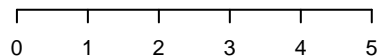


Elav_Nvec_vc1.1_XM_032383641.2

fraction genes in fg and bg

2OG-Fell_Oxy_3	p=4.3E-02	n=2
AA_permease_2	p=4.3E-02	n=2
AA_permease_C	p=4.3E-02	n=2
Cbl_N	p=4.3E-02	n=1
Cbl_N2	p=4.3E-02	n=1
Cbl_N3	p=4.3E-02	n=1
Hid1	p=4.3E-02	n=1
HJURP_C	p=4.3E-02	n=1
PDE8	p=4.3E-02	n=1
PH_RBD	p=4.3E-02	n=1
Prok-RING_4	p=4.3E-02	n=1
RHD_dimer	p=4.3E-02	n=1
RHD_DNA_bind	p=4.3E-02	n=1
RT_RNaseH	p=4.3E-02	n=1
SAND	p=4.3E-02	n=1
SRF-TF	p=4.3E-02	n=1
WIF	p=4.3E-02	n=1
Actin	p=6.7E-02	n=1
GATA	p=6.7E-02	n=1
Integrase_H2C2	p=6.7E-02	n=1
PBD	p=6.7E-02	n=1
Pkinase_Tyr	p=6.7E-02	n=2
AC_N	p=8.9E-02	n=1
RUN	p=8.9E-02	n=1
Abhydrolase_1	p=1.1E-01	n=1
PAS_9	p=1.1E-01	n=1
EF-hand_5	p=1.2E-01	n=1
MFS_1_like	p=1.2E-01	n=1
PDEase_I	p=1.2E-01	n=1
Pkinase	p=1.2E-01	n=2



-log₁₀(p)
n=40/38 input genes with annotations

fg=0.04	bg=0.00
fg=0.04	bg=0.00
fg=0.04	bg=0.00
fg=0.02	bg=0.00
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fg=0.02	bg=0.00
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fg=0.04	bg=0.01
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.04	bg=0.01



fraction