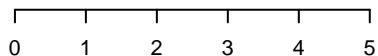


Fox_Nvec_vc1.1_XM_048722445.1

fraction genes in fg and bg

Neurensin	p=5.8E-02	n=2
OSR1_C	p=5.8E-02	n=2
ACBP	p=7.7E-02	n=2
Acetyltransf_16	p=7.7E-02	n=1
ARA70	p=7.7E-02	n=1
BCL_N	p=7.7E-02	n=1
Calx-beta	p=7.7E-02	n=2
CIAPIN1	p=7.7E-02	n=1
CSN5_C	p=7.7E-02	n=1
DÉP	p=7.7E-02	n=2
DNA_gyraseB	p=7.7E-02	n=1
DNA_topoisolV	p=7.7E-02	n=1
DUF2181	p=7.7E-02	n=1
DUF4460	p=7.7E-02	n=1
DUF4461	p=7.7E-02	n=1
DUF4795	p=7.7E-02	n=1
IML1	p=7.7E-02	n=1
JAB	p=7.7E-02	n=2
Leg1	p=7.7E-02	n=1
MCM	p=7.7E-02	n=2
MCM_lid	p=7.7E-02	n=2
MCM_OB	p=7.7E-02	n=2
MCM6_C	p=7.7E-02	n=1
Med13_C	p=7.7E-02	n=1
MID_MedPIWI	p=7.7E-02	n=1
MM_CoA_mutase	p=7.7E-02	n=1
MRG	p=7.7E-02	n=1
MRP_L53	p=7.7E-02	n=1
MYCBPAP	p=7.7E-02	n=1
Myosin-VI_CBD	p=7.7E-02	n=1

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

 $-\log_{10}(p)$

n=242/199 input genes with annotations



fraction