

Elav_Nvec_vc1.1_XM_048722445.1

fraction genes in fg and bg

Acetyltransf_16	p=6.5E-02	n=1
ARA70	p=6.5E-02	n=1
Asp_Arg_Hydrox	p=6.5E-02	n=1
BAR_3_WASP_bdg	p=6.5E-02	n=1
BCL_N	p=6.5E-02	n=1
Cation_ATPase	p=6.5E-02	n=2
CSN5_C	p=6.5E-02	n=1
DEP	p=6.5E-02	n=2
DNA_gyraseB	p=6.5E-02	n=1
DNA_topoisolv	p=6.5E-02	n=1
DUF2181	p=6.5E-02	n=1
DUF4460	p=6.5E-02	n=1
DUF4461	p=6.5E-02	n=1
DUF4795	p=6.5E-02	n=1
EIF4E-T	p=6.5E-02	n=1
IML1	p=6.5E-02	n=1
JAB	p=6.5E-02	n=2
Med13_C	p=6.5E-02	n=1
MID_MedPIWI	p=6.5E-02	n=1
MM_CoA_mutase	p=6.5E-02	n=1
MRG	p=6.5E-02	n=1
MRP_L53	p=6.5E-02	n=1
MYCBPAP	p=6.5E-02	n=1
Myosin-VI_CBD	p=6.5E-02	n=1
Na_Ca_ex_C	p=6.5E-02	n=1
NAD_kinase	p=6.5E-02	n=1
NERD	p=6.5E-02	n=1
NOPS	p=6.5E-02	n=1
OSR1_C	p=6.5E-02	n=2
P-mevalo_kinase	p=6.5E-02	n=1

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



$-\log_{10}(p)$
n=214/179 input genes with annotations



fraction