

Fox_Nvec_vc1.1_XM_032366211.2

fraction genes in fg and bg

ABC_ATPase	p=9.5E-02	n=1
Acatn	p=9.5E-02	n=1
ADK	p=9.5E-02	n=2
Apyrase	p=9.5E-02	n=1
BAT2_N	p=9.5E-02	n=1
C1_2	p=9.5E-02	n=1
C1_4	p=9.5E-02	n=1
C12orf66_like	p=9.5E-02	n=1
Cast	p=9.5E-02	n=1
CCDC50_N	p=9.5E-02	n=1
CD225	p=9.5E-02	n=2
CLN3	p=9.5E-02	n=2
Cohesin_load	p=9.5E-02	n=1
Dak1	p=9.5E-02	n=1
Dak2	p=9.5E-02	n=1
DNA_pol3_delta2	p=9.5E-02	n=1
DUF1180	p=9.5E-02	n=1
DUF2370	p=9.5E-02	n=1
DUF3598	p=9.5E-02	n=1
DUF3677	p=9.5E-02	n=1
DUF3827	p=9.5E-02	n=1
DUF4483	p=9.5E-02	n=1
DUF4490	p=9.5E-02	n=1
Ebp2	p=9.5E-02	n=1
EF-hand_5	p=9.5E-02	n=3
EHN	p=9.5E-02	n=1
F-box-like	p=9.5E-02	n=4
Filament	p=9.5E-02	n=1
GAIN	p=9.5E-02	n=2
Glyco_hydro_56	p=9.5E-02	n=1

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

 $-\log_{10}(p)$

n=290/255 input genes with annotations



fraction