

Metallophos_C	p=6.8E-02	n=3
AMP-binding	p=1.1E-01	n=2
Bap31	p=1.1E-01	n=1
Bap31_Bap29_C	p=1.1E-01	n=1
BAT2_N	p=1.1E-01	n=1
BCAS3	p=1.1E-01	n=1
Biopterin_H	p=1.1E-01	n=1
BRAP2	p=1.1E-01	n=1
CLN3	p=1.1E-01	n=2
ClpS	p=1.1E-01	n=1
Complex1_49kDa	p=1.1E-01	n=1
Cript	p=1.1E-01	n=1
DAG_kinase_N	p=1.1E-01	n=1
DDE_Tnp_1_7	p=1.1E-01	n=2
DER1	p=1.1E-01	n=1
DNA_pol_delta_4	p=1.1E-01	n=1
DUF11	p=1.1E-01	n=1
DUF1151	p=1.1E-01	n=1
DUF1387	p=1.1E-01	n=1
DUF2046	p=1.1E-01	n=1
DUF2370	p=1.1E-01	n=1
DUF4470	p=1.1E-01	n=1
DUF4471	p=1.1E-01	n=1
DUF747	p=1.1E-01	n=1
FERM_M	p=1.1E-01	n=3
FERM_N	p=1.1E-01	n=3
Glyco_transf_41	p=1.1E-01	n=1
Glycos_transf_4	p=1.1E-01	n=1
GpcrRhopsn4	p=1.1E-01	n=2
HIF-1a_CTAD	p=1.1E-01	n=1

fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.01	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.01	bg=0.00
fg=0.00	bg=0.00



-log10(p)

n=228/236 input genes with annotations



fraction