

## Ncol\_Nvec\_vc1.1\_XM\_032366211.2

## fraction genes in fg and bg

SET	p=3.1E-02	n=5
ABC_ATPase	p=8.7E-02	n=1
Acatn	p=8.7E-02	n=1
ADK	p=8.7E-02	n=2
Apyrase	p=8.7E-02	n=1
BAT2_N	p=8.7E-02	n=1
C1_2	p=8.7E-02	n=1
C1_4	p=8.7E-02	n=1
C12orf66_like	p=8.7E-02	n=1
Cast	p=8.7E-02	n=1
Cohesin_load	p=8.7E-02	n=1
CortBP2	p=8.7E-02	n=1
Dak1	p=8.7E-02	n=1
Dak2	p=8.7E-02	n=1
DNA_pol3_delta2	p=8.7E-02	n=1
DUF1180	p=8.7E-02	n=1
DUF2370	p=8.7E-02	n=1
DUF3598	p=8.7E-02	n=1
DUF3677	p=8.7E-02	n=1
DUF3827	p=8.7E-02	n=1
DUF4483	p=8.7E-02	n=1
DUF4490	p=8.7E-02	n=1
DUF4611	p=8.7E-02	n=1
Ebp2	p=8.7E-02	n=1
EF-hand_5	p=8.7E-02	n=3
EHN	p=8.7E-02	n=1
Filament	p=8.7E-02	n=1
Glyco_hydro_56	p=8.7E-02	n=1
GPS2_interact	p=8.7E-02	n=1
HHH	p=8.7E-02	n=1

fg=0.02	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=0.00	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



-log10(p)

n=280/236 input genes with annotations



fraction