

Elav\_Nvec\_vc1.1\_XM\_032374352.2

fraction genes in fg and bg

ADP_ribosyl_GH	p=8.9E-02	n=1
CDK5_activator	p=8.9E-02	n=1
CEBP_ZZ	p=8.9E-02	n=1
CS	p=8.9E-02	n=1
DUF4745	p=8.9E-02	n=1
FERM_M	p=8.9E-02	n=2
Glyco_hydro_2	p=8.9E-02	n=1
Glyco_hydro_2_C	p=8.9E-02	n=1
Glyco_hydro_2_N	p=8.9E-02	n=1
HSL_N	p=8.9E-02	n=1
Integrin_b_cyt	p=8.9E-02	n=1
Integrin_B_tail	p=8.9E-02	n=1
Integrin_beta	p=8.9E-02	n=1
MyTH4	p=8.9E-02	n=1
PSI_integrin	p=8.9E-02	n=1
PurA	p=8.9E-02	n=1
RRM_7	p=8.9E-02	n=1
Runt	p=8.9E-02	n=1
Sprouty	p=8.9E-02	n=1
Transthyretin	p=8.9E-02	n=1
VGCC_alpha2	p=8.9E-02	n=1
VWA_N	p=8.9E-02	n=1
zf-UBP	p=8.9E-02	n=1
ABC_membrane	p=1.3E-01	n=1
Abhydrolase_3	p=1.3E-01	n=1
Cpn60_TCP1	p=1.3E-01	n=1
DM	p=1.3E-01	n=1
DMA	p=1.3E-01	n=1
DUF3504	p=1.3E-01	n=1
HLH	p=1.3E-01	n=2

fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
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fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.03	bg=0.00

-log<sub>10</sub>(p)

n=70/56 input genes with annotations



fraction