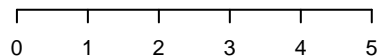


Ncol_Nvec_vc1.1_XM_032364814.2

fraction genes in fg and bg

Neur_chan_LBD	p=4.5E-03	n=10
Neur_chan_memb	p=4.5E-03	n=10
2OG-Fell_Oxy_3	p=2.4E-01	n=2
AA_kinase	p=2.4E-01	n=1
Aa_trans	p=2.4E-01	n=1
Acyltransferase	p=2.4E-01	n=1
ADAM_CR_2	p=2.4E-01	n=1
ApeC	p=2.4E-01	n=1
BACK	p=2.4E-01	n=4
BK_channel_a	p=2.4E-01	n=1
BTB	p=2.4E-01	n=5
Cadherin	p=2.4E-01	n=2
CDK5_activator	p=2.4E-01	n=1
Ceramidase	p=2.4E-01	n=1
Cu_amine_oxid	p=2.4E-01	n=1
Cu_amine_oxidN2	p=2.4E-01	n=1
Cu2_monoox_C	p=2.4E-01	n=1
Cu2_monooxygen	p=2.4E-01	n=1
Dickkopf_N	p=2.4E-01	n=1
DUF3399	p=2.4E-01	n=1
EF-hand_like	p=2.4E-01	n=2
Fes1	p=2.4E-01	n=1
Filamin	p=2.4E-01	n=3
FUN14	p=2.4E-01	n=1
Glyco_transf_92	p=2.4E-01	n=1
Hormone_recep	p=2.4E-01	n=2
lon_trans_2	p=2.4E-01	n=7
IQ	p=2.4E-01	n=1
Leg1	p=2.4E-01	n=1
Leu_zip	p=2.4E-01	n=1



-log10(p)

n=159/195 input genes with annotations

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



fraction