

gc015

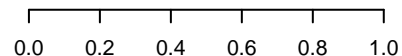
fraction genes in fg and bg

7TM_GPCR_Srx	p=1.1E-02	n=1
Homeobox_KN	p=1.7E-02	n=1
LRR_5	p=4.1E-02	n=1
Aa_trans	p=5.3E-02	n=1
7tm_1	p=6.2E-02	n=3
GPS	p=6.6E-02	n=1
Lectin_C	p=7.1E-02	n=1
7tm_2	p=9.4E-02	n=1
7tm_3	p=1.0E-01	n=1
ANF_receptor	p=1.1E-01	n=1
LRR_8	p=1.1E-01	n=1
Pkinase	p=1.5E-01	n=1

fg=0.07	bg=0.00
fg=0.07	bg=0.00
fg=0.07	bg=0.00
fg=0.07	bg=0.00
fg=0.21	bg=0.05
fg=0.07	bg=0.00
fg=0.07	bg=0.00
fg=0.07	bg=0.00
fg=0.07	bg=0.01
fg=0.07	bg=0.01
fg=0.07	bg=0.01
fg=0.07	bg=0.01

-log₁₀(p)

n=12/27 input genes with annotations



fraction