

## Elav\_Nvec\_vc1.1\_XM\_048733866.1

## fraction genes in fg and bg

5-nucleotidase	p=1.1E-01	n=1
AA_permease_C	p=1.1E-01	n=2
AAA_19	p=1.1E-01	n=1
AlbA_2	p=1.1E-01	n=1
ANAPC5	p=1.1E-01	n=1
AP3B1_C	p=1.1E-01	n=1
APCDDC	p=1.1E-01	n=1
Auts2	p=1.1E-01	n=1
AWS	p=1.1E-01	n=1
betaPIX_CC	p=1.1E-01	n=1
Bravo_FIGEY	p=1.1E-01	n=1
BTK	p=1.1E-01	n=1
C2	p=1.1E-01	n=6
C5HCH	p=1.1E-01	n=1
CiPC	p=1.1E-01	n=1
Cluap1	p=1.1E-01	n=1
cNMP_binding	p=1.1E-01	n=3
Cob_adeno_trans	p=1.1E-01	n=1
CortBP2	p=1.1E-01	n=1
CPSF_A	p=1.1E-01	n=1
DOR	p=1.1E-01	n=1
DUF2046	p=1.1E-01	n=1
DUF3689	p=1.1E-01	n=1
DUF4419	p=1.1E-01	n=1
DUF543	p=1.1E-01	n=1
EAF	p=1.1E-01	n=1
EB1	p=1.1E-01	n=1
EF-hand_4	p=1.1E-01	n=2
eIF-3c_N	p=1.1E-01	n=1
Es2	p=1.1E-01	n=1

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
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=0.02	bg=0.01
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
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



$-\log_{10}(p)$   
n=275/230 input genes with annotations



fraction