

2Fe-2S_thioredx	p=2.2E-01	n=1
6PF2K	p=2.2E-01	n=1
AA_kinase	p=2.2E-01	n=1
AcetylCoA_hyd_C	p=2.2E-01	n=1
AcetylCoA_hydro	p=2.2E-01	n=1
Aldedh	p=2.2E-01	n=2
Amino_oxidase	p=2.2E-01	n=3
Antistatin	p=2.2E-01	n=1
ApeC	p=2.2E-01	n=1
BTB_2	p=2.2E-01	n=6
Bystin	p=2.2E-01	n=1
CENP-O	p=2.2E-01	n=1
Chromo	p=2.2E-01	n=2
cNMP_binding	p=2.2E-01	n=4
CUTL	p=2.2E-01	n=1
Dicty_CAR	p=2.2E-01	n=1
DND1_DSRM	p=2.2E-01	n=1
DOR	p=2.2E-01	n=1
DUF1151	p=2.2E-01	n=1
DUF3399	p=2.2E-01	n=1
DUF3583	p=2.2E-01	n=1
DUF3808	p=2.2E-01	n=1
DUF4505	p=2.2E-01	n=1
DUF4506	p=2.2E-01	n=1
DUF4795	p=2.2E-01	n=1
DUF974	p=2.2E-01	n=1
ECH_1	p=2.2E-01	n=2
EF-hand_like	p=2.2E-01	n=2
EGF	p=2.2E-01	n=8
EIF4E-T	p=2.2E-01	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=0.00	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=0.00	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



$-\log_{10}(p)$
n=407/463 input genes with annotations



fraction