

1-acylglycerophosphocholine O-acyltransf... GO:0047184
GABA-gated chloride ion channel activity GO:0022851
GABA-A receptor activity GO:0004890
benzodiazepine receptor activity GO:0008503
methyl-CpG binding GO:0008327
TFIID-class transcription factor complex... GO:0001094
transcription coactivator binding GO:0001223
1-alkylglycerophosphocholine O-acetyltra... GO:0047192
carbon-nitrogen ligase activity, with gl... GO:0016884
protein-arginine omega-N monomethyltrans... GO:0035241
histone methyltransferase activity (H4-R... GO:0044020
protease binding GO:0002020
1-acylglycerol-3-phosphate O-acyltransfe... GO:0003841
mitogen-activated protein kinase binding GO:00051019
MAP kinase activity GO:0004707
ligand-activated transcription factor ac... GO:0098531
nuclear receptor activity GO:0004879
acylglycerol O-acyltransferase activity GO:0004879
acylglycerol O-acyltransferase activity GO:0004879
acylglycerol O-acyltransferase activity GO:0004033
2-acylglycerol-3-phosphate O-acyltransfe... GO:0047144
ion gated channel activity GO:00042239
lysophosphatidic acid acyltransferase activity... GO:0071617
structural constituent of ribosome GO:0003735
cysteine-type endopeptidase activity GO:0004197
TBP-class protein binding GO:0017025
DNA-binding transcription activator acti... GO:0001216
extracellular ligand-gated ion channel a... GO:0005230
RNA polymerase core enzyme binding GO:0043175 p=7.9E-04 | n=3 p=2.9E-03 | n=3 fg=0.01 fg=0.01 bg=0.00 bg=0.00 p=2.9E-03 n=3 p=2.9E-03 n=3 fg=0.01 bg=0.00 bg=0.00 fg=0.01 p=2.9E-03 n=3 p=6.9E-03 n=3 p=6.9E-03 n=3 fg=0.01 bg=0.00 fg=0.01 bg=0.00 fg=0.01 fg=0.01 bg=0.00 p=8.6E-03 | n=2 bq=0.00 p=8.6E-03 n=2 p=8.6E-03 n=2 fg=0.01 fg=0.01 bg=0.00 bq=0.00 bg=0.00 p=8.6E-03 fg=0.01 p=1.3F-02fa=0.03 bg=0.01 bg=0.00 l n=6 p=1.3E-02 fg=0.01 p=2.0E-02n=4fa = 0.02bg=0.01 bğ=0.00 fg=0.01 fg=0.01 p=2.1E-02 p=2.1E-02 n=3bg=0.00 n=3bg=0.00 p=2.1E-02 p=2.4E-02 fg = 0.01bg=0.00 n=3bg=0.00 n=2p=2.4E-02 p=3.0E-02 fg=0.01 n=2bg=0.00 n=10 fg=0.05 bg=0.02 p=3.1E-02 p=3.1E-02 fg=0.01 bg=0.00 n=3n=3fg=0.01 bg=0.00 p=3.4E-02 fg=0.05 n=10 bg=0.02 p=3.4E-02 fg=0.02 bg=0.01 n=4 fğ=0.02 p=3.4E-02 n=4bg=0.01 p=3.7E-02 fg=0.05 bg=0.03 bg=0.03 n=12 p=3.7E-02 p=4.2E-02 fg=0.05 n=12 n=6 bq = 0.01∥ fğ=0.01 p=4.4E-02 bg=0.00 2 5 0.0 0.2 3 0.4 0.6 8.0 1.0 fraction –log(p) n=220/802 input genes with annotations

GO:CC Ncol_Nvec_vc1.1_XM_001625588.3 fraction genes in fg and expected value

