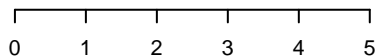


AC_N	p=6.7E-02	n=3
2-Hacid_dh	p=1.4E-01	n=2
2-Hacid_dh_C	p=1.4E-01	n=2
A1_Propeptide	p=1.4E-01	n=1
adh_short	p=1.4E-01	n=4
Ala_racemase_N	p=1.4E-01	n=1
AlaDh_PNT_C	p=1.4E-01	n=1
Alpha-amylase_C	p=1.4E-01	n=1
ANAPC16	p=1.4E-01	n=1
Ank_5	p=1.4E-01	n=3
Arg_tRNA_synt_N	p=1.4E-01	n=1
Arginosuc_synt	p=1.4E-01	n=1
ARPC4	p=1.4E-01	n=1
Asp_Arg_Hydrox	p=1.4E-01	n=1
ATP-synt_E	p=1.4E-01	n=1
ATP-synt_F	p=1.4E-01	n=1
Autophagy_act_C	p=1.4E-01	n=1
BAG6	p=1.4E-01	n=1
Biotin_lipoyl_2	p=1.4E-01	n=1
Bravo_FIGEY	p=1.4E-01	n=1
CBM_48	p=1.4E-01	n=1
CCDC142	p=1.4E-01	n=1
CDC27	p=1.4E-01	n=1
CDC37_C	p=1.4E-01	n=1
CDC37_M	p=1.4E-01	n=1
CDC37_N	p=1.4E-01	n=1
CDC73_C	p=1.4E-01	n=1
CDC73_N	p=1.4E-01	n=1
CDI	p=1.4E-01	n=1
CENP-K	p=1.4E-01	n=1



-log<sub>10</sub>(p)  
n=502/473 input genes with annotations

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=0.00	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
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



fraction