

group II metabotropic glutamate receptor... GO:0001641
scaffold protein binding GO:0097110
collagen binding GO:0005518
beta-catenin binding GO:0008013
G protein-coupled receptor activity GO:0004930
protein-hormone receptor activity GO:0016500
phosphatidylinositol binding GO:0035091
calcium channel regulator activity GO:005246
ligand-gated sodium channel activity GO:0015280
carboxylic ester hydrolase activity GO:0052689
inositol tetrakisphosphate 5-kinase acti... GO:0047326
succinyl-CoA hydrolase activity GO:00052815
glutaryl-CoA hydrolase activity GO:00052815
glutaryl-CoA hydrolase activity GO:00447466
flavonoid binding GO:0097243
hydroxymethylglutaryl-CoA hydrolase activity GO:00447994
acetoacetyl-CoA hydrolase activity GO:0047603
inositol-1,4,5-trisphosphate 6-kinase ac... GO:000823
inositol tetrakisphosphate 3-kinase aci... GO:000823
unositol tetrakisphosphate 3-kinase acti... GO:000824
UDP-galactose:beta-N-acetylglucosamine b... GO:0008499
myo-inositol-1,2,3,4,6-heptakisphosphate... GO:0102732
choloyl-CoA hydrolase activity GO:0033882
acetylcholine binding GO:0042166
vitamin binding GO:0042166
vitamin binding GO:0042166
cholinesterase activity GO:0003990
sphinganine kinase activity GO:0003990
sphinganine kinase activity GO:0003986
long-chain-fatty-acyl-CoA reductase acti... GO:00050062 p=1.9E-04 | n=3 p=5.2E-04 | n=4 fg=0.04 fg=0.06 bg=0.00 bg=0.01 p=5.2E-04 n=4 p=6.8E-04 n=4 p=1.4E-03 n=3 p=1.7E-03 n=14 p=1.9E-03 n=4 p=3.4E-03 n=5 p=3.4E-03 n=3 fğ=0.06 bg=0.01 fg=0.04 bg=0.00 fg=0.21 fg=0.06 bg=0.06 bg=0.01 fg=0.07 fg=0.04 bg=0.01 bq = 0.00p=3.4E-03 n=2 p=8.3E-03 n=4 p=9.0E-03 n=1 fg=0.03 fg=0.06 bg=0.00 bg=0.01 bg=0.00 fg=0.01 p=9.0F-03 n=1fa=0.01 bg=0.00 bg=0.00 fg=0.01 fg=0.01 p=9.0E-03 n=1 $p=9.0E-03 \mid n=1$ bg = 0.00bg=0.00 p=9.0E-03 n=1 fg=0.01 fg=0.01 n=9.0E-03 | n=1 bg=0.00 p=9.0E-03 n=1 bg=0.00 p=9.0E-03 n=1 fq = 0.01bg=0.00 p=9.0E-03 n=1 fg=0.01 bg=0.00 p=9.0E-03 n=1 p=9.0E-03 n=1 fg=0.01 bg=0.00 bg=0.00 fg=0.01 p=9.0E-03 n=1 p=1.5E-02 n=2 fg=0.01 bg=0.00 fg=0.03 bg=0.00 fğ=0.06 p=1.5E-02 p=1.6E-02 bg=0.01 n=4fg=0.03 bg=0.00 n=2p=1.6E-02 p=1.6E-02 n=2 fg=0.03 bg=0.00 fg=0.03 bg=0.00 bg=0.00 n=2p=1.8E-02 p=1.8E-02 fg=0.01 fg=0.01 n=1 A1 adenosine receptor binding GO:0031686 long-chain-fatty-acyl-CoA reductase acti... GO:0050062 bq = 0.00n=1∥ fğ=0.01 p=1.8E-02 bg=0.00 2 3 0.0 5 0.2 0.4 0.6 0.8 1.0 fraction -log(p)

n=68/113 input genes with annotations

GO:CC fraction genes in fg and expected value

