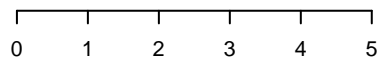


Elav_Nvec_vc1.1_XM_048726662.1

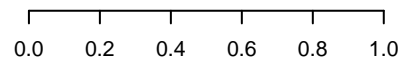
fraction genes in fg and bg

| | | |
|-----------------|-----------|-----|
| BtpA | p=7.2E-02 | n=1 |
| CaM_bdg_C0 | p=7.2E-02 | n=1 |
| CKAP2_C | p=7.2E-02 | n=1 |
| Complex1_51K | p=7.2E-02 | n=1 |
| DIM1 | p=7.2E-02 | n=1 |
| DTC | p=7.2E-02 | n=1 |
| DUF1682 | p=7.2E-02 | n=1 |
| DUF3591 | p=7.2E-02 | n=1 |
| eIF-3_zeta | p=7.2E-02 | n=1 |
| ETC_C1_NDUFA4 | p=7.2E-02 | n=1 |
| FAM117 | p=7.2E-02 | n=1 |
| Fz | p=7.2E-02 | n=2 |
| HD_3 | p=7.2E-02 | n=1 |
| Kringle | p=7.2E-02 | n=2 |
| Lipin_mid | p=7.2E-02 | n=1 |
| Lipin_N | p=7.2E-02 | n=1 |
| LNS2 | p=7.2E-02 | n=1 |
| Mago_nashi | p=7.2E-02 | n=1 |
| Med23 | p=7.2E-02 | n=1 |
| Meiosis_expr | p=7.2E-02 | n=1 |
| Myotub-related | p=7.2E-02 | n=2 |
| NADH_4Fe-4S | p=7.2E-02 | n=1 |
| Nucleoporin_FG2 | p=7.2E-02 | n=1 |
| Nup153 | p=7.2E-02 | n=1 |
| PALP | p=7.2E-02 | n=2 |
| PBC | p=7.2E-02 | n=1 |
| Peptidase_C98 | p=7.2E-02 | n=1 |
| Pirin | p=7.2E-02 | n=1 |
| Pirin_C | p=7.2E-02 | n=1 |
| RanGAP1_C | p=7.2E-02 | n=1 |

| | |
|---------|---------|
| 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=0.01 | 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 |
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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 |
| fg=0.00 | bg=0.00 |

-log₁₀(p)

n=192/171 input genes with annotations



fraction