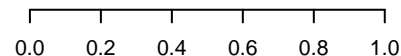


Elav\_Nvec\_vc1.1\_XM\_032364784.2

fraction genes in fg and bg

60KD_IMP	p=9.2E-02	n=1
adh_short	p=9.2E-02	n=3
AP3D1	p=9.2E-02	n=1
BAT2_N	p=9.2E-02	n=1
CybS	p=9.2E-02	n=1
Dimer_Tnp_hAT	p=9.2E-02	n=2
DSHCT	p=9.2E-02	n=1
DUF2451	p=9.2E-02	n=1
DUF3689	p=9.2E-02	n=1
DUF4371	p=9.2E-02	n=2
DUF5745	p=9.2E-02	n=1
E3_UbLigase_EDD	p=9.2E-02	n=1
FANCI_HD1	p=9.2E-02	n=1
FANCI_HD2	p=9.2E-02	n=1
FANCI_S1	p=9.2E-02	n=1
FANCI_S1-cap	p=9.2E-02	n=1
FANCI_S2	p=9.2E-02	n=1
FANCI_S3	p=9.2E-02	n=1
FANCI_S4	p=9.2E-02	n=1
GPI2	p=9.2E-02	n=1
GRAM	p=9.2E-02	n=2
HGTP_anticonodon2	p=9.2E-02	n=1
His_Phos_2	p=9.2E-02	n=2
HLH	p=9.2E-02	n=3
JHY	p=9.2E-02	n=1
MBOAT	p=9.2E-02	n=1
MCM	p=9.2E-02	n=1
MCM_lid	p=9.2E-02	n=1
MCM_OB	p=9.2E-02	n=1
Melibiase_2	p=9.2E-02	n=1

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



$-\log_{10}(p)$   
n=212/170 input genes with annotations

fraction