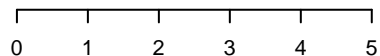


Acetyltransf_16	p=1.2E-01	n=1
APCDDC	p=1.2E-01	n=1
APP_amyloid	p=1.2E-01	n=1
APP_Cu_bd	p=1.2E-01	n=1
APP_E2	p=1.2E-01	n=1
APP_N	p=1.2E-01	n=1
CaM_bdg_C0	p=1.2E-01	n=1
CCDC158	p=1.2E-01	n=1
CDV3	p=1.2E-01	n=1
Coatomer_WDAD	p=1.2E-01	n=1
COLFI	p=1.2E-01	n=1
Cys_Met_Meta_PP	p=1.2E-01	n=1
DNA_ligase_A_C	p=1.2E-01	n=1
DNA_ligase_A_M	p=1.2E-01	n=1
DNA_ligase_A_N	p=1.2E-01	n=1
DUF4659	p=1.2E-01	n=1
DUF758	p=1.2E-01	n=1
EAF	p=1.2E-01	n=1
EF-hand_7	p=1.2E-01	n=5
ELM2	p=1.2E-01	n=2
FA_FANCE	p=1.2E-01	n=1
Fanconi_A_N	p=1.2E-01	n=1
FSIP1	p=1.2E-01	n=1
GAF_2	p=1.2E-01	n=1
GATA	p=1.2E-01	n=2
GRAM	p=1.2E-01	n=2
Guanylate_kin	p=1.2E-01	n=2
Hairy_orange	p=1.2E-01	n=1
HJURP_C	p=1.2E-01	n=1
HUN	p=1.2E-01	n=1



$-\log_{10}(p)$
n=231/211 input genes with annotations

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



fraction