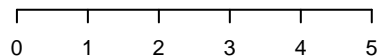


gc003b

fraction genes in fg and bg

Exo_endo_phos	p=3.8E-03	n=2
PAX	p=2.0E-02	n=1
7tm_1	p=2.8E-02	n=4
7tm_3	p=2.8E-02	n=2
ANF_receptor	p=2.8E-02	n=2
RasGEF_N	p=2.8E-02	n=1
Homeodomain	p=7.1E-02	n=1
RhoGEF	p=7.1E-02	n=1
CUB	p=8.2E-02	n=1
GAIN	p=8.2E-02	n=1
I-set	p=1.0E-01	n=1
Ig_3	p=1.1E-01	n=1
GPS	p=1.1E-01	n=1
7tm_2	p=1.5E-01	n=1
Pkinase	p=2.0E-01	n=1



-log₁₀(p)
n=15/18 input genes with annotations

fg=0.10	bg=0.00
fg=0.05	bg=0.00
fg=0.19	bg=0.04
fg=0.10	bg=0.01
fg=0.10	bg=0.01
fg=0.05	bg=0.00
fg=0.05	bg=0.00
fg=0.05	bg=0.00
fg=0.05	bg=0.00
fg=0.05	bg=0.00
fg=0.05	bg=0.00
fg=0.05	bg=0.00
fg=0.05	bg=0.00
fg=0.05	bg=0.01
fg=0.05	bg=0.01



fraction