

Fox_Nvec_vc1.1_XM_032361813.2

fraction genes in fg and bg

APP_amyloid	p=9.1E-02	n=1
APP_Cu_bd	p=9.1E-02	n=1
APP_E2	p=9.1E-02	n=1
APP_N	p=9.1E-02	n=1
Astacin	p=9.1E-02	n=2
DER1	p=9.1E-02	n=1
DNA_pol_B	p=9.1E-02	n=1
DNA_pol_B_exo1	p=9.1E-02	n=1
DUF1387	p=9.1E-02	n=1
DUF4200	p=9.1E-02	n=1
DUF758	p=9.1E-02	n=1
FAM117	p=9.1E-02	n=1
FAM219A	p=9.1E-02	n=1
FANCAA	p=9.1E-02	n=1
FH2	p=9.1E-02	n=2
Formin_GBD_N	p=9.1E-02	n=1
Glyco_hydro_99	p=9.1E-02	n=1
Glyco_transf_41	p=9.1E-02	n=1
GpcrRhopsn4	p=9.1E-02	n=2
HAUS-augmin3	p=9.1E-02	n=1
HSF_DNA-bind	p=9.1E-02	n=1
IER	p=9.1E-02	n=1
IRF-2BP1_2	p=9.1E-02	n=1
LKAAEAR	p=9.1E-02	n=1
MAM	p=9.1E-02	n=2
MM_CoA_mutase	p=9.1E-02	n=1
NCD1	p=9.1E-02	n=1
NCD2	p=9.1E-02	n=1
Njmu-R1	p=9.1E-02	n=1
Porin_3	p=9.1E-02	n=1

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=0.00	bg=0.00
fg=0.00	bg=0.00
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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=0.00	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=0.00	bg=0.00



$-\log_{10}(p)$
n=185/142 input genes with annotations



fraction