

Ncol_Nvec_vc1.1_XM_032374041.2

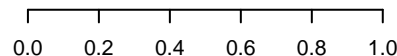
fraction genes in fg and bg

Band_7	p=1.0E-01	n=2
BTK	p=1.0E-01	n=1
bZIP_2	p=1.0E-01	n=2
CAP	p=1.0E-01	n=2
CDK5_activator	p=1.0E-01	n=1
Cobalamin_bind	p=1.0E-01	n=1
COLFI	p=1.0E-01	n=1
DUF3399	p=1.0E-01	n=1
DUF4430	p=1.0E-01	n=1
ELM2	p=1.0E-01	n=2
EMP70	p=1.0E-01	n=1
ETC_C1_NDUFA4	p=1.0E-01	n=1
Gelsolin	p=1.0E-01	n=2
HIF-1a_CTAD	p=1.0E-01	n=1
HTH_3	p=1.0E-01	n=1
MBF1	p=1.0E-01	n=1
NERD	p=1.0E-01	n=1
NHR2	p=1.0E-01	n=1
NPC1_N	p=1.0E-01	n=1
Paf67	p=1.0E-01	n=1
PH_17	p=1.0E-01	n=1
PLA2_inh	p=1.0E-01	n=1
PMT	p=1.0E-01	n=1
PMT_4TMC	p=1.0E-01	n=1
RNA_pol_Rpb4	p=1.0E-01	n=1
RNase_Zc3h12a	p=1.0E-01	n=1
ROQ_II	p=1.0E-01	n=1
Rpn3_C	p=1.0E-01	n=1
Shal-type	p=1.0E-01	n=1
SNF	p=1.0E-01	n=2

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
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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

 $-\log_{10}(p)$

n=173/155 input genes with annotations



fraction