

| | | |
|-----------------|-----------|-----|
| AA_permease_2 | p=1.0E-01 | n=3 |
| ABC1 | p=1.0E-01 | n=2 |
| Acyltransferase | p=1.0E-01 | n=2 |
| AP3D1 | p=1.0E-01 | n=1 |
| ATP_synt_E | p=1.0E-01 | n=1 |
| BAT2_N | p=1.0E-01 | n=1 |
| CAP_N | p=1.0E-01 | n=1 |
| CBM_21 | p=1.0E-01 | n=2 |
| CCDC74_C | p=1.0E-01 | n=1 |
| CCSAP | p=1.0E-01 | n=1 |
| CKAP2_C | p=1.0E-01 | n=1 |
| CoA_trans | p=1.0E-01 | n=1 |
| CybS | p=1.0E-01 | n=1 |
| Dimer_Tnp_hAT | p=1.0E-01 | n=2 |
| DUF2451 | p=1.0E-01 | n=1 |
| DUF3689 | p=1.0E-01 | n=1 |
| DUF4371 | p=1.0E-01 | n=2 |
| DUF4464 | p=1.0E-01 | n=1 |
| DUF5745 | p=1.0E-01 | n=1 |
| E3_UbLigase_EDD | p=1.0E-01 | n=1 |
| F_actin_cap_B | p=1.0E-01 | n=1 |
| FANCI_HD1 | p=1.0E-01 | n=1 |
| FANCI_HD2 | p=1.0E-01 | n=1 |
| FANCI_S1 | p=1.0E-01 | n=1 |
| FANCI_S1-cap | p=1.0E-01 | n=1 |
| FANCI_S2 | p=1.0E-01 | n=1 |
| FANCI_S3 | p=1.0E-01 | n=1 |
| FANCI_S4 | p=1.0E-01 | n=1 |
| Filament | p=1.0E-01 | n=1 |
| Formyl_trans_C | p=1.0E-01 | n=1 |

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|---------|---------|
| fg=0.01 | bg=0.00 |
| fg=0.01 | bg=0.00 |
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$-\log_{10}(p)$
n=320/279 input genes with annotations



fraction