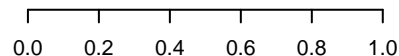


gc027

fraction genes in fg and bg

APS_kinase	p=1.4E-02	n=1
ARS2	p=1.4E-02	n=1
ATP-sulfurylase	p=1.4E-02	n=1
Bromodomain	p=1.4E-02	n=2
Btz	p=1.4E-02	n=1
CCDC32	p=1.4E-02	n=1
DSL	p=1.4E-02	n=1
DUF1077	p=1.4E-02	n=1
FKBP_C	p=1.4E-02	n=2
HAND	p=1.4E-02	n=1
Med21	p=1.4E-02	n=1
Med4	p=1.4E-02	n=1
Memo	p=1.4E-02	n=1
MNNL	p=1.4E-02	n=1
NOPS	p=1.4E-02	n=1
preSET_CXC	p=1.4E-02	n=1
PUA_2	p=1.4E-02	n=1
RPN6_C_helix	p=1.4E-02	n=1
RPN6_N	p=1.4E-02	n=1
RRM_1	p=1.4E-02	n=5
Sep15_SelM	p=1.4E-02	n=1
SERRATE_Ars2_N	p=1.4E-02	n=1
SLIDE	p=1.4E-02	n=1
SOSSC	p=1.4E-02	n=1
SPC12	p=1.4E-02	n=1
TMEM33_Pom33	p=1.4E-02	n=1
Topo_C_assoc	p=1.4E-02	n=1
Topoisom_I	p=1.4E-02	n=1
Topoisom_I_N	p=1.4E-02	n=1
TPR_9	p=1.4E-02	n=1

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



$-\log_{10}(p)$
n=81/70 input genes with annotations

fraction