

Ncol\_Nvec\_vc1.1\_XM\_032375635.2

fraction genes in fg and bg

Actin	p=8.7E-02	n=2
AlaDh_PNT_C	p=8.7E-02	n=1
APOBEC_N	p=8.7E-02	n=1
BAR_3_WASP_bdg	p=8.7E-02	n=1
BCL_N	p=8.7E-02	n=1
BNIP3	p=8.7E-02	n=1
BTB_2	p=8.7E-02	n=3
Ca_hom_mod	p=8.7E-02	n=1
CARD_2	p=8.7E-02	n=1
Cellulase	p=8.7E-02	n=1
COX4	p=8.7E-02	n=1
CSRNP_N	p=8.7E-02	n=1
DNA_pol_phi	p=8.7E-02	n=1
DSPn	p=8.7E-02	n=1
DUF2052	p=8.7E-02	n=1
DUF4464	p=8.7E-02	n=1
Enkurin	p=8.7E-02	n=1
FAD_binding_2	p=8.7E-02	n=1
FAM221	p=8.7E-02	n=1
Fe-S_biosyn	p=8.7E-02	n=1
Flavokinase	p=8.7E-02	n=1
Forkhead	p=8.7E-02	n=3
FTHFS	p=8.7E-02	n=1
Fz	p=8.7E-02	n=2
GCV_H	p=8.7E-02	n=1
Gly_rich	p=8.7E-02	n=1
Glyco_transf_43	p=8.7E-02	n=1
Homeodomain	p=8.7E-02	n=3
HSF_DNA-bind	p=8.7E-02	n=1
I-set	p=8.7E-02	n=3

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
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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=0.01	bg=0.00



-log10(p)

n=199/179 input genes with annotations



fraction