

Fox_Nvec_v1g93844

fraction genes in fg and bg

CBX7_C	p=1.1E-01	n=2
CTP_transf_like	p=1.1E-01	n=2
AC_N	p=1.2E-01	n=2
AcetylCoA_hyd_C	p=1.2E-01	n=1
AcetylCoA_hydro	p=1.2E-01	n=1
AP3B1_C	p=1.2E-01	n=1
ATP-synt_E	p=1.2E-01	n=1
Biotin_lipoyl_2	p=1.2E-01	n=1
BOP1NT	p=1.2E-01	n=1
Cation_ATPase_C	p=1.2E-01	n=2
Cation_ATPase_N	p=1.2E-01	n=2
CD20	p=1.2E-01	n=2
CDC50	p=1.2E-01	n=1
CDK5_activator	p=1.2E-01	n=1
cEGF	p=1.2E-01	n=1
CTP_synth_N	p=1.2E-01	n=1
CUPID	p=1.2E-01	n=1
DinB_2	p=1.2E-01	n=1
DIRP	p=1.2E-01	n=1
Dis3l2_C_term	p=1.2E-01	n=1
DOR	p=1.2E-01	n=1
DUF1325	p=1.2E-01	n=1
DUF3543	p=1.2E-01	n=1
DUF4460	p=1.2E-01	n=1
DUF4461	p=1.2E-01	n=1
DUF4485	p=1.2E-01	n=1
DUF4500	p=1.2E-01	n=1
DUF4706	p=1.2E-01	n=1
EGF_CA	p=1.2E-01	n=3
FANCAA	p=1.2E-01	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
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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

 $-\log_{10}(p)$

n=299/237 input genes with annotations



fraction