

## Fox\_Nvec\_vc1.1\_XM\_001631403.3

## fraction genes in fg and bg

Atg8	p=4.7E-02	n=2
DUF2370	p=5.7E-02	n=1
DUF719	p=5.7E-02	n=1
GAF_2	p=5.7E-02	n=1
Gly_rich	p=5.7E-02	n=1
Glycos_transf_4	p=5.7E-02	n=1
KIAA1430	p=5.7E-02	n=1
NHR2	p=5.7E-02	n=1
p25-alpha	p=5.7E-02	n=1
Peptidase_M16_M	p=5.7E-02	n=1
RBB1NT	p=5.7E-02	n=1
Ribonuc_2-5A	p=5.7E-02	n=1
SPRY	p=5.7E-02	n=2
zf-MYND	p=5.7E-02	n=2
Annexin	p=5.9E-02	n=1
ANTH	p=5.9E-02	n=1
EF-hand_13	p=5.9E-02	n=1
ERp29	p=5.9E-02	n=1
I_LWEQ	p=5.9E-02	n=1
Intu_longin_1	p=5.9E-02	n=1
Ras	p=5.9E-02	n=3
Ribosomal_S9	p=5.9E-02	n=1
TAFH	p=5.9E-02	n=1
tRNA_anti-codon	p=5.9E-02	n=1
tRNA-synt_2	p=5.9E-02	n=1
U-box	p=5.9E-02	n=1
V-set	p=5.9E-02	n=1
CBM_21	p=7.0E-02	n=1
EF-hand_1	p=7.0E-02	n=1
HSP90	p=7.0E-02	n=1

fg=0.02	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	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=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.03	bg=0.01
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00



-log10(p)

n=80/70 input genes with annotations



fraction