

Elav_Nvec_vc1.1_XM_001626717.3

fraction genes in fg and bg

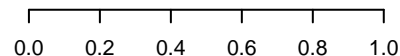
ADH_N	p=1.4E-01	n=1
ADH_zinc_N	p=1.4E-01	n=1
BK_channel_a	p=1.4E-01	n=1
CK_II_beta	p=1.4E-01	n=1
DUF1154	p=1.4E-01	n=1
DUF3350	p=1.4E-01	n=1
ELL	p=1.4E-01	n=1
ELM2	p=1.4E-01	n=2
Hid1	p=1.4E-01	n=1
HTH_Tnp_Tc5	p=1.4E-01	n=1
Ig_3	p=1.4E-01	n=2
IMD	p=1.4E-01	n=1
Kringle	p=1.4E-01	n=1
Occludin_ELL	p=1.4E-01	n=1
PALP	p=1.4E-01	n=1
PH_14	p=1.4E-01	n=1
PI-PLC-X	p=1.4E-01	n=2
PI-PLC-Y	p=1.4E-01	n=2
PID	p=1.4E-01	n=2
PTPlake_phytase	p=1.4E-01	n=1
RabGAP-TBC	p=1.4E-01	n=2
SCAPER_N	p=1.4E-01	n=1
SIX1_SD	p=1.4E-01	n=1
Ski_Sno	p=1.4E-01	n=1
START	p=1.4E-01	n=1
VIT_2	p=1.4E-01	n=1
VWA_3	p=1.4E-01	n=1
zf-C2H2_6	p=1.4E-01	n=1
zf-U11-48K	p=1.4E-01	n=1
Homeodomain	p=1.6E-01	n=3

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



-log10(p)

n=93/74 input genes with annotations



fraction