

Elav_Nvec_vc1.1_XM_032381020.2

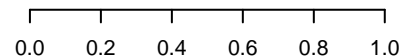
fraction genes in fg and bg

AAA_5	p=1.1E-01	n=1
AHD	p=1.1E-01	n=1
AhpC-TSA_2	p=1.1E-01	n=1
ANAPC5	p=1.1E-01	n=1
AOX	p=1.1E-01	n=1
Arm_vescicular	p=1.1E-01	n=1
ASD2	p=1.1E-01	n=1
ATP_Ca_trans_C	p=1.1E-01	n=1
ATP-synt_ab_Xtn	p=1.1E-01	n=1
Band_7	p=1.1E-01	n=2
Bcl-2	p=1.1E-01	n=2
Bin3	p=1.1E-01	n=1
BTB_3	p=1.1E-01	n=1
C1_1	p=1.1E-01	n=4
C1_2	p=1.1E-01	n=1
CBF_beta	p=1.1E-01	n=1
CBFB_NFYA	p=1.1E-01	n=1
CH	p=1.1E-01	n=4
DENN	p=1.1E-01	n=2
DNA_ligase_A_C	p=1.1E-01	n=1
DNA_ligase_A_M	p=1.1E-01	n=1
DNA_ligase_A_N	p=1.1E-01	n=1
DUF1356	p=1.1E-01	n=1
DUF4502	p=1.1E-01	n=1
DUF4503	p=1.1E-01	n=1
Dynein_IC2	p=1.1E-01	n=1
ELL	p=1.1E-01	n=1
EloA-BP1	p=1.1E-01	n=1
FAM117	p=1.1E-01	n=1
FCH	p=1.1E-01	n=2

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=0.00	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=263/227 input genes with annotations



fraction