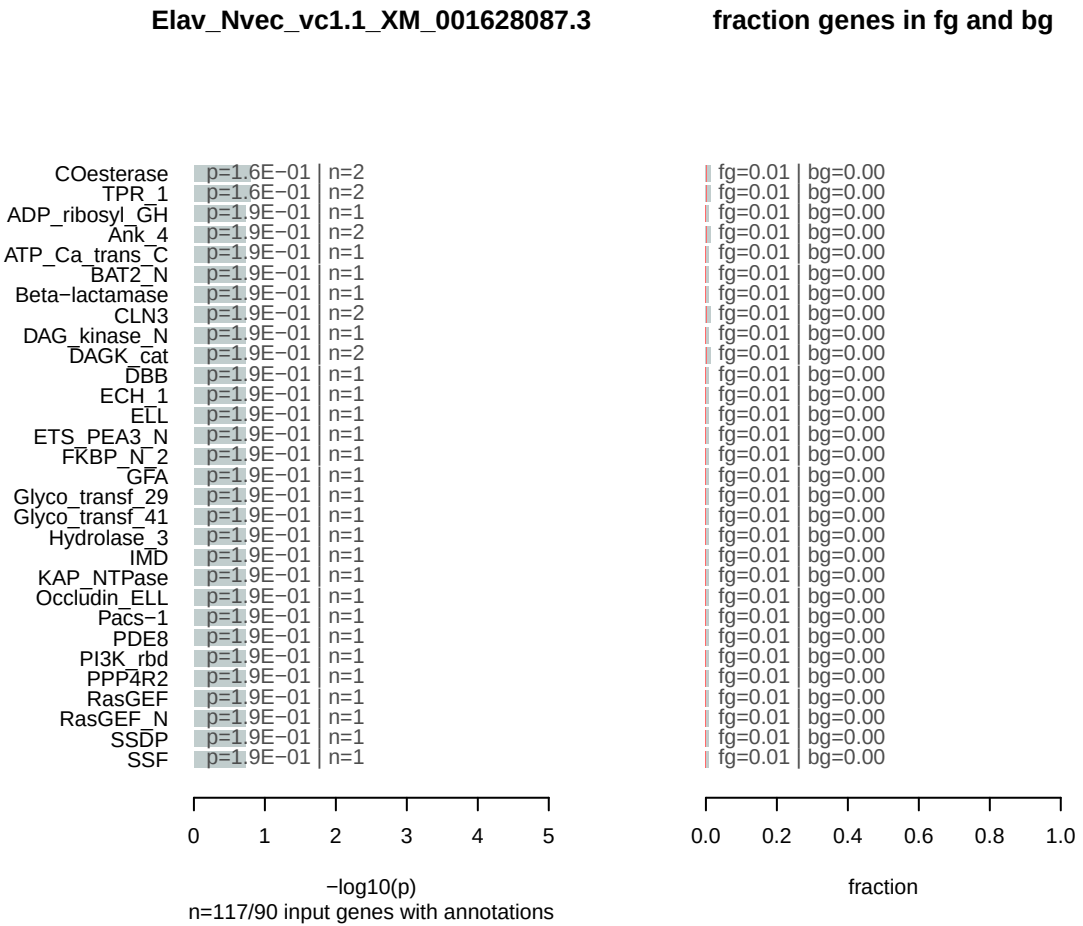
Elav_Nvec_vc1.1_XM_001628087.3
fraction genes in fg and bg
COesterase p=1.6E−01 n=2
TPR_1 p=1.6E−01 n=2
ADP_ribosyl_GH p=1.9E−01 n=1
Ank_4 p=1.9E−01 n=2
ATP_Ca_trans_C p=1.9E−01 n=1
BAT2_N p=1.9E−01 n=1
Beta−lactamase p=1.9E−01 n=1
CLN3 p=1.9E−01 n=2
DAG_kinase_N p=1.9E−01 n=1
DAGK_cat p=1.9E−01 n=2
DBB p=1.9E−01 n=1
ECH_1 p=1.9E−01 n=1
ELL p=1.9E−01 n=1
ETS_PEA3_N p=1.9E−01 n=1
FKBP_N_2 p=1.9E−01 n=1
GFA p=1.9E−01 n=1
Glyco_transf_29 p=1.9E−01 n=1
Glyco_transf_41 p=1.9E−01 n=1
Hydrolase_3 p=1.9E−01 n=1
IMD p=1.9E−01 n=1
KAP_NTPase p=1.9E−01 n=1
Occludin_ELL p=1.9E−01 n=1
Pacs−1 p=1.9E−01 n=1
PDE8 p=1.9E−01 n=1
PI3K_rbd p=1.9E−01 n=1
PPP4R2 p=1.9E−01 n=1
RasGEF p=1.9E−01 n=1
RasGEF_N p=1.9E−01 n=1
SSDP p=1.9E−01 n=1
SSF p=1.9E−01 n=1
fg=0.01 bg=0.00
0 1 2 3 4 5
−log10(p)
n=117/90 input genes with annotations
0.0 0.2 0.4 0.6 0.8 1.0
fraction