

Fox_Nvec_vc1.1_XM_001622708.3

fraction genes in fg and bg

5-nucleotidase	p=8.4E-02	n=1
AAA_33	p=8.4E-02	n=2
APCDDC	p=8.4E-02	n=1
BAAT_C	p=8.4E-02	n=1
Bile_Hydr_Trans	p=8.4E-02	n=1
BtpA	p=8.4E-02	n=1
CASP_C	p=8.4E-02	n=1
CDC73_C	p=8.4E-02	n=1
CDC73_N	p=8.4E-02	n=1
Complex1_LYR	p=8.4E-02	n=2
CortBP2	p=8.4E-02	n=1
DAP	p=8.4E-02	n=1
DEP	p=8.4E-02	n=2
DHR10	p=8.4E-02	n=1
DNA_pol_D_N	p=8.4E-02	n=1
DOR	p=8.4E-02	n=1
DUF1771	p=8.4E-02	n=1
DUF2181	p=8.4E-02	n=1
DUF2451	p=8.4E-02	n=1
Elong_Iki1	p=8.4E-02	n=1
ETS_PEA3_N	p=8.4E-02	n=1
FAM219A	p=8.4E-02	n=1
FANCL_C	p=8.4E-02	n=1
FANCL_d1	p=8.4E-02	n=1
FANCL_d2	p=8.4E-02	n=1
FANCL_d3	p=8.4E-02	n=1
Fascin	p=8.4E-02	n=1
Fer2_4	p=8.4E-02	n=1
FHA_2	p=8.4E-02	n=1
Galactosyl_T	p=8.4E-02	n=3

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



-log10(p)

n=280/214 input genes with annotations



fraction