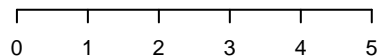


Elav_Nvec_vc1.1_XM_001639164.3

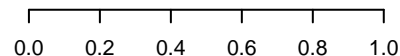
fraction genes in fg and bg

ELM2	p=4.2E-02	n=2
ATP-synt_ab	p=7.2E-02	n=1
ATP-synt_ab_N	p=7.2E-02	n=1
ATP-synt_ab_Xtn	p=7.2E-02	n=1
Cu_amine_oxid	p=7.2E-02	n=1
Cu_amine_oxidN2	p=7.2E-02	n=1
DDHD	p=7.2E-02	n=1
DUF2370	p=7.2E-02	n=1
DUF4495	p=7.2E-02	n=1
DUF758	p=7.2E-02	n=1
DUSP	p=7.2E-02	n=1
EF-hand_5	p=7.2E-02	n=2
FAM53	p=7.2E-02	n=1
Fer4_24	p=7.2E-02	n=1
Fra10Ac1	p=7.2E-02	n=1
KIND	p=7.2E-02	n=1
NT-C2	p=7.2E-02	n=1
PALP	p=7.2E-02	n=1
Pkinase	p=7.2E-02	n=4
PPP4R2	p=7.2E-02	n=1
PTPlike_phytase	p=7.2E-02	n=1
zf-C2H2_6	p=7.2E-02	n=1
adh_short	p=1.0E-01	n=1
ARID	p=1.0E-01	n=1
BAH	p=1.0E-01	n=1
DUF3585	p=1.0E-01	n=1
GATA	p=1.0E-01	n=1
HLH	p=1.0E-01	n=2
IP_trans	p=1.0E-01	n=1
TIR_2	p=1.0E-01	n=1

-log₁₀(p)

n=59/52 input genes with annotations

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
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=0.06	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



fraction