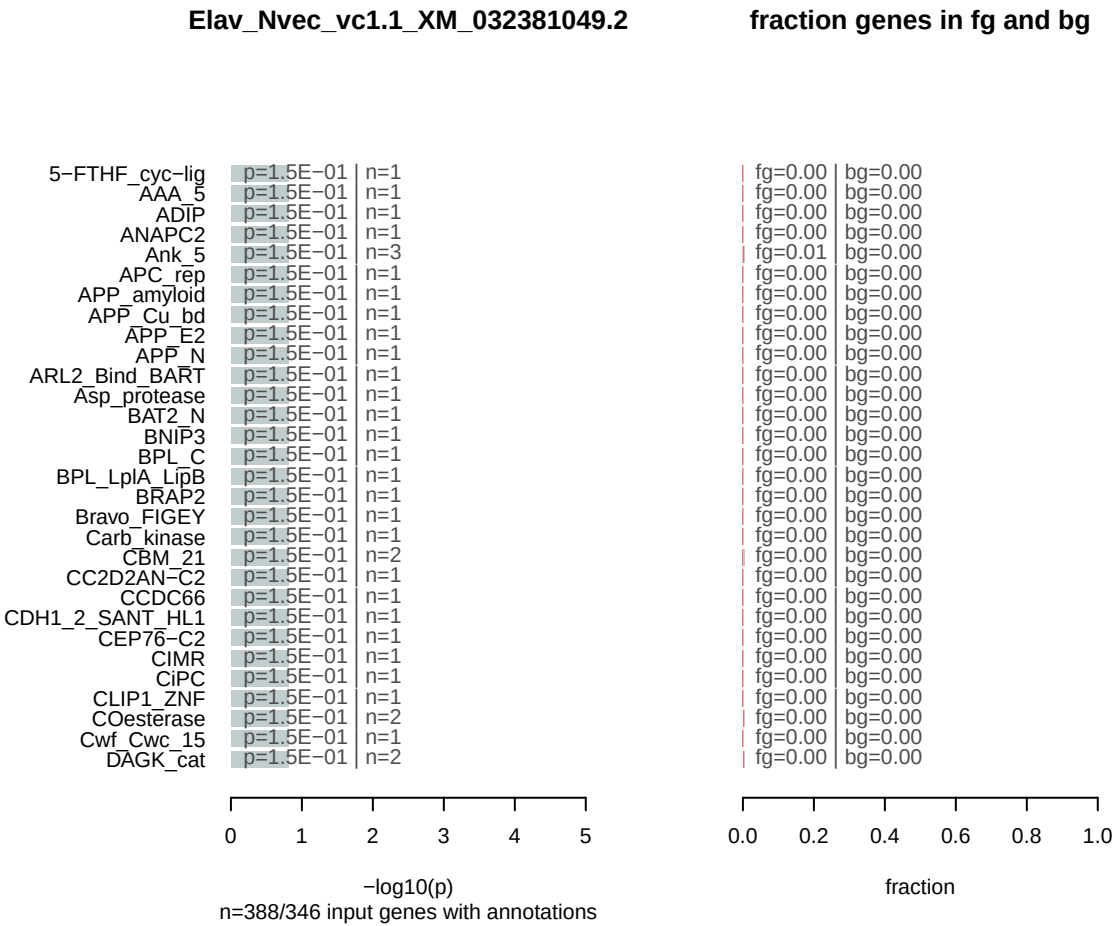
Elav_Nvec_vc1.1_XM_032381049.2
fraction genes in fg and bg
5−FTHF_cyc−lig p=1.5E−01 n=1 fg=0.00 bg=0.00
AAA_5 p=1.5E−01 n=1 fg=0.00 bg=0.00
ADIP p=1.5E−01 n=1 fg=0.00 bg=0.00
ANAPC2 p=1.5E−01 n=1 fg=0.00 bg=0.00
Ank_5 p=1.5E−01 n=3 fg=0.01 bg=0.00
APC_rep p=1.5E−01 n=1 fg=0.00 bg=0.00
APP_amyloid p=1.5E−01 n=1 fg=0.00 bg=0.00
APP_Cu_bd p=1.5E−01 n=1 fg=0.00 bg=0.00
APP_E2 p=1.5E−01 n=1 fg=0.00 bg=0.00
APP_N p=1.5E−01 n=1 fg=0.00 bg=0.00
ARL2_Bind_BART p=1.5E−01 n=1 fg=0.00 bg=0.00
Asp_protease p=1.5E−01 n=1 fg=0.00 bg=0.00
BAT2_N p=1.5E−01 n=1 fg=0.00 bg=0.00
BNIP3 p=1.5E−01 n=1 fg=0.00 bg=0.00
BPL_C p=1.5E−01 n=1 fg=0.00 bg=0.00
BPL_LplA_LipB p=1.5E−01 n=1 fg=0.00 bg=0.00
BRAP2 p=1.5E−01 n=1 fg=0.00 bg=0.00
Bravo_FIGEY p=1.5E−01 n=1 fg=0.00 bg=0.00
Carb_kinase p=1.5E−01 n=1 fg=0.00 bg=0.00
CBM_21 p=1.5E−01 n=2 fg=0.00 bg=0.00
CC2D2AN−C2 p=1.5E−01 n=1 fg=0.00 bg=0.00
CCDC66 p=1.5E−01 n=1 fg=0.00 bg=0.00
CDH1_2_SANT_HL1 p=1.5E−01 n=1 fg=0.00 bg=0.00
CEP76−C2 p=1.5E−01 n=1 fg=0.00 bg=0.00
CIMR p=1.5E−01 n=1 fg=0.00 bg=0.00
CiPC p=1.5E−01 n=1 fg=0.00 bg=0.00
CLIP1_ZNF p=1.5E−01 n=1 fg=0.00 bg=0.00
COesterase p=1.5E−01 n=2 fg=0.00 bg=0.00
Cwf_Cwc_15 p=1.5E−01 n=1 fg=0.00 bg=0.00
DAGK_cat p=1.5E−01 n=2 fg=0.00 bg=0.00
0 1 2 3 4 5
−log10(p)
n=388/346 input genes with annotations
0.0 0.2 0.4 0.6 0.8 1.0
fraction