

Elav_Nvec_vc1.1_XM_032377792.2

fraction genes in fg and bg

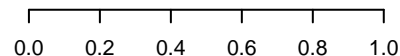
FA	p=6.9E-02	n=3
AAA_30	p=1.1E-01	n=1
Aminotran_5	p=1.1E-01	n=2
Arv1	p=1.1E-01	n=1
BCL_N	p=1.1E-01	n=1
BET	p=1.1E-01	n=1
BRD4_CDT	p=1.1E-01	n=1
BRO1	p=1.1E-01	n=2
BTD	p=1.1E-01	n=1
CBF_beta	p=1.1E-01	n=1
CCDC154	p=1.1E-01	n=1
CCDC158	p=1.1E-01	n=1
Cys_Met_Meta_PP	p=1.1E-01	n=1
DOR	p=1.1E-01	n=1
DOT1	p=1.1E-01	n=1
DUF1180	p=1.1E-01	n=1
DUF21	p=1.1E-01	n=1
EIF4E-T	p=1.1E-01	n=1
FERM_C	p=1.1E-01	n=3
FERM_M	p=1.1E-01	n=4
FERM_N	p=1.1E-01	n=3
GAF_2	p=1.1E-01	n=1
GFA	p=1.1E-01	n=1
Glyco_transf_90	p=1.1E-01	n=1
GOLGA2L5	p=1.1E-01	n=1
HA2	p=1.1E-01	n=2
HEAT	p=1.1E-01	n=2
Hrs_helial	p=1.1E-01	n=1
Hydrolase	p=1.1E-01	n=2
IQ_SEC7_PH	p=1.1E-01	n=1

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



$-\log_{10}(p)$

n=237/224 input genes with annotations



fraction