

Elav_Nvec_vc1.1_XM_032386992.2

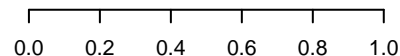
fraction genes in fg and bg

Actin	p=1.1E-01	n=2
Aldose_epim	p=1.1E-01	n=1
AMMECR1	p=1.1E-01	n=1
ANAPC5	p=1.1E-01	n=1
ATP_synt_H	p=1.1E-01	n=1
ATP-grasp_2	p=1.1E-01	n=1
Atthog	p=1.1E-01	n=1
AtuA	p=1.1E-01	n=1
Bravo_FIGEY	p=1.1E-01	n=1
BTB_2	p=1.1E-01	n=5
BTD	p=1.1E-01	n=1
CaM_bdg_C0	p=1.1E-01	n=1
CD225	p=1.1E-01	n=2
CHDCT2	p=1.1E-01	n=1
CHDNT	p=1.1E-01	n=1
Chromo	p=1.1E-01	n=2
CLN3	p=1.1E-01	n=2
DOR	p=1.1E-01	n=1
DUF1086	p=1.1E-01	n=1
DUF1087	p=1.1E-01	n=1
DUF11	p=1.1E-01	n=1
DUF1115	p=1.1E-01	n=1
DUF2370	p=1.1E-01	n=1
DUF4419	p=1.1E-01	n=1
Dynactin	p=1.1E-01	n=1
EGF_CA	p=1.1E-01	n=3
ETS_PEA3_N	p=1.1E-01	n=1
FGAR-AT_linker	p=1.1E-01	n=1
FGAR-AT_N	p=1.1E-01	n=1
FTHFS	p=1.1E-01	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
fg=0.00	bg=0.00
fg=0.01	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
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00

 $-\log_{10}(p)$

n=275/220 input genes with annotations



fraction