

Ncol_Nvec_vc1.1_XM_032371032.2

fraction genes in fg and bg

Amino_oxidase	p=7.2E-02	n=2
ANF_receptor	p=7.2E-02	n=4
BtpA	p=7.2E-02	n=1
CaM_bdg_C0	p=7.2E-02	n=1
CKAP2_C	p=7.2E-02	n=1
Complex1_51K	p=7.2E-02	n=1
CUB	p=7.2E-02	n=3
DIM1	p=7.2E-02	n=1
DTC	p=7.2E-02	n=1
DUF1682	p=7.2E-02	n=1
DUF2052	p=7.2E-02	n=1
DUF3591	p=7.2E-02	n=1
DUF4495	p=7.2E-02	n=1
DUF758	p=7.2E-02	n=1
Dynactin_p22	p=7.2E-02	n=1
EF-hand_5	p=7.2E-02	n=3
elf-3_zeta	p=7.2E-02	n=1
ETC_C1_NDUFA4	p=7.2E-02	n=1
FAM117	p=7.2E-02	n=1
GIY-YIG	p=7.2E-02	n=1
Laminin_EGF	p=7.2E-02	n=2
LepA_C	p=7.2E-02	n=1
Lig_chan	p=7.2E-02	n=2
Lig_chan-Glu_bd	p=7.2E-02	n=2
Lipin_mid	p=7.2E-02	n=1
Lipin_N	p=7.2E-02	n=1
LNS2	p=7.2E-02	n=1
Mago_nashi	p=7.2E-02	n=1
Med10	p=7.2E-02	n=1
Meiosis_expr	p=7.2E-02	n=1

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

-log₁₀(p)

n=202/223 input genes with annotations



fraction