

Fox_Nvec_vc1.1_XM_032364121.2

fraction genes in fg and bg

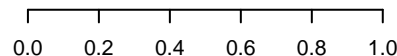
4HBT_2	p=5.2E-02	n=1
7tm_3	p=5.2E-02	n=3
ABC_ATPase	p=5.2E-02	n=1
ALS2CR8	p=5.2E-02	n=1
ANF_receptor	p=5.2E-02	n=3
BAT2_N	p=5.2E-02	n=1
BNIP3	p=5.2E-02	n=1
DNA_ligase_A_C	p=5.2E-02	n=1
DNA_ligase_A_M	p=5.2E-02	n=1
DNA_ligase_A_N	p=5.2E-02	n=1
DUF3384	p=5.2E-02	n=1
DUF4200	p=5.2E-02	n=1
FOXP-CC	p=5.2E-02	n=1
LIG3_BRCT	p=5.2E-02	n=1
Macoilin	p=5.2E-02	n=1
MORN	p=5.2E-02	n=2
N_BRCA1_IG	p=5.2E-02	n=1
Ndr	p=5.2E-02	n=1
PFU	p=5.2E-02	n=1
PUL	p=5.2E-02	n=1
SKIP_SNW	p=5.2E-02	n=1
SPA	p=5.2E-02	n=1
Tuberin	p=5.2E-02	n=1
zf-PARP	p=5.2E-02	n=1
Zmiz1_N	p=5.2E-02	n=1
7tm_1	p=7.0E-02	n=8
DUF1154	p=7.0E-02	n=1
Glyco_hydro_15	p=7.0E-02	n=1
HTH_Tnp_Tc5	p=7.0E-02	n=1
Mpv17_PMP22	p=7.0E-02	n=1

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=0.03	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
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fg=0.02	bg=0.00
fg=0.01	bg=0.00
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fg=0.01	bg=0.00
fg=0.07	bg=0.03
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00



-log10(p)

n=102/81 input genes with annotations



fraction