

Ncol_Nvec_vc1.1_XM_032375721.2

fraction genes in fg and bg

Meis_PKNOX_N	p=8.1E-02	n=2
AAA_5	p=8.1E-02	n=1
AP1AR	p=8.1E-02	n=1
APOBEC_N	p=8.1E-02	n=1
ATP-synt_ab_C	p=8.1E-02	n=1
Auts2	p=8.1E-02	n=1
Cul7	p=8.1E-02	n=1
DNA_pol_alpha_N	p=8.1E-02	n=1
DSPc	p=8.1E-02	n=3
Efg1	p=8.1E-02	n=1
eIF-3c_N	p=8.1E-02	n=1
eIF-5a	p=8.1E-02	n=1
Ependymin	p=8.1E-02	n=1
FAM92	p=8.1E-02	n=1
Fe-S_biosyn	p=8.1E-02	n=1
Fer2_3	p=8.1E-02	n=1
Fer4_17	p=8.1E-02	n=1
Ferric_reduct	p=8.1E-02	n=1
His_biosynth	p=8.1E-02	n=1
Homeobox_KN	p=8.1E-02	n=2
IFT46_B_C	p=8.1E-02	n=1
ILEI	p=8.1E-02	n=2
KBP_C	p=8.1E-02	n=1
Laminin_G_3	p=8.1E-02	n=3
Leg1	p=8.1E-02	n=1
Lipase	p=8.1E-02	n=1
Mt_ATP-synt_B	p=8.1E-02	n=1
Nup54	p=8.1E-02	n=1
Nup54_C	p=8.1E-02	n=1
PAN_1	p=8.1E-02	n=4

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=0.00	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
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
fg=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.01	bg=0.00
fg=0.00	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=0.00	bg=0.00
fg=0.02	bg=0.00



-log₁₀(p)
n=213/177 input genes with annotations



fraction