

lipid phosphatase activity GO:0042577
translation elongation factor activity GO:0003746
oxidoreductase activity, acting on a sul... GO:0016670
ubiquitin binding GO:0043130
phospholipase activity GO:0004620
phosphatidic acid binding GO:0070300
carbohydrate kinase activity GO:0019200 p=4.6E-03 | n=2 p=4.6E-03 | n=2 fg=0.02 fg=0.02 bg=0.00 bg=0.00 p=1.3E-02 n=2 p=1.9E-02 n=3 fg=0.02 bg=0.00 bg=0.01 fg=0.03 fğ=0.04 p=1.9E-02 p=2.5E-02 n=4 bg=0.01 fg=0.02 bg=0.00 n=2p=2.5E-02 p=3.4E-02 fg=0.02 fg=0.03 n=2 bg=0.00 ubiquitin–like protein binding GO:0032182
lipase activity GO:0016298
JUN kinase binding GO:0008432
G–protein beta/gamma–subunit complex bin... GO:0031683
transcription coregulator activity GO:0003712
transcription corepressor activity GO:0003714 n=3bg=0.01 p=3.5E-02 p=4.0E-02 n=4fg=0.04 bg=0.01 fg=0.02 bg=0.00 n=2p=4.0E-02 p=5.1E-02 bg=0.00 fg=0.02 bg=0.06 bg=0.02 fg=0.10 n = 11p=5.4E-02 p=5.7E-02 fg=0.05 transcription corepressor activity GO:0003714
HMG box domain binding GO:0071837
poly-pyrimidine tract binding GO:0008187
chromatin binding GO:0008187
4-hydroxy-2-oxoglutarate aldolase activi... GO:0008700
sphingomyelin phosphodiesterase D activi... GO:0050290
1-alkenylglycerophosphocholine O-acyltra... GO:0047159
testosterone 17-beta-dehydrogenase (NADP... GO:0047045
cupric ion binding GO:1903135
P-Man:Man3GlcNAc2-PP-Dol alpha-1,2-man... GO:0004377
manganese ion transmembrane transporter ... GO:0005384
5'-3' DNA helicase activity GO:0043139 fa=0.02 n=2bg = 0.00fg=0.02 p=5.7E-02 bg=0.00 p=6.5E-02n = 10fg=0.09 bg=0.05 p=6.8E-02 p=6.8E-02 fg=0.01 bg=0.00 n=1n=1fg=0.01 bg=0.00 p=6.8E-02 p=6.8E-02 fg=0.01 bg=0.00 n=1fg=0.01 p=6.8E n=1bg=0.00 p=6.8E-02 fg=0.01 bg=0.00 n=1p=6.8E-02 p=6.8E-02 n=1fg=0.01 bg=0.00 fg=0.01 bg=0.00 n=15'-3' DNA helicase activity GO:0043139
nickel cation transmembrane transporter ... GO:0015099
8-oxo-7,8-dihydroguanine DNA N-glycosyla... GO:0034039
calcium-independent phospholipase A2 act... GO:0047499
steroid dehydrogenase activity GO:0016229 p=6.8E-02 fg=0.01 bg=0.00 n=1p=6.8E-02 fg=0.01 bg=0.00 n=1fg=0.01 p=6.8E-02 n=1bg=0.00 p=6.8E-02 p=6.8E-02 p=6.8E-02 bg=0.00 bg=0.00 fg=0.01 n=1fg=0.01 fg=0.01 n=1 cyclic nucleotide-dependent protein kina... GO:0004690 6-phosphofructo-2-kinase activity GO:0003873 n=1bq = 0.00fg=0.01 p=6.8E-02 bg=0.00 0 2 0.0 1 3 5 0.2 0.4 0.6 8.0 1.0 fraction –log(p) n=108/384 input genes with annotations

GO:CC Elav_Nvec_vc1.1_XM_048728399.1

fraction genes in fg and expected valu

