

Elav_Nvec_vc1.1_XM_001639164.3

fraction genes in fg and bg

| | | |
|-----------------|-----------|-----|
| Ashwin | p=8.7E-02 | n=1 |
| ATP-synt_ab_Xtn | p=8.7E-02 | n=1 |
| Atthog | p=8.7E-02 | n=2 |
| BIRC6 | p=8.7E-02 | n=1 |
| CASP_C | p=8.7E-02 | n=1 |
| CBFD_NFYB_HMF | p=8.7E-02 | n=2 |
| COQ7 | p=8.7E-02 | n=1 |
| CTF_NFI | p=8.7E-02 | n=1 |
| Cu_amine_oxid | p=8.7E-02 | n=1 |
| Cu_amine_oxidN2 | p=8.7E-02 | n=1 |
| DFP | p=8.7E-02 | n=1 |
| DNA_methylase | p=8.7E-02 | n=1 |
| DUF21 | p=8.7E-02 | n=1 |
| DUF2370 | p=8.7E-02 | n=1 |
| DUF3454 | p=8.7E-02 | n=1 |
| DUF3715 | p=8.7E-02 | n=1 |
| DUF4495 | p=8.7E-02 | n=1 |
| DUF4611 | p=8.7E-02 | n=1 |
| DUF4795 | p=8.7E-02 | n=1 |
| DUF726 | p=8.7E-02 | n=1 |
| DUF758 | p=8.7E-02 | n=1 |
| E3_UbLigase_R4 | p=8.7E-02 | n=1 |
| EF-hand_5 | p=8.7E-02 | n=3 |
| ELM2 | p=8.7E-02 | n=2 |
| FAM53 | p=8.7E-02 | n=1 |
| FDF | p=8.7E-02 | n=1 |
| Fer4_24 | p=8.7E-02 | n=1 |
| FHA_2 | p=8.7E-02 | n=1 |
| Fra10Ac1 | p=8.7E-02 | n=1 |
| HAND | p=8.7E-02 | n=1 |

| | |
|---------|---------|
| 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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$-\log_{10}(p)$
n=263/233 input genes with annotations



fraction