

Ncol_Nvec_vc1.1_XM_048722339.1

fraction genes in fg and bg

Adaptin_N	p=9.5E-02	n=2
Alpha_adaptin_C	p=9.5E-02	n=1
AP3B1_C	p=9.5E-02	n=1
CBF_beta	p=9.5E-02	n=1
CCDC158	p=9.5E-02	n=1
CEBP_ZZ	p=9.5E-02	n=1
CID	p=9.5E-02	n=2
Coatomer_WDAD	p=9.5E-02	n=1
Cobalamin_bind	p=9.5E-02	n=1
CRAL_TRIO	p=9.5E-02	n=2
Creatinase_N	p=9.5E-02	n=1
Creatinase_N_2	p=9.5E-02	n=1
CREPT	p=9.5E-02	n=1
CTF_NFI	p=9.5E-02	n=1
DBB	p=9.5E-02	n=1
DIRP	p=9.5E-02	n=1
DMAP1	p=9.5E-02	n=1
Dopey_N	p=9.5E-02	n=1
DOT1	p=9.5E-02	n=1
DUF1387	p=9.5E-02	n=1
DUF3543	p=9.5E-02	n=1
DUF3583	p=9.5E-02	n=1
DUF3689	p=9.5E-02	n=1
DUF4430	p=9.5E-02	n=1
DUF4555	p=9.5E-02	n=1
DUF4598	p=9.5E-02	n=1
DUF4611	p=9.5E-02	n=1
EF-hand_9	p=9.5E-02	n=1
eIF3g	p=9.5E-02	n=1
Glycos_transf_2	p=9.5E-02	n=2

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

 $-\log_{10}(p)$

n=300/274 input genes with annotations



fraction