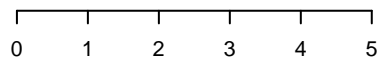


Elav\_Nvec\_vc1.1\_XM\_032374864.2

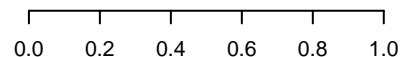
fraction genes in fg and bg

ABC_ATPase	p=8.9E-02	n=1
Acatn	p=8.9E-02	n=1
ADK	p=8.9E-02	n=2
Apyrase	p=8.9E-02	n=1
ArfGap	p=8.9E-02	n=2
ASF1_hist_chap	p=8.9E-02	n=1
BAT2_N	p=8.9E-02	n=1
C1_2	p=8.9E-02	n=1
C1_4	p=8.9E-02	n=1
C12orf66_like	p=8.9E-02	n=1
Cast	p=8.9E-02	n=1
CLN3	p=8.9E-02	n=2
Cohesin_load	p=8.9E-02	n=1
Dak1	p=8.9E-02	n=1
Dak2	p=8.9E-02	n=1
DNA_pol3_delta2	p=8.9E-02	n=1
DUF1180	p=8.9E-02	n=1
DUF2370	p=8.9E-02	n=1
DUF3598	p=8.9E-02	n=1
DUF3827	p=8.9E-02	n=1
DUF4483	p=8.9E-02	n=1
DUF4611	p=8.9E-02	n=1
Ebp2	p=8.9E-02	n=1
EHN	p=8.9E-02	n=1
Elf1	p=8.9E-02	n=1
F-box-like	p=8.9E-02	n=4
Filament	p=8.9E-02	n=1
GAIN	p=8.9E-02	n=2
Glyco_hydro_56	p=8.9E-02	n=1
Glyoxalase_4	p=8.9E-02	n=1

fg=0.00	bg=0.00
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fg=0.01	bg=0.00
fg=0.00	bg=0.00
fg=0.01	bg=0.00
fg=0.00	bg=0.00
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fg=0.01	bg=0.00
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$-\log_{10}(p)$   
n=279/234 input genes with annotations



fraction