

Ncol_Nvec_vc1.1_XM_032384300.2

fraction genes in fg and bg

| | | |
|-----------------|-----------|-----|
| 2OG-Fell_Oxy | p=7.8E-02 | n=2 |
| DIOX_N | p=7.8E-02 | n=2 |
| Glyco_transf_92 | p=7.8E-02 | n=2 |
| UMPH-1 | p=7.8E-02 | n=2 |
| 5-nucleotidase | p=1.2E-01 | n=1 |
| Acetyltransf_13 | p=1.2E-01 | n=1 |
| Aida_C2 | p=1.2E-01 | n=1 |
| Aida_N | p=1.2E-01 | n=1 |
| ATG16 | p=1.2E-01 | n=1 |
| Autophagy_act_C | p=1.2E-01 | n=1 |
| Biotin_lipoyl_2 | p=1.2E-01 | n=1 |
| BTB_2 | p=1.2E-01 | n=4 |
| CAP-ZIP_m | p=1.2E-01 | n=1 |
| CCSAP | p=1.2E-01 | n=1 |
| CDC50 | p=1.2E-01 | n=1 |
| CENP-T_C | p=1.2E-01 | n=1 |
| Cl-B14_5a | p=1.2E-01 | n=1 |
| CLU | p=1.2E-01 | n=1 |
| CLU_N | p=1.2E-01 | n=1 |
| CTF_NFI | p=1.2E-01 | n=1 |
| DHFR_1 | p=1.2E-01 | n=1 |
| DNA_primase_lrg | p=1.2E-01 | n=1 |
| DUF2205 | p=1.2E-01 | n=1 |
| DUF4485 | p=1.2E-01 | n=1 |
| DUF4611 | p=1.2E-01 | n=1 |
| EF-hand_5 | p=1.2E-01 | n=3 |
| eIF3_p135 | p=1.2E-01 | n=1 |
| ERO1 | p=1.2E-01 | n=1 |
| Es2 | p=1.2E-01 | n=1 |
| FAM72 | p=1.2E-01 | n=1 |

| | |
|---------|---------|
| 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 |
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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=322/304 input genes with annotations



fraction