

Elav_Nvec_vc1.1_XM_001631673.3

fraction genes in fg and bg

ATP-synt_ab_Xtn	p=7.7E-02	n=1
Atthog	p=7.7E-02	n=1
BIRC6	p=7.7E-02	n=1
BolA	p=7.7E-02	n=1
CASP_C	p=7.7E-02	n=1
COQ7	p=7.7E-02	n=1
Cu_amine_oxid	p=7.7E-02	n=1
Cu_amine_oxidN2	p=7.7E-02	n=1
DALR_1	p=7.7E-02	n=1
DFP	p=7.7E-02	n=1
DUF21	p=7.7E-02	n=1
DUF2370	p=7.7E-02	n=1
DUF3454	p=7.7E-02	n=1
DUF4495	p=7.7E-02	n=1
DUF4611	p=7.7E-02	n=1
DUF726	p=7.7E-02	n=1
DUF758	p=7.7E-02	n=1
E3_UbLigase_R4	p=7.7E-02	n=1
EF-hand_5	p=7.7E-02	n=3
ELM2	p=7.7E-02	n=2
FAM53	p=7.7E-02	n=1
FDF	p=7.7E-02	n=1
Fer4_24	p=7.7E-02	n=1
FHA_2	p=7.7E-02	n=1
Fra10Ac1	p=7.7E-02	n=1
HAND	p=7.7E-02	n=1
HECW_N	p=7.7E-02	n=1
HECW1_helix	p=7.7E-02	n=1
HLH	p=7.7E-02	n=3
KIND	p=7.7E-02	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
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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=0.01	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=0.01	bg=0.00
fg=0.00	bg=0.00

0 1 2 3 4 5

0.0 0.2 0.4 0.6 0.8 1.0

-log₁₀(p)

fraction

n=182/153 input genes with annotations