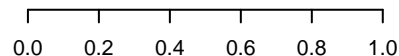


Elav\_Nvec\_vc1.1\_XM\_032367288.2

fraction genes in fg and bg

Anticodon_1	p=8.3E-02	n=1
Arm_2	p=8.3E-02	n=1
Calsequestrin	p=8.3E-02	n=1
Cbl_N	p=8.3E-02	n=1
Cbl_N2	p=8.3E-02	n=1
Cbl_N3	p=8.3E-02	n=1
CDC73_C	p=8.3E-02	n=1
CDC73_N	p=8.3E-02	n=1
CDV3	p=8.3E-02	n=1
COQ7	p=8.3E-02	n=1
COX4	p=8.3E-02	n=1
DIM1	p=8.3E-02	n=1
DUF2340	p=8.3E-02	n=1
DUF3399	p=8.3E-02	n=1
DUF4485	p=8.3E-02	n=1
eIF-3c_N	p=8.3E-02	n=1
F-box	p=8.3E-02	n=2
FERM_f0	p=8.3E-02	n=1
Glyco_transf_8	p=8.3E-02	n=2
GMP_synt_C	p=8.3E-02	n=1
GPS2_interact	p=8.3E-02	n=1
HR1	p=8.3E-02	n=1
LTV	p=8.3E-02	n=1
MCM3AP_GANP	p=8.3E-02	n=1
Med1	p=8.3E-02	n=1
MKLP1_Arf_bdg	p=8.3E-02	n=1
mRNA_cap_C	p=8.3E-02	n=1
mRNA_cap_enzyme	p=8.3E-02	n=1
Nramp	p=8.3E-02	n=1
Nup153	p=8.3E-02	n=1

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



$-\log_{10}(p)$   
n=242/195 input genes with annotations

fraction