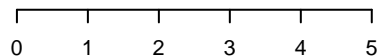


Elav_Nvec_vc1.1_XM_001632368.2

fraction genes in fg and bg

ACC_central	p=6.6E-02	n=1
ADP_ribosyl_GH	p=6.6E-02	n=2
APCDDC	p=6.6E-02	n=1
Auts2	p=6.6E-02	n=1
BBIP10	p=6.6E-02	n=1
CUPID	p=6.6E-02	n=1
DMAP1	p=6.6E-02	n=1
DUF2205	p=6.6E-02	n=1
EHN	p=6.6E-02	n=1
F-box	p=6.6E-02	n=2
FAIM1	p=6.6E-02	n=1
FAM117	p=6.6E-02	n=1
GFA	p=6.6E-02	n=1
Glyco_transf_24	p=6.6E-02	n=1
Glyco_transf_29	p=6.6E-02	n=1
Glycos_transf_1	p=6.6E-02	n=2
Kazal_2	p=6.6E-02	n=2
MARVEL	p=6.6E-02	n=2
Mgm101p	p=6.6E-02	n=1
Microtub_bind	p=6.6E-02	n=1
NOB1_Zn_bind	p=6.6E-02	n=1
OSCP	p=6.6E-02	n=1
p53-inducible11	p=6.6E-02	n=1
PAP2_3	p=6.6E-02	n=1
Pet191_N	p=6.6E-02	n=1
PIN_6	p=6.6E-02	n=1
Ribosomal_L31e	p=6.6E-02	n=1
Ribosomal_L37ae	p=6.6E-02	n=1
Ribosomal_L50	p=6.6E-02	n=1
Ribosomal_S27	p=6.6E-02	n=1

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



-log₁₀(p)
n=181/145 input genes with annotations



fraction