

Elav\_Nvec\_vc1.1\_XM\_032382466.2

fraction genes in fg and bg

YqaJ	p=3.5E-03	n=3
5-nucleotidase	p=8.1E-02	n=1
A_deaminase	p=8.1E-02	n=1
AdoHcyase	p=8.1E-02	n=1
AdoHcyase_NAD	p=8.1E-02	n=1
AIP3	p=8.1E-02	n=1
ARA70	p=8.1E-02	n=1
B12-binding_2	p=8.1E-02	n=1
CaM_bdg_C0	p=8.1E-02	n=1
CLN3	p=8.1E-02	n=2
COMPASS-Shg1	p=8.1E-02	n=1
DAZAP2	p=8.1E-02	n=1
DC_STAMP	p=8.1E-02	n=1
DUF3105	p=8.1E-02	n=1
DUF3454	p=8.1E-02	n=1
DUF4502	p=8.1E-02	n=1
DUF4503	p=8.1E-02	n=1
eIF-3_zeta	p=8.1E-02	n=1
ETF_alpha	p=8.1E-02	n=1
FACT-Spt16_Nlob	p=8.1E-02	n=1
FAM92	p=8.1E-02	n=1
GKAP	p=8.1E-02	n=1
GTP_EFTU_D2	p=8.1E-02	n=2
Headcase	p=8.1E-02	n=1
HECA	p=8.1E-02	n=1
HJURP_C	p=8.1E-02	n=1
HSNSD	p=8.1E-02	n=1
IRF-2BP1_2	p=8.1E-02	n=1
Mab-21	p=8.1E-02	n=2
Mago-bind	p=8.1E-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
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
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=0.00	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

0 1 2 3 4 5

0.0 0.2 0.4 0.6 0.8 1.0

-log<sub>10</sub>(p)

fraction

n=197/182 input genes with annotations