

Elav_Nvec_vc1.1_XM_001640781.3

fraction genes in fg and bg

CSRNP_N	p=1.1E-01	n=1
Dishevelled	p=1.1E-01	n=1
DIX	p=1.1E-01	n=1
Dsh_C	p=1.1E-01	n=1
DUF758	p=1.1E-01	n=1
I-set	p=1.1E-01	n=2
Insulin	p=1.1E-01	n=1
NHR2	p=1.1E-01	n=1
Nucleic_acid_bd	p=1.1E-01	n=1
PEHE	p=1.1E-01	n=1
Pkinase_Tyr	p=1.1E-01	n=3
RAWUL	p=1.1E-01	n=1
Runt	p=1.1E-01	n=1
SAND	p=1.1E-01	n=1
START	p=1.1E-01	n=1
TAFH	p=1.1E-01	n=1
VIT_2	p=1.1E-01	n=1
VWA_3	p=1.1E-01	n=1
zf-C2H2_6	p=1.1E-01	n=1
SAM_1	p=1.1E-01	n=2
Cation_ATPase_C	p=1.2E-01	n=1
Cation_ATPase_N	p=1.2E-01	n=1
DED	p=1.2E-01	n=1
DUF3504	p=1.2E-01	n=1
EF-hand_1	p=1.2E-01	n=1
ELM2	p=1.2E-01	n=1
EphA2_TM	p=1.2E-01	n=1
Ephrin_rec_like	p=1.2E-01	n=1
FAM181	p=1.2E-01	n=1
Frag1	p=1.2E-01	n=1

fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.03	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.04	bg=0.01
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.03	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00

 $-\log_{10}(p)$

n=70/56 input genes with annotations



fraction