

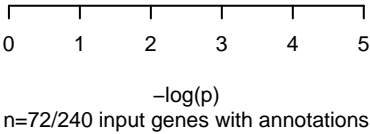
GO:BP
Ncol_Nvec_vc1.1_XM_032385089.2

fraction genes in fg and expected value

integrin-mediated signaling pathway	GO:0007229
positive regulation of JUN kinase activi...	GO:0043507
positive regulation of JNK cascade	GO:0046330
regulation of multicellular organismal p...	GO:0051239
dendritic spine organization	GO:0097061
vascular endothelial growth factor recep...	GO:0048010
synaptic vesicle transport	GO:0048489
substrate adhesion-dependent cell spread...	GO:0034446
regulation of clathrin-dependent endocyt...	GO:2000369
interleukin-2-mediated signaling pathway	GO:0038110
negative regulation of anoikis	GO:2000811
regulation of protein autophosphorylatio...	GO:0031952
negative regulation of CD4-positive, alp...	GO:0043371
growth hormone receptor signaling pathwa...	GO:0060396
signal complex assembly	GO:0007172
positive regulation of response to stimu...	GO:0048584
cell adhesion mediated by integrin	GO:0033627
ephrin receptor signaling pathway	GO:0048013
Fc-gamma receptor signaling pathway invo...	GO:0038096
synaptic vesicle cycle	GO:0099504
B cell activation	GO:0042113
positive regulation of exocytosis	GO:0045921
cell morphogenesis involved in neuron di...	GO:0048667
nuclear-transcribed mRNA catabolic proce...	GO:0000288
regulation of hematopoietic stem cell di...	GO:1902036
cell maturation	GO:0048469
epidermal growth factor receptor signali...	GO:0007173
regulation of alpha-beta T cell prolifer...	GO:0046640
pyrimidine nucleotide-sugar transmembran...	GO:0090481
peptidyl-tyrosine autophosphorylation	GO:0038083

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p=4.2E-03	n=6
p=4.2E-03	n=3
p=4.2E-03	n=3
p=4.5E-03	n=2
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p=4.5E-03	n=2
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p=4.5E-03	n=2
p=5.4E-03	n=23
p=5.9E-03	n=3
p=5.9E-03	n=3
p=5.9E-03	n=3
p=6.0E-03	n=6
p=6.5E-03	n=5
p=6.8E-03	n=4
p=7.1E-03	n=14
p=7.9E-03	n=3
p=7.9E-03	n=3
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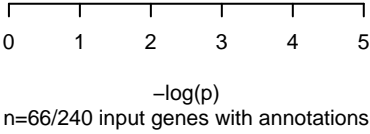
GO:MF
Ncol_Nvec_vc1.1_XM_032385089.2

fraction genes in fg and expected value

signaling adaptor activity	GO:0035591
signaling receptor complex adaptor activ...	GO:0030159
amide transmembrane transporter activity	GO:0042887
SH2 domain binding	GO:0042169
RNA cap binding	GO:0000339
protein transmembrane transporter activi...	GO:0008320
pyrimidine nucleotide-sugar transmembran...	GO:0015165
ubiquitin-like protein ligase binding	GO:0044389
non-membrane spanning protein tyrosine k...	GO:0004715
integrin binding	GO:0005178
GTPase activity	GO:0003924
JUN kinase binding	GO:0008432
BH domain binding	GO:0051400
purine ribonucleoside binding	GO:0032550
ribonucleoside binding	GO:0032549
GTP binding	GO:0005525
quaternary ammonium group binding	GO:0050997
phosphatidylcholine binding	GO:0031210
peptide transmembrane transporter activi...	GO:1904680
protein heterodimerization activity	GO:0046982
nucleoside binding	GO:0001882
purine nucleoside binding	GO:0001883
guanyl nucleotide binding	GO:0019001
guanyl ribonucleotide binding	GO:0032561
L-leucine transmembrane transporter acti...	GO:0015190
L-methionine transmembrane transporter a...	GO:0015191
glycerol-3-phosphate dehydrogenase [NAD(...	GO:0047952
1-phosphatidylinositol-3-phosphate 4-kin...	GO:0052811
Roundabout binding	GO:0048495
Toll-like receptor binding	GO:0035325

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p=4.8E-03	n=3
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p=4.9E-03	n=2
p=6.9E-03	n=8
p=9.6E-03	n=2
p=1.1E-02	n=3
p=1.5E-02	n=5
p=1.6E-02	n=2
p=1.6E-02	n=2
p=2.9E-02	n=4
p=2.9E-02	n=4
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GO:CC
Ncol_Nvec_vc1.1_XM_032385089.2

fraction genes in fg and expected value

stress fiber	GO:0001725
axon terminus	GO:0043679
cleavage furrow	GO:0032154
mRNA cap binding complex	GO:0005845
P-body	GO:0000932
sperm annulus	GO:0097227
septin complex	GO:0031105
septin ring	GO:0005940
endosome	GO:0005768
extrinsic component of plasma membrane	GO:0019897
cell leading edge	GO:0031252
cell body	GO:0044297
somatodendritic compartment	GO:0036477
extrinsic component of cytoplasmic side ...	GO:0031234
focal adhesion	GO:0005925
neuronal cell body	GO:0043025
Golgi membrane	GO:0000139
intracellular vesicle	GO:0097708
cytoplasmic vesicle	GO:0031410
dendrite	GO:0030425
dendritic tree	GO:0097447
cytosol	GO:0005829
transport vesicle membrane	GO:0030658
cell-substrate junction	GO:0030055
transport vesicle	GO:0030133
terminal bouton	GO:0043195
Sin3 complex	GO:0016580
tRNA-intron endonuclease complex	GO:0000214
ESCRT-0 complex	GO:0033565
septin filament array	GO:0032160

p=2.1E-04	n=4
p=2.8E-04	n=8
p=3.4E-04	n=4
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p=4.2E-03	n=2
p=4.2E-03	n=2
p=5.6E-03	n=13
p=7.1E-03	n=4
p=1.0E-02	n=7
p=1.3E-02	n=11
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p=2.3E-02	n=10
p=2.3E-02	n=10
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