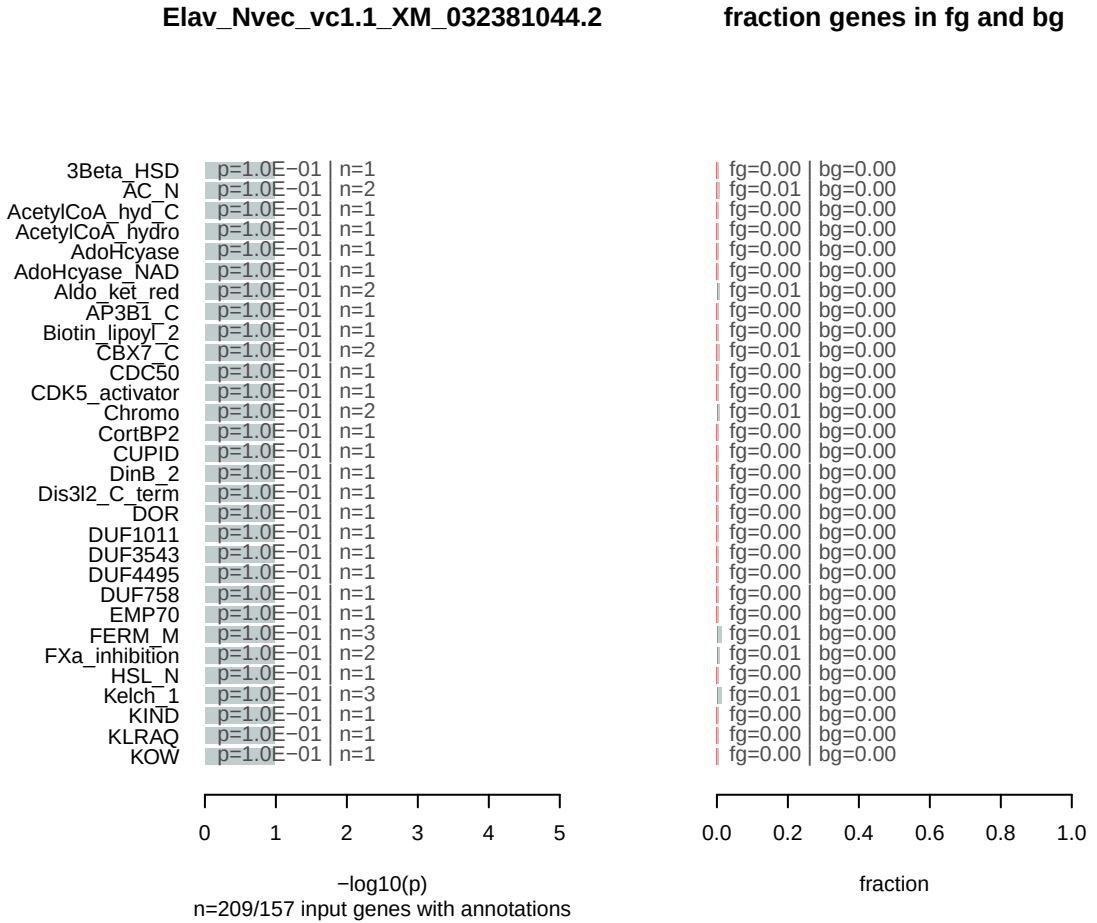
Elav_Nvec_vc1.1_XM_032381044.2
fraction genes in fg and bg
3Beta_HSD p=1.0E−01 n=1 fg=0.00 bg=0.00
AC_N p=1.0E−01 n=2 fg=0.01 bg=0.00
AcetylCoA_hyd_C p=1.0E−01 n=1 fg=0.00 bg=0.00
AcetylCoA_hydro p=1.0E−01 n=1 fg=0.00 bg=0.00
AdoHcyase p=1.0E−01 n=1 fg=0.00 bg=0.00
AdoHcyase_NAD p=1.0E−01 n=1 fg=0.00 bg=0.00
Aldo_ket_red p=1.0E−01 n=2 fg=0.01 bg=0.00
AP3B1_C p=1.0E−01 n=1 fg=0.00 bg=0.00
Biotin_lipoyl_2 p=1.0E−01 n=1 fg=0.00 bg=0.00
CBX7_C p=1.0E−01 n=2 fg=0.01 bg=0.00
CDC50 p=1.0E−01 n=1 fg=0.00 bg=0.00
CDK5_activator p=1.0E−01 n=1 fg=0.00 bg=0.00
Chromo p=1.0E−01 n=2 fg=0.01 bg=0.00
CortBP2 p=1.0E−01 n=1 fg=0.00 bg=0.00
CUPID p=1.0E−01 n=1 fg=0.00 bg=0.00
DinB_2 p=1.0E−01 n=1 fg=0.00 bg=0.00
Dis3l2_C_term p=1.0E−01 n=1 fg=0.00 bg=0.00
DOR p=1.0E−01 n=1 fg=0.00 bg=0.00
DUF1011 p=1.0E−01 n=1 fg=0.00 bg=0.00
DUF3543 p=1.0E−01 n=1 fg=0.00 bg=0.00
DUF4495 p=1.0E−01 n=1 fg=0.00 bg=0.00
DUF758 p=1.0E−01 n=1 fg=0.00 bg=0.00
EMP70 p=1.0E−01 n=1 fg=0.00 bg=0.00
FERM_M p=1.0E−01 n=3 fg=0.01 bg=0.00
FXa_inhibition p=1.0E−01 n=2 fg=0.01 bg=0.00
HSL_N p=1.0E−01 n=1 fg=0.00 bg=0.00
Kelch_1 p=1.0E−01 n=3 fg=0.01 bg=0.00
KIND p=1.0E−01 n=1 fg=0.00 bg=0.00
KLRAQ p=1.0E−01 n=1 fg=0.00 bg=0.00
KOW p=1.0E−01 n=1 fg=0.00 bg=0.00
0 1 2 3 4 5
−log10(p)
n=209/157 input genes with annotations
0.0 0.2 0.4 0.6 0.8 1.0
fraction