

Ncol_Nvec_vc1.1_XM_032376608.2

fraction genes in fg and bg

Amino_oxidase	p=1.3E-01	n=2
Bap31	p=1.3E-01	n=1
Bap31_Bap29_C	p=1.3E-01	n=1
BTK	p=1.3E-01	n=1
Bystin	p=1.3E-01	n=1
CENP-O	p=1.3E-01	n=1
CUTL	p=1.3E-01	n=1
Dicty_CAR	p=1.3E-01	n=1
DND1_DSRM	p=1.3E-01	n=1
DOR	p=1.3E-01	n=1
DUF1151	p=1.3E-01	n=1
DUF2052	p=1.3E-01	n=1
DUF3583	p=1.3E-01	n=1
EF-hand_like	p=1.3E-01	n=2
FAM53	p=1.3E-01	n=1
FAM92	p=1.3E-01	n=1
Fe-S_biosyn	p=1.3E-01	n=1
Fer4_24	p=1.3E-01	n=1
Gelsolin	p=1.3E-01	n=2
GOLD_2	p=1.3E-01	n=1
HIG_1_N	p=1.3E-01	n=1
Hint	p=1.3E-01	n=1
hnRNP_Q_AcD	p=1.3E-01	n=1
Homeodomain	p=1.3E-01	n=4
IMD	p=1.3E-01	n=1
Ion_trans_2	p=1.3E-01	n=5
Laminin_G_2	p=1.3E-01	n=2
NRBF2_MIT	p=1.3E-01	n=1
ORMDL	p=1.3E-01	n=1
Peptidase_M2	p=1.3E-01	n=1

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

 $-\log_{10}(p)$

n=209/198 input genes with annotations



fraction