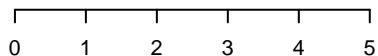


Fox\_Nvec\_vc1.1\_XM\_032371461.2

fraction genes in fg and bg

2Fe-2S_thioredx	p=7.9E-02	n=1
2OG-FeII_Oxy_2	p=7.9E-02	n=2
Aldolase_II	p=7.9E-02	n=2
ALG11_N	p=7.9E-02	n=1
ATP-synt_ab_C	p=7.9E-02	n=1
ATP-synt_F	p=7.9E-02	n=1
CAS_CSE1	p=7.9E-02	n=1
Cation_ATPase_C	p=7.9E-02	n=2
Cation_ATPase_N	p=7.9E-02	n=3
CCSAP	p=7.9E-02	n=1
CEBP_ZZ	p=7.9E-02	n=1
CLN3	p=7.9E-02	n=2
Cmc1	p=7.9E-02	n=1
COX7C	p=7.9E-02	n=1
Ctf8	p=7.9E-02	n=1
CTNNBL	p=7.9E-02	n=1
Cul7	p=7.9E-02	n=1
DAGAT	p=7.9E-02	n=2
Dis3l2_C_term	p=7.9E-02	n=1
DNA_pol3_delta2	p=7.9E-02	n=1
DnaJ_C	p=7.9E-02	n=2
DUF1387	p=7.9E-02	n=1
DUF2205	p=7.9E-02	n=1
DUF4612	p=7.9E-02	n=1
E1-E2_ATPase	p=7.9E-02	n=3
FANCM-MHF_bd	p=7.9E-02	n=1
Fcf1	p=7.9E-02	n=1
GEMIN8	p=7.9E-02	n=1
Helicase_PWI	p=7.9E-02	n=1
HHH_2	p=7.9E-02	n=1

fg=0.00	bg=0.00
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fg=0.01	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=251/199 input genes with annotations



fraction