

Fox_Nvec_vc1.1_XM_032372264.2

fraction genes in fg and bg

AF-4	p=9.4E-02	n=1
AF-4_C	p=9.4E-02	n=1
Ank_2	p=9.4E-02	n=6
APP_amyloid	p=9.4E-02	n=1
APP_Cu_bd	p=9.4E-02	n=1
APP_E2	p=9.4E-02	n=1
APP_N	p=9.4E-02	n=1
Arm_vesicular	p=9.4E-02	n=1
Astacin	p=9.4E-02	n=3
ASXH	p=9.4E-02	n=1
B12D	p=9.4E-02	n=1
Band_7	p=9.4E-02	n=2
CaM_bdg_C0	p=9.4E-02	n=1
CAP	p=9.4E-02	n=2
Cation_ATPase	p=9.4E-02	n=2
CDC50	p=9.4E-02	n=1
CEBP_ZZ	p=9.4E-02	n=1
CLAMP	p=9.4E-02	n=1
Cys_Met_Meta_PP	p=9.4E-02	n=1
DNA_pol_delta_4	p=9.4E-02	n=1
Dopey_N	p=9.4E-02	n=1
DUF2370	p=9.4E-02	n=1
DUF3583	p=9.4E-02	n=1
EAF	p=9.4E-02	n=1
ELM2	p=9.4E-02	n=2
FA_FANCE	p=9.4E-02	n=1
Hairy_orange	p=9.4E-02	n=1
ICA69	p=9.4E-02	n=1
MAM	p=9.4E-02	n=2
MgtE	p=9.4E-02	n=1

fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.03	bg=0.01
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.01	bg=0.00
fg=0.00	bg=0.00
fg=0.01	bg=0.00
fg=0.00	bg=0.00
fg=0.01	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.01	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.01	bg=0.00
fg=0.00	bg=0.00

 $-\log_{10}(p)$

n=194/153 input genes with annotations



fraction