

Elav_Nvec_vc1.1_XM_032377390.2

fraction genes in fg and bg

ADIP	p=1.3E-01	n=1
AP1AR	p=1.3E-01	n=1
Arm_2	p=1.3E-01	n=1
ASF1_hist_chap	p=1.3E-01	n=1
ATP_Ca_trans_C	p=1.3E-01	n=1
ATP-cone	p=1.3E-01	n=1
BCCT	p=1.3E-01	n=1
C2	p=1.3E-01	n=6
C2-set_2	p=1.3E-01	n=1
CBFB_NFYA	p=1.3E-01	n=1
CDH1_2_SANT_HL1	p=1.3E-01	n=1
Complex1_49kDa	p=1.3E-01	n=1
COQ7	p=1.3E-01	n=1
DAZAP2	p=1.3E-01	n=1
DBP10CT	p=1.3E-01	n=1
DHR10	p=1.3E-01	n=1
DNA_primase_lrg	p=1.3E-01	n=1
DUF2040	p=1.3E-01	n=1
DUF3399	p=1.3E-01	n=1
DUF3454	p=1.3E-01	n=1
DUF3668	p=1.3E-01	n=1
DUF4149	p=1.3E-01	n=1
DUF4208	p=1.3E-01	n=1
DUF726	p=1.3E-01	n=1
DUF89	p=1.3E-01	n=1
E3_UbLigase_R4	p=1.3E-01	n=1
EF-hand_5	p=1.3E-01	n=4
EF-hand_9	p=1.3E-01	n=1
F-box-like	p=1.3E-01	n=5
Fam20C	p=1.3E-01	n=1

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
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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=0.00	bg=0.00



$-\log_{10}(p)$
n=336/319 input genes with annotations



fraction