

Ncol_Nvec_vc1.1_XM_001639692.3

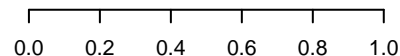
fraction genes in fg and bg

BTB_2	p=3.0E-03	n=9
Neur_chan_LBD	p=3.0E-03	n=9
Neur_chan_memb	p=3.0E-03	n=9
Oxysterol_BP	p=8.3E-02	n=3
Ion_trans_2	p=1.2E-01	n=7
P4Ha_N	p=1.2E-01	n=2
Abhydrolase_3	p=1.8E-01	n=2
ADAM_CR_2	p=1.8E-01	n=1
AIRS	p=1.8E-01	n=1
ANF_receptor	p=1.8E-01	n=4
Arm_vesicular	p=1.8E-01	n=1
Cadherin	p=1.8E-01	n=3
CaM_bdg_C0	p=1.8E-01	n=1
CDK5_activator	p=1.8E-01	n=1
CEBP_ZZ	p=1.8E-01	n=1
Ceramidase	p=1.8E-01	n=1
CG-1	p=1.8E-01	n=1
Chromo	p=1.8E-01	n=2
Cu_amine_oxid	p=1.8E-01	n=1
Cu_amine_oxidN2	p=1.8E-01	n=1
Dickkopf_N	p=1.8E-01	n=1
DUF1977	p=1.8E-01	n=1
DUF3399	p=1.8E-01	n=1
DUF3583	p=1.8E-01	n=1
EF-hand_7	p=1.8E-01	n=6
EF-hand_like	p=1.8E-01	n=2
EMI	p=1.8E-01	n=1
ERO1	p=1.8E-01	n=1
Exo_endo_phos_2	p=1.8E-01	n=1
Fes1	p=1.8E-01	n=1

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

 $-\log_{10}(p)$

n=292/356 input genes with annotations



fraction