

Elav_Nvec_vc1.1_XM_048719424.1

fraction genes in fg and bg

G-alpha	p=4.4E-03	n=4
3Beta_HSD	p=1.3E-01	n=1
AAA_19	p=1.3E-01	n=1
Actin	p=1.3E-01	n=2
AhpC-TSA_2	p=1.3E-01	n=1
ANAPC5	p=1.3E-01	n=1
Ank	p=1.3E-01	n=3
Arm_vescicular	p=1.3E-01	n=1
ATP-synt_ab_Xtn	p=1.3E-01	n=1
ATP-synt_F	p=1.3E-01	n=1
Auts2	p=1.3E-01	n=1
BTB_3	p=1.3E-01	n=1
C1_2	p=1.3E-01	n=1
CDH1_2_SANT_HL1	p=1.3E-01	n=1
CG-1	p=1.3E-01	n=1
Chorein_N	p=1.3E-01	n=2
CRIM	p=1.3E-01	n=1
DAG_kinase_N	p=1.3E-01	n=1
DED	p=1.3E-01	n=2
DNA_ligase_A_C	p=1.3E-01	n=1
DNA_ligase_A_M	p=1.3E-01	n=1
DNA_ligase_A_N	p=1.3E-01	n=1
DUF1977	p=1.3E-01	n=1
DUF2370	p=1.3E-01	n=1
DUF4208	p=1.3E-01	n=1
DUF4419	p=1.3E-01	n=1
DUF953	p=1.3E-01	n=1
Dynein_IC2	p=1.3E-01	n=1
EphA2_TM	p=1.3E-01	n=2
Ephrin_Ibd	p=1.3E-01	n=2

fg=0.01	bg=0.00
fg=0.00	bg=0.00
fg=0.00	bg=0.00
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fg=0.01	bg=0.00
fg=0.01	bg=0.00

-log₁₀(p)

n=296/246 input genes with annotations



fraction