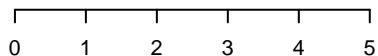


Fox_Nvec_vc1.1_XM_048719424.1

fraction genes in fg and bg

Actin	p=9.4E-02	n=2
ANAPC5	p=9.4E-02	n=1
Arm_vesicular	p=9.4E-02	n=1
Auts2	p=9.4E-02	n=1
BTB_3	p=9.4E-02	n=1
C1_2	p=9.4E-02	n=1
CAP	p=9.4E-02	n=2
Cation_ATPase	p=9.4E-02	n=3
CDH1_2_SANT_HL1	p=9.4E-02	n=1
Chorein_N	p=9.4E-02	n=2
DAG_kinase_N	p=9.4E-02	n=1
DNA_ligase_A_C	p=9.4E-02	n=1
DNA_ligase_A_M	p=9.4E-02	n=1
DNA_ligase_A_N	p=9.4E-02	n=1
DUF4208	p=9.4E-02	n=1
DUF4419	p=9.4E-02	n=1
IRF-2BP1_2	p=9.4E-02	n=1
Kazal_1	p=9.4E-02	n=1
LIG3_BRCT	p=9.4E-02	n=1
Med25	p=9.4E-02	n=1
Med25_VWA	p=9.4E-02	n=1
MNHL	p=9.4E-02	n=1
Mo25	p=9.4E-02	n=1
Na_Ca_ex	p=9.4E-02	n=2
PDE4_UCR	p=9.4E-02	n=1
PDEase_I	p=9.4E-02	n=2
PGK	p=9.4E-02	n=1
PhoLip_ATPase_C	p=9.4E-02	n=2
PhoLip_ATPase_N	p=9.4E-02	n=2
Rep_fac_C	p=9.4E-02	n=1

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

-log₁₀(p)

n=178/139 input genes with annotations



fraction