

AAA_11	p=1.2E-01	n=2
AAA_12	p=1.2E-01	n=2
AcetylCoA_hyd_C	p=1.2E-01	n=1
AcetylCoA_hydro	p=1.2E-01	n=1
Actin	p=1.2E-01	n=3
Agenet	p=1.2E-01	n=1
ANAPC4_WD40	p=1.2E-01	n=3
BolA	p=1.2E-01	n=1
CCDC50_N	p=1.2E-01	n=1
CID	p=1.2E-01	n=2
Clathrin_lg_ch	p=1.2E-01	n=1
COX5B	p=1.2E-01	n=1
CTC1	p=1.2E-01	n=1
CUTL	p=1.2E-01	n=1
Cwf_Cwc_15	p=1.2E-01	n=1
Diphthamide_syn	p=1.2E-01	n=1
DOR	p=1.2E-01	n=1
DUF2236	p=1.2E-01	n=1
DUF3454	p=1.2E-01	n=1
DUF3512	p=1.2E-01	n=1
DUF3543	p=1.2E-01	n=1
DUF3583	p=1.2E-01	n=1
DUF3808	p=1.2E-01	n=1
DUF92	p=1.2E-01	n=1
EF-hand_2	p=1.2E-01	n=2
EF-hand_3	p=1.2E-01	n=2
EF-hand_9	p=1.2E-01	n=1
eIF-1a	p=1.2E-01	n=1
ELM2	p=1.2E-01	n=2
ETF_alpha	p=1.2E-01	n=1

fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.01	bg=0.00
fg=0.01	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=0.01	bg=0.00
fg=0.01	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



$-\log_{10}(p)$
n=299/275 input genes with annotations



fraction