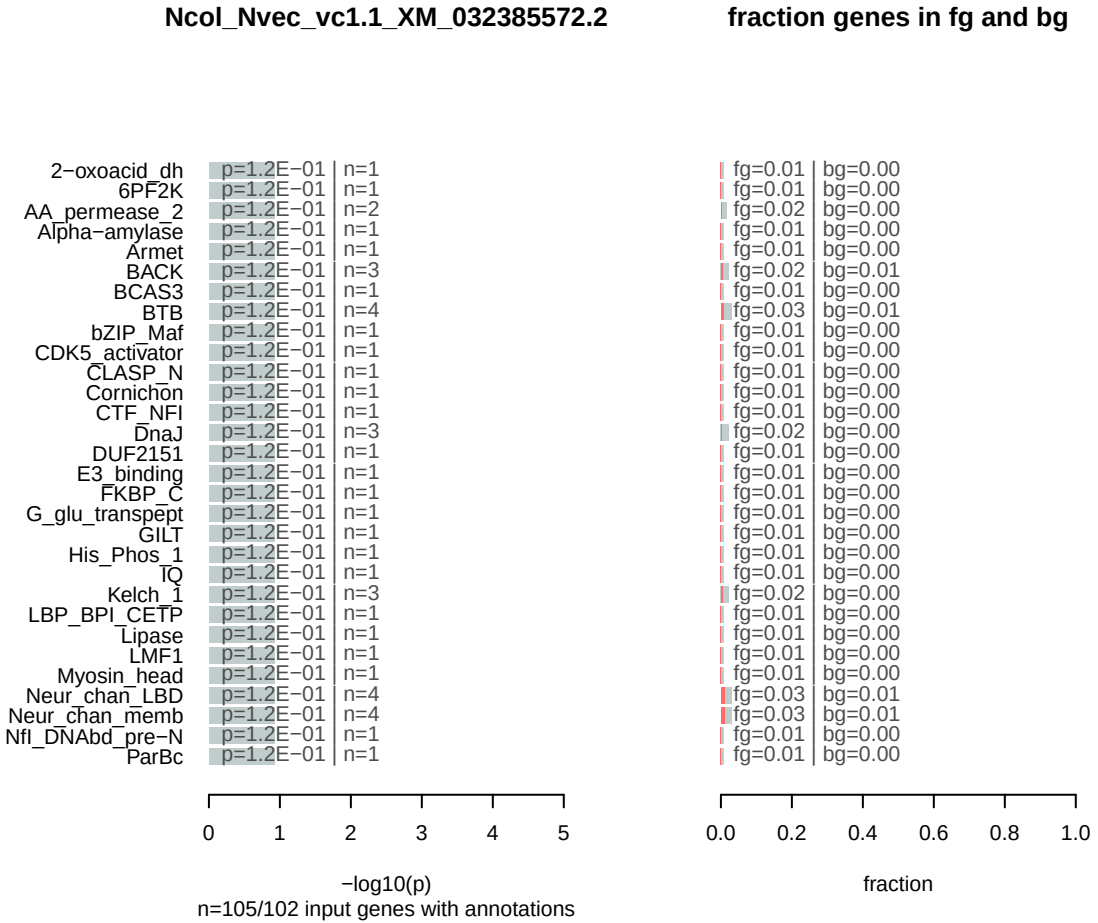
Ncol_Nvec_vc1.1_XM_032385572.2
fraction genes in fg and bg
2−oxoacid_dh p=1.2E−01 n=1 fg=0.01 bg=0.00
6PF2K p=1.2E−01 n=1 fg=0.01 bg=0.00
AA_permease_2 p=1.2E−01 n=2 fg=0.02 bg=0.00
Alpha−amylase p=1.2E−01 n=1 fg=0.01 bg=0.00
Armet p=1.2E−01 n=1 fg=0.01 bg=0.00
BACK p=1.2E−01 n=3 fg=0.02 bg=0.01
BCAS3 p=1.2E−01 n=1 fg=0.01 bg=0.00
BTB p=1.2E−01 n=4 fg=0.03 bg=0.01
bZIP_Maf p=1.2E−01 n=1 fg=0.01 bg=0.00
CDK5_activator p=1.2E−01 n=1 fg=0.01 bg=0.00
CLASP_N p=1.2E−01 n=1 fg=0.01 bg=0.00
Cornichon p=1.2E−01 n=1 fg=0.01 bg=0.00
CTF_NFI p=1.2E−01 n=1 fg=0.01 bg=0.00
DnaJ p=1.2E−01 n=3 fg=0.02 bg=0.00
DUF2151 p=1.2E−01 n=1 fg=0.01 bg=0.00
E3_binding p=1.2E−01 n=1 fg=0.01 bg=0.00
FKBP_C p=1.2E−01 n=1 fg=0.01 bg=0.00
G_glu_transpept p=1.2E−01 n=1 fg=0.01 bg=0.00
GILT p=1.2E−01 n=1 fg=0.01 bg=0.00
His_Phos_1 p=1.2E−01 n=1 fg=0.01 bg=0.00
IQ p=1.2E−01 n=1 fg=0.01 bg=0.00
Kelch_1 p=1.2E−01 n=3 fg=0.02 bg=0.00
LBP_BPI_CETP p=1.2E−01 n=1 fg=0.01 bg=0.00
Lipase p=1.2E−01 n=1 fg=0.01 bg=0.00
LMF1 p=1.2E−01 n=1 fg=0.01 bg=0.00
Myosin_head p=1.2E−01 n=1 fg=0.01 bg=0.00
Neur_chan_LBD p=1.2E−01 n=4 fg=0.03 bg=0.01
Neur_chan_memb p=1.2E−01 n=4 fg=0.03 bg=0.01
NfI_DNAbd_pre−N p=1.2E−01 n=1 fg=0.01 bg=0.00
ParBc p=1.2E−01 n=1 fg=0.01 bg=0.00
0 1 2 3 4 5
−log10(p)
n=105/102 input genes with annotations
0.0 0.2 0.4 0.6 0.8 1.0
fraction