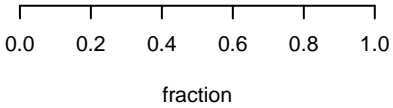
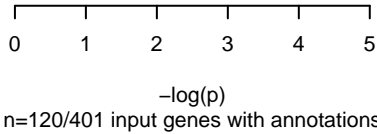
GO:BP
Ncol_Nvec_vc1.1_XM_048719424.1

fraction genes in fg and expected value

plasma membrane invagination	GO:0099024
positive regulation of multicellular org...	GO:0040018
plasma membrane raft assembly	GO:0044854
mesectoderm development	GO:0048383
cellular response to exogenous dsRNA	GO:0071360
establishment or maintenance of cytoskel...	GO:0003380
negative regulation of sequestering of t...	GO:0010891
pole cell migration	GO:0007280
positive regulation of skeletal muscle t...	GO:0048643
regulation of mitochondrial outer membra...	GO:1901028
negative regulation of transport	GO:0051051
chemical synaptic transmission	GO:0007268
regulation of mitotic cell cycle phase t...	GO:1901990
negative regulation of mitochondrion org...	GO:0010823
negative regulation of cytosolic calcium...	GO:0051481
regulation of cell communication by elec...	GO:0010649
positive regulation of interferon-gamma ...	GO:0032729
positive regulation of protein sumoylati...	GO:0033235
post-embryonic body morphogenesis	GO:0040032
microvillus assembly	GO:0030033
alpha-amino acid catabolic process	GO:1901606
axon guidance	GO:0007411
regulation of endocytosis	GO:0030100
response to light stimulus	GO:0009416
receptor-mediated endocytosis	GO:0006898
regulation of heart rate by cardiac cond...	GO:0086091
protein localization to endosome	GO:0036010
cardiac muscle cell membrane repolarizat...	GO:0099622
regulation of membrane repolarization	GO:0060306
eye photoreceptor cell differentiation	GO:0001754

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p=4.6E-03	n=12
p=5.2E-03	n=15
p=5.6E-03	n=11
p=5.6E-03	n=4
p=5.6E-03	n=4
p=5.7E-03	n=2
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p=5.7E-03	n=2
p=5.7E-03	n=2
p=5.9E-03	n=5
p=6.0E-03	n=11
p=6.3E-03	n=9
p=6.9E-03	n=11
p=7.2E-03	n=8
p=8.2E-03	n=3
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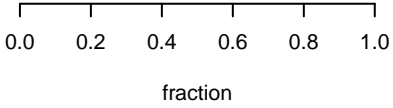
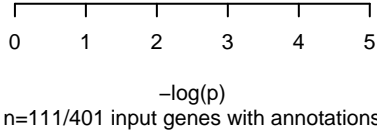
GO:MF
Ncol_Nvec_vc1.1_XM_048719424.1

fraction genes in fg and expected value

structural constituent of cytoskeleton	GO:0005200
phosphatidylcholine binding	GO:0031210
spectrin binding	GO:0030507
S100 protein binding	GO:0044548
anion binding	GO:0043168
signaling receptor binding	GO:0005102
phosphatidylethanolamine binding	GO:0008429
transcription regulator activity	GO:0140110
disordered domain specific binding	GO:0097718
ion channel regulator activity	GO:0099106
5S rRNA binding	GO:0008097
G-protein beta/gamma-subunit complex bin...	GO:0031683
ammonium ion binding	GO:0070405
potassium channel regulator activity	GO:0015459
transcription coregulator activity	GO:0003712
channel regulator activity	GO:0016247
phospholipid transporter activity	GO:0005548
calcium channel regulator activity	GO:0005246
transcription coactivator activity	GO:0003713
proteoglycan binding	GO:0043394
BH domain binding	GO:0051400
phospholipid binding	GO:0005543
ATPase-coupled ion transmembrane transpo...	GO:0042625
amino acid binding	GO:0016597
ubiquitin protein ligase binding	GO:0031625
phosphatidylinositol phosphate kinase ac...	GO:0016307
1-phosphatidylinositol-4-phosphate 5-kin...	GO:0016308
1-phosphatidylinositol-5-phosphate 4-kin...	GO:0016309
oxidative DNA demethylase activity	GO:0035516
purine nucleoside transmembrane transpor...	GO:0015211

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p=1.4E-02	n=19
p=1.6E-02	n=3
p=1.9E-02	n=5
p=2.0E-02	n=2
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p=2.0E-02	n=3
p=2.1E-02	n=11
p=2.4E-02	n=5
p=2.5E-02	n=3
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p=2.5E-02	n=8
p=2.9E-02	n=2
p=2.9E-02	n=2
p=2.9E-02	n=11
p=3.6E-02	n=3
p=3.6E-02	n=3
p=4.3E-02	n=8
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GO:CC
Ncol_Nvec_vc1.1_XM_048719424.1

fraction genes in fg and expected value

heterotrimeric G-protein complex	GO:0005834
uropod	GO:0001931
collagen and cuticulin-based cuticle ext...	GO:0060102
I band	GO:0031674
sarcolemma	GO:0042383
clathrin-coated pit	GO:0005905
cytoplasmic side of membrane	GO:0098562
asymmetric synapse	GO:0032279
recycling endosome membrane	GO:0055038
eukaryotic translation initiation factor...	GO:0071541
Z disc	GO:0030018
cytoplasmic side of plasma membrane	GO:0009898
T-tubule	GO:0030315
pericentriolar material	GO:0000242
sperm fibrous sheath	GO:0035686
sensory dendrite	GO:0071683
extrinsic component of plasma membrane	GO:0019897
postsynaptic density	GO:0014069
axon cytoplasm	GO:1904115
eukaryotic translation initiation factor...	GO:0005852
postsynaptic membrane	GO:0045211
cell projection membrane	GO:0031253
cytosolic large ribosomal subunit	GO:0022625
postsynaptic specialization	GO:0099572
postsynapse	GO:0098794
plasma membrane protein complex	GO:0098797
extrinsic component of membrane	GO:0019898
microvillus membrane	GO:0031528
membrane microdomain	GO:0098857
membrane raft	GO:0045121

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p=1.9E-03	n=2
p=1.9E-03	n=2
p=4.7E-03	n=6
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p=5.6E-03	n=3
p=8.8E-03	n=10
p=1.0E-02	n=7
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p=1.5E-02	n=9
p=1.7E-02	n=3
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p=2.0E-02	n=7
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p=2.6E-02	n=5
p=2.7E-02	n=8
p=2.8E-02	n=4
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