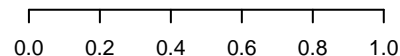


Elav_Nvec_vc1.1_XM_032369367.2

fraction genes in fg and bg

Adaptin_N	p=9.1E-02	n=2
Alpha_adaptin_C	p=9.1E-02	n=1
AP3B1_C	p=9.1E-02	n=1
CBF_beta	p=9.1E-02	n=1
CCDC158	p=9.1E-02	n=1
CEBP_ZZ	p=9.1E-02	n=1
CHDCT2	p=9.1E-02	n=1
CHDNT	p=9.1E-02	n=1
CID	p=9.1E-02	n=2
Coatomer_WDAD	p=9.1E-02	n=1
CREPT	p=9.1E-02	n=1
DBB	p=9.1E-02	n=1
DIRP	p=9.1E-02	n=1
DMAP1	p=9.1E-02	n=1
Dopey_N	p=9.1E-02	n=1
DOT1	p=9.1E-02	n=1
DUF1086	p=9.1E-02	n=1
DUF1087	p=9.1E-02	n=1
DUF3512	p=9.1E-02	n=1
DUF3543	p=9.1E-02	n=1
DUF3583	p=9.1E-02	n=1
DUF4598	p=9.1E-02	n=1
DUF4611	p=9.1E-02	n=1
EF_assoc_1	p=9.1E-02	n=1
EF_assoc_2	p=9.1E-02	n=1
EF-hand_9	p=9.1E-02	n=1
elf3g	p=9.1E-02	n=1
Glycos_transf_2	p=9.1E-02	n=3
GOLGA2L5	p=9.1E-02	n=1
HAUS6_N	p=9.1E-02	n=1

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



-log₁₀(p)
n=280/248 input genes with annotations

fraction