

Acetyltransf_3	p=1.3E-01	n=1
ACT_7	p=1.3E-01	n=1
Aminotran_1_2	p=1.3E-01	n=2
ASF1_hist_chap	p=1.3E-01	n=1
ATP_Ca_trans_C	p=1.3E-01	n=1
BCCT	p=1.3E-01	n=1
Castor1_N	p=1.3E-01	n=1
CBFB_NFYA	p=1.3E-01	n=1
CDH1_2_SANT_HL1	p=1.3E-01	n=1
COQ7	p=1.3E-01	n=1
DHC_N1	p=1.3E-01	n=2
DHR10	p=1.3E-01	n=1
DUF3668	p=1.3E-01	n=1
DUF3719	p=1.3E-01	n=1
DUF4149	p=1.3E-01	n=1
DUF4208	p=1.3E-01	n=1
DUF719	p=1.3E-01	n=1
DUF758	p=1.3E-01	n=1
DUF89	p=1.3E-01	n=1
EF-hand_7	p=1.3E-01	n=7
EF-hand_9	p=1.3E-01	n=1
EloA-BP1	p=1.3E-01	n=1
Fam20C	p=1.3E-01	n=1
Fer4_24	p=1.3E-01	n=1
FHA_2	p=1.3E-01	n=1
GAF_2	p=1.3E-01	n=1
HAUS4	p=1.3E-01	n=1
HSL_N	p=1.3E-01	n=1
JIP_LZII	p=1.3E-01	n=1
Jnk-SapK_ap_N	p=1.3E-01	n=1

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
fg=0.00	bg=0.00
fg=0.00	bg=0.00
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fg=0.01	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
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.02	bg=0.01
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
fg=0.00	bg=0.00
fg=0.00	bg=0.00



$-\log_{10}(p)$
n=308/282 input genes with annotations



fraction