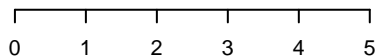


Elav\_Nvec\_vc1.1\_XM\_001640726.3

fraction genes in fg and bg

HLH	p=1.4E-02	n=13
PHD	p=1.4E-02	n=12
FERM_C	p=1.0E-01	n=7
Frizzled	p=1.0E-01	n=4
zf-C2HC	p=1.0E-01	n=4
FERM_N	p=2.1E-01	n=8
AC_N	p=2.5E-01	n=3
CBM_21	p=2.5E-01	n=3
DDT	p=2.5E-01	n=3
DIX	p=2.5E-01	n=3
Mito_carr	p=2.5E-01	n=10
RNB	p=2.5E-01	n=3
WSD	p=2.5E-01	n=3
FERM_M	p=2.5E-01	n=9
Pkinase	p=2.7E-01	n=28
CH	p=2.9E-01	n=8
6PGD	p=3.4E-01	n=1
7tm_1	p=3.4E-01	n=54
7tm_3	p=3.4E-01	n=5
A_deaminase	p=3.4E-01	n=1
AAA_30	p=3.4E-01	n=1
Actin	p=3.4E-01	n=3
Activin_recpt	p=3.4E-01	n=2
Adaptin_binding	p=3.4E-01	n=1
AF-4	p=3.4E-01	n=1
AF-4_C	p=3.4E-01	n=1
AP4E_app_platf	p=3.4E-01	n=1
APC_rep	p=3.4E-01	n=1
APH	p=3.4E-01	n=1
ARA70	p=3.4E-01	n=1



$-\log_{10}(p)$   
n=791/1005 input genes with annotations

fg=0.01	bg=0.00
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=0.01	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
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=0.01	bg=0.00
fg=0.02	bg=0.01
fg=0.01	bg=0.00
fg=0.00	bg=0.00
fg=0.04	bg=0.03
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



fraction