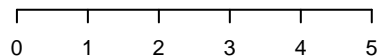


Elav_Nvec_vc1.1_XM_032378928.2

fraction genes in fg and bg

CBX7_C	p=5.2E-02	n=2
Chromo	p=5.2E-02	n=2
3Beta_HSD	p=1.0E-01	n=1
AC_N	p=1.0E-01	n=2
CDC50	p=1.0E-01	n=1
CDK5_activator	p=1.0E-01	n=1
CLASP_N	p=1.0E-01	n=1
CSRNP_N	p=1.0E-01	n=1
DOR	p=1.0E-01	n=1
DUF1154	p=1.0E-01	n=1
DUF3543	p=1.0E-01	n=1
DUF4495	p=1.0E-01	n=1
DUF758	p=1.0E-01	n=1
HSL_N	p=1.0E-01	n=1
KIND	p=1.0E-01	n=1
Mab-21	p=1.0E-01	n=1
MyTH4	p=1.0E-01	n=1
P66_CC	p=1.0E-01	n=1
PH_14	p=1.0E-01	n=1
PHD	p=1.0E-01	n=2
RasGEF	p=1.0E-01	n=1
RasGEF_N	p=1.0E-01	n=1
SAND	p=1.0E-01	n=1
Synapsin	p=1.0E-01	n=1
Synapsin_C	p=1.0E-01	n=1
Tau95	p=1.0E-01	n=1
Tau95_N	p=1.0E-01	n=1
WAC_Acf1_DNA_bd	p=1.0E-01	n=1
zf-HC5HC2H_2	p=1.0E-01	n=1
LRR_8	p=1.1E-01	n=2



-log₁₀(p)
n=80/66 input genes with annotations

fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.01	bg=0.00
fg=0.02	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=0.01	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=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.02	bg=0.00



fraction