

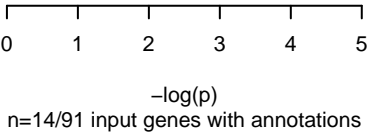
GO:BP
Ncol_Nvec_vc1.1_XM_032375635.2

fraction genes in fg and expected value

negative regulation of striated muscle c...	GO:0010664
regulation of cardiac muscle cell apopto...	GO:0010665
toxin transport	GO:1901998
cellular response to mechanical stimulus	GO:0071260
regulation of cellular response to heat	GO:1900034
positive regulation of cellular cataboli...	GO:0031331
negative regulation of mitochondrion org...	GO:0010823
cellular response to hydrogen peroxide	GO:0070301
response to peptide	GO:1901652
response to fungus	GO:0009620
sodium-independent organic anion transpo...	GO:0043252
alpha-ketoglutarate transport	GO:0015742
urate transport	GO:0015747
cellular response to water deprivation	GO:0042631
positive regulation of mitochondrial cal...	GO:0051561
renal urate salt excretion	GO:0097744
response to methotrexate	GO:0031427
fatty acid derivative transport	GO:1901571
regulation of systemic arterial blood pr...	GO:0001992
negative regulation of double-strand bre...	GO:2001033
positive regulation of microtubule bindi...	GO:1904528
cellular response to sodium arsenite	GO:1903936
negative regulation of mitochondrial mem...	GO:1902109
maternal aggressive behavior	GO:0002125
sodium-independent prostaglandin transpo...	GO:0071720
nicotinate transport	GO:2001142
negative regulation of protein targeting...	GO:1903215
negative regulation of female receptivit...	GO:0007621
cellular response to diamide	GO:0072738
short-chain fatty acid transport	GO:0015912

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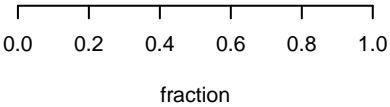
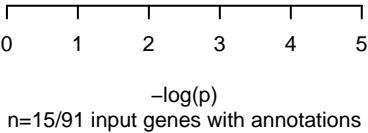
GO:MF
Ncol_Nvec_vc1.1_XM_032375635.2

fraction genes in fg and expected value

protein kinase C binding	GO:0005080
prostaglandin receptor activity	GO:0004955
urate transmembrane transporter activity	GO:0015143
D-xylose 1-dehydrogenase (NADP+) activit...	GO:0047837
nicotinate transmembrane transporter act...	GO:0090416
V1A vasopressin receptor binding	GO:0031894
short-chain fatty acid transmembrane tra...	GO:0015636
sodium-independent organic anion transme...	GO:0015347
translation elongation factor binding	GO:0061770
chloride ion binding	GO:0031404
NAD(P)+ transhydrogenase activity	GO:0008746
vasopressin receptor activity	GO:0005000
toxin transmembrane transporter activity	GO:0019534
quaternary ammonium group transmembrane ...	GO:0015651
STAT family protein binding	GO:0097677
galactosylgalactosylxylosylprotein 3-bet...	GO:0015018
sequence-specific single stranded DNA bi...	GO:0098847
dipeptidase activity	GO:0016805
peptide hormone binding	GO:0017046
AMP binding	GO:0016208
adenyl-nucleotide exchange factor activi...	GO:0000774
RNA cap binding	GO:0000339
carboxypeptidase activity	GO:0004180
organic cation transmembrane transporter...	GO:0015101
RNA polymerase II intronic transcription...	GO:0001162
anion:anion antiporter activity	GO:0015301
phosphatase activator activity	GO:0019211
inorganic anion exchanger activity	GO:0005452
ammonium transmembrane transporter activ...	GO:0008519
solute:anion antiporter activity	GO:0140323

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GO:CC
Ncol_Nvec_vc1.1_XM_032375635.2

fraction genes in fg and expected value

Lsm1-7-Pat1 complex	GO:1990726
mRNA cap binding complex	GO:0005845
nuclear stress granule	GO:0097165
intrinsic component of mitochondrial out...	GO:0031306
integral component of mitochondrial oute...	GO:0031307
kinetochore microtubule	GO:0005828
RNA cap binding complex	GO:0034518
supramolecular complex	GO:0099080
ESCRT III complex	GO:0000815
messenger ribonucleoprotein complex	GO:1990124
mitotic spindle pole	GO:0097431
pronucleus	GO:0045120
PML body	GO:0016605
ribonucleoprotein granule	GO:0035770
integral component of mitochondrial memb...	GO:0032592
Golgi lumen	GO:0005796
euchromatin	GO:0000791
intrinsic component of mitochondrial mem...	GO:0098573
spindle	GO:0005819
ESCRT complex	GO:0036452
spindle microtubule	GO:0005876
integral component of membrane	GO:0016021
brush border membrane	GO:0031526
integral component of plasma membrane	GO:0005887
intrinsic component of membrane	GO:0031224
intrinsic component of plasma membrane	GO:0031226
kinetochore	GO:0000776
cis-Golgi network	GO:0005801
caveola	GO:0005901
mitotic spindle	GO:0072686

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