

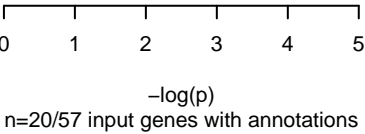
GO:BP  
gc014

fraction genes in fg and expected value

positive regulation of neuroblast prolifer...	GO:0002052
adenylate cyclase-activating G protein-c...	GO:0007189
visual perception	GO:0007601
G protein-coupled receptor signaling pat...	GO:0007186
cGMP-mediated signaling	GO:0019934
pyramidal neuron migration to cerebral c...	GO:0021852
neurohypophysis development	GO:0021985
regulation of cyclase activity	GO:0031279
insulin receptor signaling pathway via p...	GO:0038028
cysteine biosynthetic process from serin...	GO:0006535
axon midline choice point recognition	GO:0016199
hypothalamus cell differentiation	GO:0021979
regulation of heart rate	GO:0002027
protein transmembrane transport	GO:0071806
regulation of lyase activity	GO:0051339
vasodilation	GO:0042311
cysteine biosynthetic process via cystat...	GO:0019343
homoiothermy	GO:0042309
neuron fate determination	GO:0048664
oxygen transport	GO:0015671
response to folic acid	GO:0051593
hydrogen sulfide biosynthetic process	GO:0070814
positive regulation of transforming grow...	GO:0032914
transsulfuration	GO:0019346
regulation of fever generation	GO:0031620
homocysteine catabolic process	GO:0043418
negative regulation of neuron maturation	GO:0014043
response to endothelin	GO:1990839
homoserine metabolic process	GO:0009092
midgut development	GO:0007494

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p=8.0E-04	n=3
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p=5.1E-03	n=2
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p=5.8E-03	n=2
p=6.6E-03	n=2
p=7.9E-03	n=1
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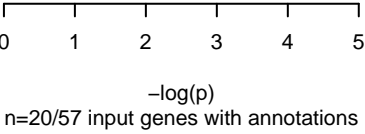
GO:MF  
gc014

fraction genes in fg and expected value

G protein-coupled receptor activity	GO:0004930
oxygen binding	GO:0019825
extracellular matrix structural constitu...	GO:0030021
cystathionine beta-synthase activity	GO:0004122
carbon monoxide binding	GO:0070025
melanocyte-stimulating hormone receptor ...	GO:0004980
nitric oxide binding	GO:0070026
oxygen carrier activity	GO:0005344
endothelin receptor activity	GO:0004962
oxidoreductase activity, acting on other...	GO:0016662
nitrite reductase (NO-forming) activity	GO:0050421
peptide hormone binding	GO:0017046
nitrite reductase activity	GO:0098809
hormone binding	GO:0042562
oxidoreductase activity, acting on other...	GO:0016661
peptide binding	GO:0042277
type 1 angiotensin receptor binding	GO:0031702
metallopeptidase activity	GO:0008237
vitamin binding	GO:0019842
amide binding	GO:0033218
heme binding	GO:0020037
metallocarboxypeptidase activity	GO:0004181
angiotensin receptor binding	GO:0031701
tetrapyrrole binding	GO:0046906
S-adenosyl-L-methionine binding	GO:1904047
calcium-dependent phospholipid binding	GO:0005544
extracellular matrix protein binding	GO:1990430
neuropeptide binding	GO:0042923
G protein-coupled receptor activity invo...	GO:0099530
G protein-coupled neurotransmitter recep...	GO:0150047

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p=7.9E-03	n=1
p=1.1E-02	n=1
p=1.3E-02	n=1
p=1.3E-02	n=1
p=1.3E-02	n=2
p=1.6E-02	n=1
p=1.7E-02	n=2
p=1.8E-02	n=1
p=1.9E-02	n=3
p=2.1E-02	n=1
p=2.5E-02	n=2
p=3.2E-02	n=2
p=3.4E-02	n=3
p=3.6E-02	n=2
p=3.6E-02	n=1
p=3.9E-02	n=1
p=4.1E-02	n=2
p=4.2E-02	n=1
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GO:CC  
gc014

fraction genes in fg and expected value

sperm head plasma membrane	GO:1990913
fibrillar collagen trimer	GO:0005583
leading edge membrane	GO:0031256
external side of plasma membrane	GO:0009897
Schaffer collateral - CA1 synapse	GO:0098685
cell projection membrane	GO:0031253
cell surface	GO:0009986
G protein-coupled receptor complex	GO:0097648
G protein-coupled receptor dimeric compl...	GO:0038037
G protein-coupled receptor heterodimeric...	GO:0038039
GABA receptor complex	GO:1902710
G protein-coupled GABA receptor complex	GO:1902712
ruffle membrane	GO:0032587
cell periphery	GO:0071944
plasma membrane	GO:0005886
axolemma	GO:0030673
trans-Golgi network membrane	GO:0032588
neuron projection membrane	GO:0032589
main axon	GO:0044304
ruffle	GO:0001726
side of membrane	GO:0098552
cell projection	GO:0042995
plasma membrane signaling receptor compl...	GO:0098802
membrane raft	GO:0045121
membrane microdomain	GO:0098857
dendritic shaft	GO:0043198
perikaryon	GO:0043204
centriolar satellite	GO:0034451
intrinsic component of plasma membrane	GO:0031226
integral component of plasma membrane	GO:0005887

p=1.3E-03	n=2
p=2.6E-03	n=1
p=9.9E-03	n=2
p=2.5E-02	n=2
p=3.2E-02	n=2
p=4.5E-02	n=2
p=5.3E-02	n=3
p=5.4E-02	n=1
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p=5.4E-02	n=1
p=5.4E-02	n=1
p=5.6E-02	n=1
p=7.0E-02	n=12
p=7.0E-02	n=11
p=7.1E-02	n=1
p=8.6E-02	n=1
p=8.8E-02	n=1
p=9.8E-02	n=1
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p=1.3E-01	n=2
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