

Elav_Nvec_vc1.1_XM_032385556.2

fraction genes in fg and bg

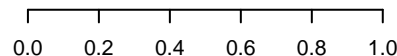
APCDDC	p=8.6E-02	n=1
BAR_3	p=8.6E-02	n=1
DUF229	p=8.6E-02	n=1
FERM_f0	p=8.6E-02	n=1
I_LWEQ	p=8.6E-02	n=1
IRS	p=8.6E-02	n=1
LIM	p=8.6E-02	n=2
PALP	p=8.6E-02	n=1
PDZ_6	p=8.6E-02	n=1
PH	p=8.6E-02	n=3
SCAPER_N	p=8.6E-02	n=1
Talin_middle	p=8.6E-02	n=1
UNC-79	p=8.6E-02	n=1
VBS	p=8.6E-02	n=1
CD20	p=1.2E-01	n=1
COesterase	p=1.2E-01	n=1
Patched	p=1.2E-01	n=1
PH_19	p=1.2E-01	n=1
WD40_2	p=1.2E-01	n=1
zf-met	p=1.2E-01	n=1
7tm_1	p=1.4E-01	n=8
Guanylate_kin	p=1.4E-01	n=1
Peptidase_C1	p=1.4E-01	n=1
Pou	p=1.4E-01	n=1
SH3_9	p=1.4E-01	n=1
ASC	p=1.5E-01	n=1
Homeodomain	p=1.5E-01	n=2
P_proprotein	p=1.5E-01	n=1
Peptidase_S8	p=1.5E-01	n=1
RhoGAP	p=1.5E-01	n=1

fg=0.02	bg=0.00
fg=0.02	bg=0.00
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fg=0.12	bg=0.07
fg=0.02	bg=0.00
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fg=0.02	bg=0.00
fg=0.03	bg=0.01
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00



$-\log_{10}(p)$

n=50/53 input genes with annotations



fraction