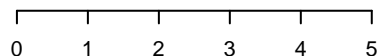


Ncol_Nvec_vc1.1_XM_032386356.2

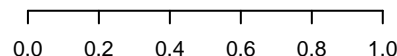
fraction genes in fg and bg

Neur_chan_LBD	p=1.2E-02	n=7
Neur_chan_memb	p=1.2E-02	n=7
BTB_2	p=6.2E-02	n=6
FA_desaturase	p=6.3E-02	n=2
2Fe-2S_thioredx	p=1.2E-01	n=1
Acyl_transf_3	p=1.2E-01	n=1
Acyltransferase	p=1.2E-01	n=2
ANAPC5	p=1.2E-01	n=1
Asparaginase_2	p=1.2E-01	n=2
ATP_synt_H	p=1.2E-01	n=1
ATP_synt_F	p=1.2E-01	n=1
CDK5_activator	p=1.2E-01	n=1
Cobalamin_bind	p=1.2E-01	n=1
Cu_amine_oxid	p=1.2E-01	n=1
Cu_amine_oxidN2	p=1.2E-01	n=1
DUF1977	p=1.2E-01	n=1
DUF2040	p=1.2E-01	n=1
DUF3543	p=1.2E-01	n=1
DUF3808	p=1.2E-01	n=1
DUF3987	p=1.2E-01	n=1
DUF4430	p=1.2E-01	n=1
DUF773	p=1.2E-01	n=1
EF-hand_7	p=1.2E-01	n=5
EGF	p=1.2E-01	n=5
Glyco_tran_10_N	p=1.2E-01	n=2
Glyco_transf_10	p=1.2E-01	n=2
GRASP55_65	p=1.2E-01	n=1
HLH	p=1.2E-01	n=4
HMG_box	p=1.2E-01	n=3
Importin_rep	p=1.2E-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=0.00	bg=0.00
fg=0.00	bg=0.00
fg=0.01	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
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=0.02	bg=0.01
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=0.00	bg=0.00

-log₁₀(p)

n=224/227 input genes with annotations



fraction