

Elav\_Nvec\_vc1.1\_XM\_001633398.3

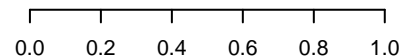
fraction genes in fg and bg

40S_S4_C	p=1.1E-01	n=1
ABC_transp_aux	p=1.1E-01	n=1
ANAPC8	p=1.1E-01	n=1
Armet	p=1.1E-01	n=1
BAAT_C	p=1.1E-01	n=1
BACK	p=1.1E-01	n=4
BAT2_N	p=1.1E-01	n=1
Bile_Hydr_Trans	p=1.1E-01	n=1
Clat_adaptor_s	p=1.1E-01	n=2
Cnd3	p=1.1E-01	n=1
COX5B	p=1.1E-01	n=1
Csm1	p=1.1E-01	n=1
DNA_pol3_delta2	p=1.1E-01	n=1
DUF1682	p=1.1E-01	n=1
DUF2205	p=1.1E-01	n=1
DUF2462	p=1.1E-01	n=1
DUF3337	p=1.1E-01	n=1
DUF3754	p=1.1E-01	n=1
DUF4217	p=1.1E-01	n=2
DUF4497	p=1.1E-01	n=1
DUF4517	p=1.1E-01	n=1
Dynamitin	p=1.1E-01	n=1
EHN	p=1.1E-01	n=1
Fanconi_C	p=1.1E-01	n=1
FPL	p=1.1E-01	n=1
FragX_IP	p=1.1E-01	n=1
GAF	p=1.1E-01	n=2
GARS_A	p=1.1E-01	n=1
GARS_C	p=1.1E-01	n=1
GARS_N	p=1.1E-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=0.01	bg=0.00
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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



$-\log_{10}(p)$   
n=308/289 input genes with annotations



fraction