

Acyl-CoA_dh_1	p=1.5E-01	n=2
Acyl-CoA_dh_N	p=1.5E-01	n=2
ANAPC5	p=1.5E-01	n=1
ARPC4	p=1.5E-01	n=1
Biotin_lipoyl	p=1.5E-01	n=2
Bravo_FIGEY	p=1.5E-01	n=1
bZIP_1	p=1.5E-01	n=4
CENP-O	p=1.5E-01	n=1
Cobalamin_bind	p=1.5E-01	n=1
Coiled-coil_56	p=1.5E-01	n=1
CRAL_TRIO	p=1.5E-01	n=2
Cript	p=1.5E-01	n=1
Cyt-b5	p=1.5E-01	n=2
Cytochrom_C1	p=1.5E-01	n=1
DDDD	p=1.5E-01	n=1
DUF2152	p=1.5E-01	n=1
DUF3105	p=1.5E-01	n=1
DUF4430	p=1.5E-01	n=1
DUF719	p=1.5E-01	n=1
Dynactin_p62	p=1.5E-01	n=1
ERO1	p=1.5E-01	n=1
Exo_endo_phos_2	p=1.5E-01	n=1
FAM221	p=1.5E-01	n=1
Fer4_7	p=1.5E-01	n=1
Fic	p=1.5E-01	n=1
FragX_IP	p=1.5E-01	n=1
Gaa1	p=1.5E-01	n=1
GCN5L1	p=1.5E-01	n=1
Glucosamine_iso	p=1.5E-01	n=1
Glyco_hydro_47	p=1.5E-01	n=2

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



$-\log_{10}(p)$
n=302/279 input genes with annotations



fraction