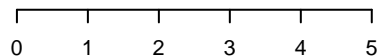


Elav\_Nvec\_vc1.1\_XM\_048734482.1

fraction genes in fg and bg

A_deaminase	p=1.7E-01	n=1
Adap_comp_sub	p=1.7E-01	n=2
Adaptin_binding	p=1.7E-01	n=1
Adaptin_N	p=1.7E-01	n=2
Amino_oxidase	p=1.7E-01	n=2
AP4E_app_platf	p=1.7E-01	n=1
ASD2	p=1.7E-01	n=1
AWS	p=1.7E-01	n=1
B2-adapt-app_C	p=1.7E-01	n=1
Bax1-I	p=1.7E-01	n=2
Bromodomain	p=1.7E-01	n=4
BTB_2	p=1.7E-01	n=5
BTK	p=1.7E-01	n=1
C5HCH	p=1.7E-01	n=1
CBF_beta	p=1.7E-01	n=1
CCDC50_N	p=1.7E-01	n=1
Clat_adaptor_s	p=1.7E-01	n=2
CLU	p=1.7E-01	n=1
CLU_N	p=1.7E-01	n=1
COX5A	p=1.7E-01	n=1
CUB	p=1.7E-01	n=4
CybS	p=1.7E-01	n=1
DDT	p=1.7E-01	n=2
DEP	p=1.7E-01	n=2
Dis3l2_C_term	p=1.7E-01	n=1
Dishevelled	p=1.7E-01	n=1
DOR	p=1.7E-01	n=1
Dsh_C	p=1.7E-01	n=1
DUF1280	p=1.7E-01	n=1
DUF3399	p=1.7E-01	n=1

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=0.01	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=372/358 input genes with annotations



fraction