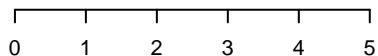


Ncol_Nvec_vc1.1_XM_001627423.3

fraction genes in fg and bg

BTB_2	p=1.9E-03	n=9
lon_trans_2	p=2.0E-03	n=9
lon_trans	p=1.4E-02	n=10
COX6C	p=1.1E-01	n=2
7tm_2	p=1.3E-01	n=3
ADAM_CR_2	p=1.3E-01	n=1
AIRS	p=1.3E-01	n=1
APP_amyloid	p=1.3E-01	n=1
APP_Cu_bd	p=1.3E-01	n=1
APP_E2	p=1.3E-01	n=1
APP_N	p=1.3E-01	n=1
ATP_synt_H	p=1.3E-01	n=1
BACK	p=1.3E-01	n=4
BTB	p=1.3E-01	n=6
CaM_bdg_C0	p=1.3E-01	n=1
CDK5_activator	p=1.3E-01	n=1
CEBP_ZZ	p=1.3E-01	n=1
cNMP_binding	p=1.3E-01	n=3
Coatomeer_WDAD	p=1.3E-01	n=1
Cu_amine_oxid	p=1.3E-01	n=1
Cu_amine_oxidN2	p=1.3E-01	n=1
DUF1180	p=1.3E-01	n=1
DUF3399	p=1.3E-01	n=1
DUF3583	p=1.3E-01	n=1
DUF719	p=1.3E-01	n=1
EAF	p=1.3E-01	n=1
ECH_2	p=1.3E-01	n=1
EF-hand_7	p=1.3E-01	n=6
EF-hand_like	p=1.3E-01	n=2
ELL	p=1.3E-01	n=1

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

-log₁₀(p)

n=254/278 input genes with annotations



fraction