

Fox_Nvec_vc1.1_XM_032374843.2

fraction genes in fg and bg

MAM	p=8.6E-02	n=3
Astacin	p=8.6E-02	n=2
Casc1_N	p=8.6E-02	n=1
CID	p=8.6E-02	n=2
CREPT	p=8.6E-02	n=1
CTC1	p=8.6E-02	n=1
DUF2236	p=8.6E-02	n=1
DUF3543	p=8.6E-02	n=1
DUF3583	p=8.6E-02	n=1
DUF4419	p=8.6E-02	n=1
DUF92	p=8.6E-02	n=1
EF-hand_9	p=8.6E-02	n=1
F-box_4	p=8.6E-02	n=1
Hamartin	p=8.6E-02	n=1
HLH	p=8.6E-02	n=3
KASH_CCD	p=8.6E-02	n=1
malic	p=8.6E-02	n=1
Malic_M	p=8.6E-02	n=1
MamL-1	p=8.6E-02	n=1
MBD_C	p=8.6E-02	n=1
MBDa	p=8.6E-02	n=1
mit_SMPDase	p=8.6E-02	n=1
MITF_TFEB_C_3_N	p=8.6E-02	n=1
MRV11	p=8.6E-02	n=1
MYCBPAP	p=8.6E-02	n=1
Ndr	p=8.6E-02	n=1
p53-inducible11	p=8.6E-02	n=1
PAW	p=8.6E-02	n=1
Pecanex_C	p=8.6E-02	n=1
Peptidase_M48_N	p=8.6E-02	n=1

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
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=0.00	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=0.00	bg=0.00



$-\log_{10}(p)$
n=178/167 input genes with annotations



fraction