

## Fox\_Nvec\_vc1.1\_XM\_001635172.3

## fraction genes in fg and bg

Adaptin_N	p=8.9E-02	n=2
Alpha_adaptin_C	p=8.9E-02	n=1
AP3B1_C	p=8.9E-02	n=1
CBF_beta	p=8.9E-02	n=1
CCDC158	p=8.9E-02	n=1
CID	p=8.9E-02	n=2
Coatomer_WDAD	p=8.9E-02	n=1
CREPT	p=8.9E-02	n=1
CTF_NFI	p=8.9E-02	n=1
DBB	p=8.9E-02	n=1
DIRP	p=8.9E-02	n=1
DMAP1	p=8.9E-02	n=1
Dopey_N	p=8.9E-02	n=1
DOT1	p=8.9E-02	n=1
DUF11	p=8.9E-02	n=1
DUF1387	p=8.9E-02	n=1
DUF3543	p=8.9E-02	n=1
DUF4598	p=8.9E-02	n=1
DUF4611	p=8.9E-02	n=1
EF-hand_9	p=8.9E-02	n=1
eIF3g	p=8.9E-02	n=1
F-box	p=8.9E-02	n=2
Glycos_transf_2	p=8.9E-02	n=2
GOLGA2L5	p=8.9E-02	n=1
HAUS6_N	p=8.9E-02	n=1
HEAT	p=8.9E-02	n=2
HPS3_C	p=8.9E-02	n=1
HPS3_Mid	p=8.9E-02	n=1
HPS3_N	p=8.9E-02	n=1
HTH_23	p=8.9E-02	n=1

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=0.00	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
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=0.01	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

 $-\log_{10}(p)$ 

n=281/246 input genes with annotations



fraction