

gc002b

fraction genes in fg and bg

DUF5600	p=5.0E-03	n=2
EHD_N	p=5.0E-03	n=2
WD40_2	p=5.0E-03	n=2
EF-hand_4	p=2.2E-02	n=2
Ank_2	p=3.9E-02	n=5
CEP170_C	p=4.5E-02	n=1
Dynamin_N	p=4.5E-02	n=2
Esterase	p=4.5E-02	n=1
PRT_C	p=4.5E-02	n=1
Trp_Tyr_perm	p=4.5E-02	n=1
GAT	p=5.1E-02	n=1
Ndc80_HEC	p=5.1E-02	n=1
PH_14	p=5.1E-02	n=1
Ras	p=5.1E-02	n=3
SH3BP5	p=5.1E-02	n=1
TB2_DP1_HVA22	p=5.1E-02	n=1
zf-LITAF-like	p=5.1E-02	n=1
AAA_11	p=7.5E-02	n=1
AAA_12	p=7.5E-02	n=1
Anoctamin	p=7.5E-02	n=1
DUF4440	p=7.5E-02	n=1
EF-hand_like	p=7.5E-02	n=1
FA_hydroxylase	p=7.5E-02	n=1
PI-PLC-X	p=7.5E-02	n=1
PI-PLC-Y	p=7.5E-02	n=1
PXA	p=7.5E-02	n=1
TspO_MBR	p=7.5E-02	n=1
VHS	p=7.5E-02	n=1
zf-AN1	p=7.5E-02	n=1
C2	p=8.1E-02	n=2

fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.06	bg=0.01
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=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.03	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
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
fg=0.02	bg=0.00

 $-\log_{10}(p)$

n=75/61 input genes with annotations



fraction