

Ncol_Nvec_vc1.1_XM_032383717.2

fraction genes in fg and bg

2Fe-2S_thioredx	p=8.1E-02	n=1
AA_kinase	p=8.1E-02	n=1
AA_permease_2	p=8.1E-02	n=2
Alg14	p=8.1E-02	n=1
Asparaginase_2	p=8.1E-02	n=2
AZUL	p=8.1E-02	n=1
BACK	p=8.1E-02	n=4
BTB	p=8.1E-02	n=4
bZIP_1	p=8.1E-02	n=3
bZIP_2	p=8.1E-02	n=2
C1_2	p=8.1E-02	n=1
CAP	p=8.1E-02	n=2
CBF_beta	p=8.1E-02	n=1
CSN5_C	p=8.1E-02	n=1
DHHA1	p=8.1E-02	n=1
Dopey_N	p=8.1E-02	n=1
DUF1211	p=8.1E-02	n=1
DUF2052	p=8.1E-02	n=1
DUF2340	p=8.1E-02	n=1
DUF3543	p=8.1E-02	n=1
DUF4706	p=8.1E-02	n=1
DUF727	p=8.1E-02	n=1
eIF2_C	p=8.1E-02	n=1
Fes1	p=8.1E-02	n=1
Fic	p=8.1E-02	n=1
Glyco_transf_22	p=8.1E-02	n=1
Glyco_transf_41	p=8.1E-02	n=1
Glycohydro_20b2	p=8.1E-02	n=1
Heme_oxygenase	p=8.1E-02	n=1
HSP70	p=8.1E-02	n=2

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=0.01	bg=0.00
fg=0.00	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.00	bg=0.00
fg=0.01	bg=0.00
fg=0.00	bg=0.00
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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



-log10(p)

n=192/196 input genes with annotations



fraction