

Elav_Nvec_vc1.1_XM_032385556.2

fraction genes in fg and bg

Acetyltransf_16	p=1.2E-01	n=1
Arm_vescicular	p=1.2E-01	n=1
CaM_bdg_C0	p=1.2E-01	n=1
CCDC142	p=1.2E-01	n=1
CDV3	p=1.2E-01	n=1
CEBP_ZZ	p=1.2E-01	n=1
COLFI	p=1.2E-01	n=1
CortBP2	p=1.2E-01	n=1
Cys_Met_Meta_PP	p=1.2E-01	n=1
DNA_ligase_A_C	p=1.2E-01	n=1
DNA_ligase_A_M	p=1.2E-01	n=1
DNA_ligase_A_N	p=1.2E-01	n=1
DUF4659	p=1.2E-01	n=1
DUF4686	p=1.2E-01	n=1
DUF758	p=1.2E-01	n=1
DUF773	p=1.2E-01	n=1
DUF974	p=1.2E-01	n=1
EAF	p=1.2E-01	n=1
ELM2	p=1.2E-01	n=2
ETS_PEA3_N	p=1.2E-01	n=1
FMO-like	p=1.2E-01	n=1
FSIP1	p=1.2E-01	n=1
GAF_2	p=1.2E-01	n=1
GATA	p=1.2E-01	n=2
Hairy_orange	p=1.2E-01	n=1
HJURP_C	p=1.2E-01	n=1
HUN	p=1.2E-01	n=1
INSIG	p=1.2E-01	n=1
Isy1	p=1.2E-01	n=1
KOW	p=1.2E-01	n=1

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



$-\log_{10}(p)$
n=232/211 input genes with annotations



fraction