

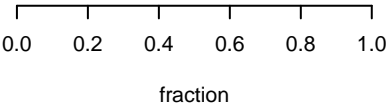
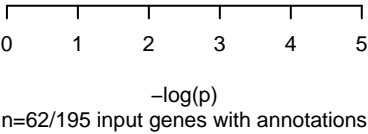
GO:BP
Ncol_Nvec_vc1.1_XM_032365977.2

fraction genes in fg and expected value

negative regulation of peptidyl-serine p... GO:0033137
behavioral response to ethanol GO:0048149
positive regulation of cholesterol biosy... GO:0045542
sevenless signaling pathway GO:0045500
regulation of extracellular exosome asse... GO:1903551
ephrin receptor signaling pathway GO:0048013
mitotic cell cycle checkpoint signaling GO:0007093
insulin receptor signaling pathway GO:0008286
nuclear-transcribed mRNA catabolic proce... GO:0000288
Ras protein signal transduction GO:0007265
positive regulation of exosomal secretio... GO:1903543
regulation of receptor signaling pathway... GO:0046425
mitotic DNA damage checkpoint signaling GO:0044773
regulation of receptor signaling pathway... GO:1904892
negative regulation of cell division GO:0051782
RNA phosphodiester bond hydrolysis, exon... GO:0090503
phosphatidylinositol dephosphorylation GO:0046856
short-term memory GO:0007614
mitotic DNA integrity checkpoint signali... GO:0044774
receptor signaling pathway via STAT GO:0097696
RNA phosphodiester bond hydrolysis GO:0090501
receptor signaling pathway via JAK-STAT GO:0007259
regulation of release of sequestered cal... GO:0010880
release of sequestered calcium ion into ... GO:0014808
mitotic G1/S transition checkpoint signa... GO:0044819
mitotic G1 DNA damage checkpoint signali... GO:0031571
torso signaling pathway GO:0008293
release of sequestered calcium ion into ... GO:1903514
sarcoplasmic reticulum calcium ion trans... GO:0070296
positive regulation of hormone biosynthe... GO:0046886

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p=3.8E-03	n=3
p=4.0E-03	n=4
p=5.1E-03	n=5
p=5.2E-03	n=3
p=5.2E-03	n=7
p=6.6E-03	n=2
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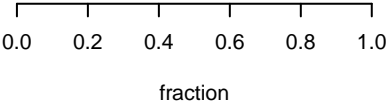
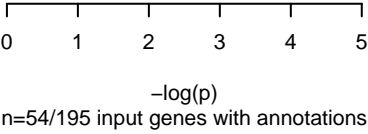
GO:MF
Ncol_Nvec_vc1.1_XM_032365977.2

fraction genes in fg and expected value

phospholipid binding GO:0005543
phosphatidylinositol phosphate 4-phospha... GO:0034596
acetyltransferase activity GO:0016407
ribonuclease activity GO:0004540
exoribonuclease activity, producing 5'-p... GO:0016896
MAP kinase activity GO:0004707
3'-5'-exoribonuclease activity GO:0000175
exoribonuclease activity GO:0004532
transmembrane transporter binding GO:0044325
phosphatidylinositol binding GO:0035091
exonuclease activity, active with either... GO:0016796
N-acetyltransferase activity GO:0008080
3'-5' exonuclease activity GO:0008408
N-acyltransferase activity GO:0016410
ATPase binding GO:0051117
primary active transmembrane transporter... GO:0015399
protein heterodimerization activity GO:0046982
glycogen binding GO:2001069
phosphatidylinositol-4,5-bisphosphate 4-... GO:0034597
S100 protein binding GO:0044548
[phosphorylase] phosphatase activity GO:0050196
open rectifier potassium channel activit... GO:0005252
platelet-derived growth factor receptor ... GO:0005161
heparan sulfate sulfotransferase activit... GO:0034483
protein O-GlcNAc transferase activity GO:0097363
poly(A)-specific ribonuclease activity GO:0004535
heparan sulfate 2-O-sulfotransferase act... GO:0004394
sulfotransferase activity GO:0008146
ABC-type sterol transporter activity GO:0034041
7SK snRNA binding GO:0097322

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p=1.8E-02	n=5
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p=2.8E-02	n=2
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p=2.8E-02	n=3
p=3.1E-02	n=4
p=3.3E-02	n=7
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GO:CC
Ncol_Nvec_vc1.1_XM_032365977.2

fraction genes in fg and expected value

fusome GO:0045169
vacuolar proton-transporting V-type ATPa... GO:0016471
intrinsic component of plasma membrane GO:0031226
proton-transporting V-type ATPase comple... GO:0033176
proton-transporting two-sector ATPase co... GO:0016469
telomerase holoenzyme complex GO:0005697
interleukin-5 receptor complex GO:0005895
dendritic microtubule GO:1901588
ESCRT-0 complex GO:0033565
CCR4-NOT complex GO:0030014
CCR4-NOT core complex GO:0030015
sarcoplasmic reticulum lumen GO:0033018
exocytic vesicle GO:0070382
ESCRT complex GO:0036452
secretory vesicle GO:0099503
intracellular vesicle GO:0097708
cytoplasmic vesicle GO:0031410
acetylcholine-gated channel complex GO:0005892
clathrin complex GO:0071439
signal peptidase complex GO:0005787
anchored component of plasma membrane GO:0046658
vesicle GO:0031982
side of membrane GO:0098552
synaptic vesicle GO:0008021
plasma membrane signaling receptor compl... GO:0098802
endosome GO:0005768
integral component of plasma membrane GO:0005887
muscle cell projection GO:0036194
muscle cell projection membrane GO:0036195
lamellar body GO:0042599

p=2.0E-03	n=3
p=2.2E-02	n=2
p=2.6E-02	n=10
p=2.9E-02	n=2
p=2.9E-02	n=2
p=3.5E-02	n=1
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p=3.5E-02	n=1
p=3.5E-02	n=1
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p=4.6E-02	n=9
p=4.8E-02	n=17
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p=6.8E-02	n=1
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p=7.4E-02	n=17
p=7.6E-02	n=5
p=8.3E-02	n=4
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