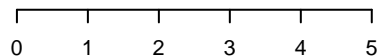


Ncol_Nvec_vc1.1_XM_032380374.2

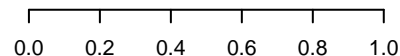
fraction genes in fg and bg

7tm_2	p=2.0E-01	n=3
Acyltransferase	p=2.0E-01	n=1
ADAM_CR_2	p=2.0E-01	n=1
BK_channel_a	p=2.0E-01	n=1
BTB	p=2.0E-01	n=4
BTB_2	p=2.0E-01	n=5
bZIP_1	p=2.0E-01	n=2
CDK5_activator	p=2.0E-01	n=1
Cu_amine_oxid	p=2.0E-01	n=1
Cu_amine_oxidN2	p=2.0E-01	n=1
Cu2_monoox_C	p=2.0E-01	n=1
Cu2_monooxygen	p=2.0E-01	n=1
DUF3399	p=2.0E-01	n=1
EF-hand_7	p=2.0E-01	n=5
EF-hand_like	p=2.0E-01	n=2
Glyco_transf_92	p=2.0E-01	n=1
HJURP_C	p=2.0E-01	n=1
HlyIII	p=2.0E-01	n=2
Ion_trans_2	p=2.0E-01	n=7
IQ	p=2.0E-01	n=1
Jun	p=2.0E-01	n=1
Laminin_G_3	p=2.0E-01	n=4
Leg1	p=2.0E-01	n=1
Leu_zip	p=2.0E-01	n=1
LMF1	p=2.0E-01	n=1
Mab-21	p=2.0E-01	n=1
MFS_1	p=2.0E-01	n=4
Myosin_head	p=2.0E-01	n=1
Neur_chan_LBD	p=2.0E-01	n=5
Neur_chan_memb	p=2.0E-01	n=5

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=0.02	bg=0.01
fg=0.03	bg=0.01
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=0.03	bg=0.01
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.04	bg=0.01
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.02	bg=0.01
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.02	bg=0.01
fg=0.01	bg=0.00
fg=0.03	bg=0.01
fg=0.03	bg=0.01

 $-\log_{10}(p)$

n=129/144 input genes with annotations



fraction