

p=3.1E-03 | n=2 p=6.0E-03 | n=2 fg=0.03 fg=0.03 bg=0.00 bg=0.00 cAMP response element binding GO:0035497 iron ion transmembrane transporter activit... GO:0005381
histone acetyltransferase activity GO:0004002
core promoter sequence—specific DNA bind... GO:0001046
acetyltransferase activity GO:0016407
peptide—lysine—N—acetyltransferase activit... GO:00041733
polynucleotide adenylyltransferase activit... GO:0004652 p=1.4E-02 n=2 p=1.9E-02 n=3 ■fg=0.03 bg=0.00 bg=0.01 fg=0.04 fğ=0.04 n=3p=1.9E-02 bg=0.01 p=2.0E-02 fg=0.03 bg=0.00 n=2fg=0.03 fg=0.03 n=2 p=2.0E-02 bg=0.00 peptide N-acetyltransferase activity GO:004652
peptide N-acetyltransferase activity GO:0034212
glycogen (starch) synthase activity GO:0004373
histone pre-mRNA stem-loop binding GO:0071207
malate dehydrogenase (NADP+) activity GO:0046554
netrin receptor activity GO:0005042
adiponectin binding GO:0055100
uanyl nucleotide exchange factor inhibi GO:1990624 p=2.0E-02n=2bg=0.00 p=3.3E-02 p=3.3E-02 fg=0.01 fg=0.01 n=1bg=0.00 bq=0.00 n=1bg=0.00 p=3.3E-02 fg=0.01 p=3.3E-02fa = 0.01bg=0.00 bg=0.00 n=1p=3.3E fg=0.01 adiponectin binding GO:0055100
guanyl nucleotide exchange factor inhibi... GO:1990624
polynucleotide 5'-phosphatase activity GO:0004651
voltage-gated proton channel activity GO:0030171
nickel cation transmembrane transporter ... GO:0015099
open rectifier potassium channel activit... GO:0005252
polynucleotide phosphatase activity GO:0098518
beta-galactosidase activity GO:0004565
gamma-glutamylaminecyclotransferase acti... GO:0061929
glycogen synthase activity, transferring... GO:0061547
DNA end binding GO:0045027
cobalt ion transmembrane transporter act... GO:0015087 p=3.3E-02 $f_{\alpha} = 0.01$ n=1bg = 0.00p=3.3E-02 bg=0.00 fg=0.01 fg=0.01 p=3.3E-02n=1bg=0.00 p=3.3E bg=0.00 n=1 $f\bar{q} = 0.01$ p=3.3E-02n=1bg=0.00 p=3.3E-02 fg=0.01 bg=0.00 n=1fg=0.01 p=3.3E-02n=1bg=0.00 p=3.3E-02 fg=0.01 bg=0.00 n=1p=3.3E-02 p=3.3E-02 n=1fg=0.01 bg=0.00 fg=0.01 bg=0.00 n=1cobalt ion transmembrane transporter act... GO:0015087
DP-N-acetylglucosamine-dolichyl-phospha... GO:0003975
RNA polymerase I general transcription i... GO:0001181
D-xylose 1-dehydrogenase (NADP+) activit... GO:0047837
N-acylneuraminate-9-phosphatase activity GO:0050124 p=3.3E-02 n=1fg=0.01 bg=0.00 p=3.3E-02 fg=0.01 bg=0.00 UDP n=1fg=0.01 p=3.3E-02 n=1bg=0.00 bg=0.00 bg=0.00 fg=0.01 p=3.3E-02n=1p=3.3E-02 p=3.3E-02 fg=0.01 fg=0.01 n=1 arylformamidase activity GO:0004061 adipokinetic hormone receptor activity GO:0097003 n=1bq = 0.00p=3.3E-02 fg=0.01 bg=0.00 2 0.0 1 3 5 0.2 0.4 0.6 0.8 1.0 fraction –log(p)

GO:CC Elav_Nvec_vc1.1_XM_032375670.2

n=77/274 input genes with annotations

fraction genes in fg and expected valu

