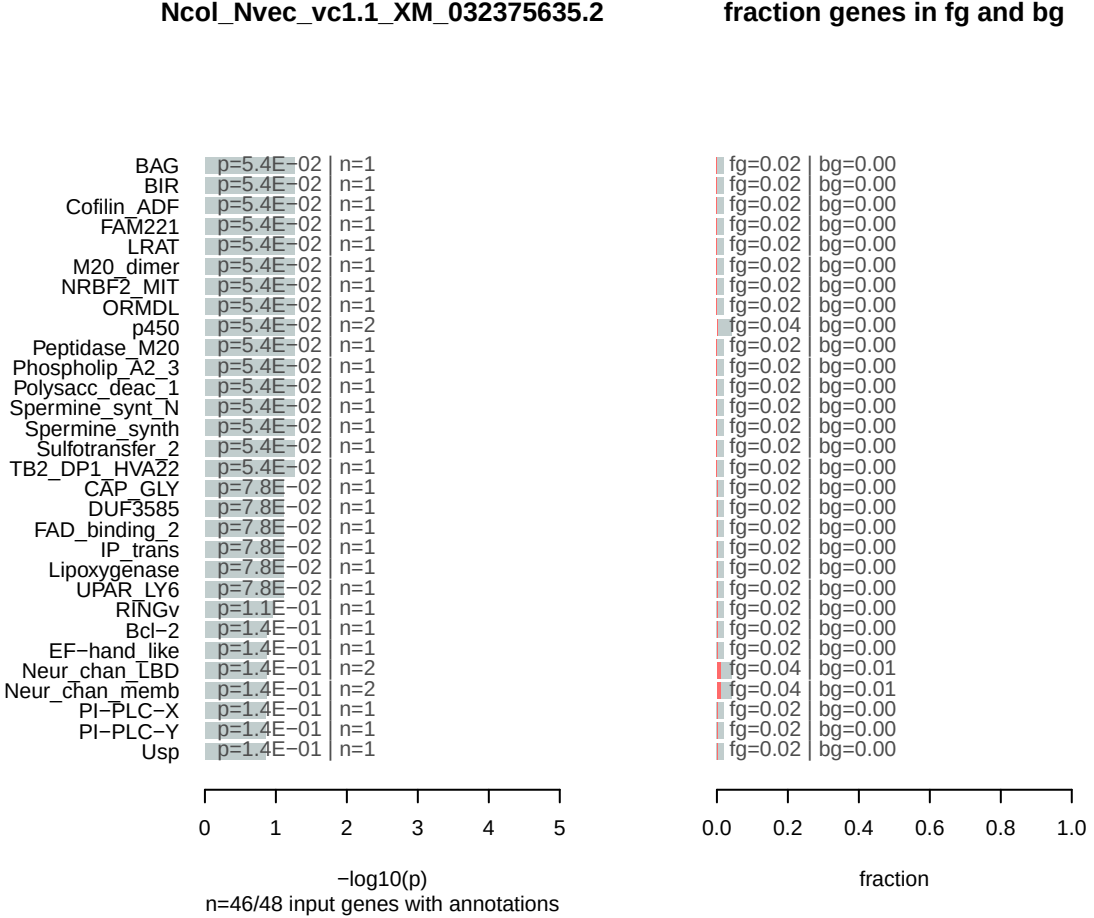
Ncol_Nvec_vc1.1_XM_032375635.2
fraction genes in fg and bg
BAG p=5.4E−02 n=1 fg=0.02 bg=0.00
BIR p=5.4E−02 n=1 fg=0.02 bg=0.00
Cofilin_ADF p=5.4E−02 n=1 fg=0.02 bg=0.00
FAM221 p=5.4E−02 n=1 fg=0.02 bg=0.00
LRAT p=5.4E−02 n=1 fg=0.02 bg=0.00
M20_dimer p=5.4E−02 n=1 fg=0.02 bg=0.00
NRBF2_MIT p=5.4E−02 n=1 fg=0.02 bg=0.00
ORMDL p=5.4E−02 n=1 fg=0.02 bg=0.00
p450 p=5.4E−02 n=2 fg=0.04 bg=0.00
Peptidase_M20 p=5.4E−02 n=1 fg=0.02 bg=0.00
Phospholip_A2_3 p=5.4E−02 n=1 fg=0.02 bg=0.00
Polysacc_deac_1 p=5.4E−02 n=1 fg=0.02 bg=0.00
Spermine_synt_N p=5.4E−02 n=1 fg=0.02 bg=0.00
Spermine_synth p=5.4E−02 n=1 fg=0.02 bg=0.00
Sulfotransfer_2 p=5.4E−02 n=1 fg=0.02 bg=0.00
TB2_DP1_HVA22 p=5.4E−02 n=1 fg=0.02 bg=0.00
CAP_GLY p=7.8E−02 n=1 fg=0.02 bg=0.00
DUF3585 p=7.8E−02 n=1 fg=0.02 bg=0.00
FAD_binding_2 p=7.8E−02 n=1 fg=0.02 bg=0.00
IP_trans p=7.8E−02 n=1 fg=0.02 bg=0.00
Lipoxygenase p=7.8E−02 n=1 fg=0.02 bg=0.00
UPAR_LY6 p=7.8E−02 n=1 fg=0.02 bg=0.00
RINGv p=1.1E−01 n=1 fg=0.02 bg=0.00
Bcl−2 p=1.4E−01 n=1 fg=0.02 bg=0.00
EF−hand_like p=1.4E−01 n=1 fg=0.02 bg=0.00
Neur_chan_LBD p=1.4E−01 n=2 fg=0.04 bg=0.01
Neur_chan_memb p=1.4E−01 n=2 fg=0.04 bg=0.01
PI−PLC−X p=1.4E−01 n=1 fg=0.02 bg=0.00
PI−PLC−Y p=1.4E−01 n=1 fg=0.02 bg=0.00
Usp p=1.4E−01 n=1 fg=0.02 bg=0.00
0 1 2 3 4 5
−log10(p)
0.0 0.2 0.4 0.6 0.8 1.0
fraction
n=46/48 input genes with annotations