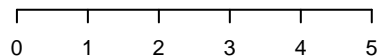


Elav_Nvec_vc1.1_XM_032367573.2

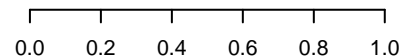
fraction genes in fg and bg

7tm_1	p=8.5E-02	n=14
Arginosuc_synth	p=8.5E-02	n=1
ASCH	p=8.5E-02	n=1
ATP_Ca_trans_C	p=8.5E-02	n=1
B3_4	p=8.5E-02	n=1
B5	p=8.5E-02	n=1
Cation_ATPase_C	p=8.5E-02	n=2
Cation_ATPase_N	p=8.5E-02	n=2
CCCAP	p=8.5E-02	n=1
CDK5_activator	p=8.5E-02	n=1
CFAP298	p=8.5E-02	n=1
CLN3	p=8.5E-02	n=3
Complex1_51K	p=8.5E-02	n=1
COQ7	p=8.5E-02	n=1
DAG_kinase_N	p=8.5E-02	n=1
DUF1736	p=8.5E-02	n=2
DUF21	p=8.5E-02	n=1
DUF2181	p=8.5E-02	n=1
DUF2340	p=8.5E-02	n=1
DUF4686	p=8.5E-02	n=1
DUF4745	p=8.5E-02	n=1
DUF547	p=8.5E-02	n=1
eIF3g	p=8.5E-02	n=1
ETS_PEA3_N	p=8.5E-02	n=1
FAM117	p=8.5E-02	n=1
Fascin	p=8.5E-02	n=1
FYVE	p=8.5E-02	n=3
Glyco_trans_1_4	p=8.5E-02	n=1
HR1	p=8.5E-02	n=1
INTS2	p=8.5E-02	n=1



-log₁₀(p)
n=263/237 input genes with annotations

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



fraction