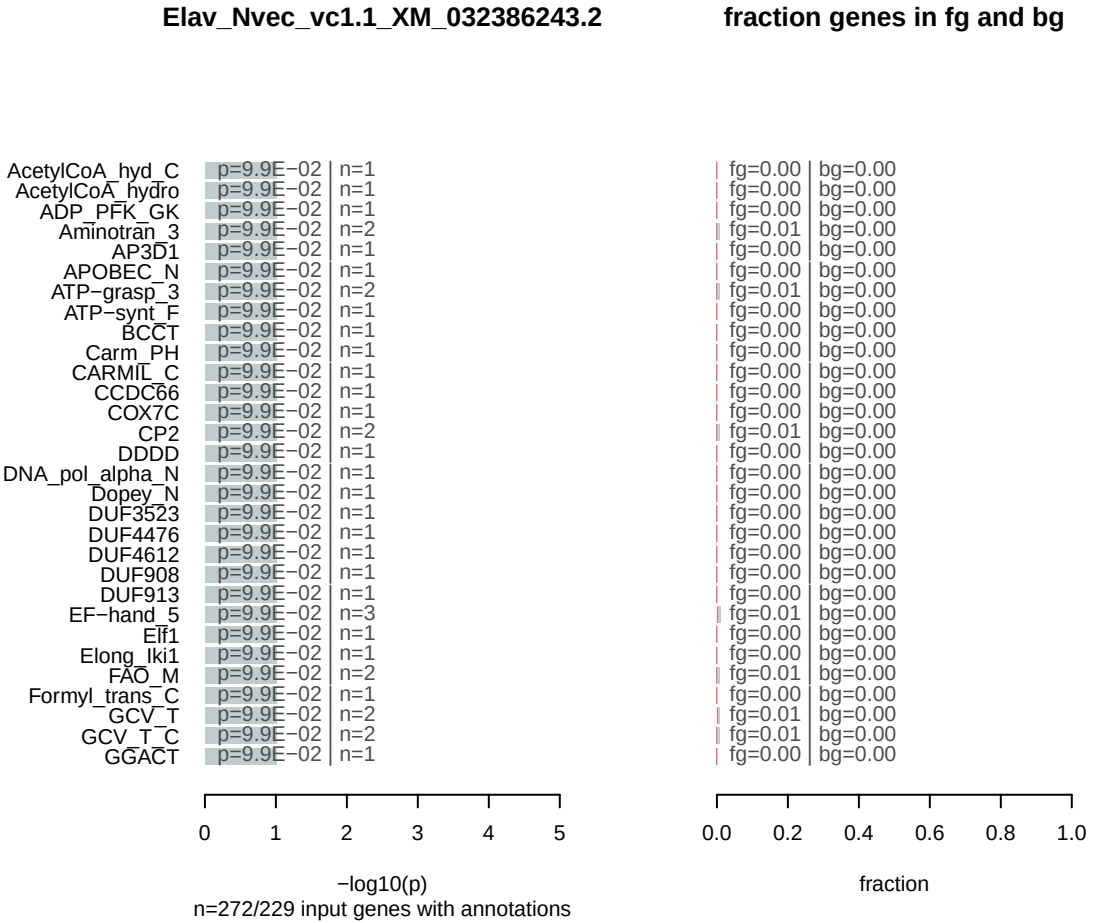
Elav_Nvec_vc1.1_XM_032386243.2
fraction genes in fg and bg
AcetylCoA_hyd_C p=9.9E−02 n=1 fg=0.00 bg=0.00
AcetylCoA_hydro p=9.9E−02 n=1 fg=0.00 bg=0.00
ADP_PFK_GK p=9.9E−02 n=1 fg=0.00 bg=0.00
Aminotran_3 p=9.9E−02 n=2 fg=0.01 bg=0.00
AP3D1 p=9.9E−02 n=1 fg=0.00 bg=0.00
APOBEC_N p=9.9E−02 n=1 fg=0.00 bg=0.00
ATP−grasp_3 p=9.9E−02 n=2 fg=0.01 bg=0.00
ATP−synt_F p=9.9E−02 n=1 fg=0.00 bg=0.00
BCCT p=9.9E−02 n=1 fg=0.00 bg=0.00
Carm_PH p=9.9E−02 n=1 fg=0.00 bg=0.00
CARMIL_C p=9.9E−02 n=1 fg=0.00 bg=0.00
CCDC66 p=9.9E−02 n=1 fg=0.00 bg=0.00
COX7C p=9.9E−02 n=1 fg=0.00 bg=0.00
CP2 p=9.9E−02 n=2 fg=0.01 bg=0.00
DDDD p=9.9E−02 n=1 fg=0.00 bg=0.00
DNA_pol_alpha_N p=9.9E−02 n=1 fg=0.00 bg=0.00
Dopey_N p=9.9E−02 n=1 fg=0.00 bg=0.00
DUF3523 p=9.9E−02 n=1 fg=0.00 bg=0.00
DUF4476 p=9.9E−02 n=1 fg=0.00 bg=0.00
DUF4612 p=9.9E−02 n=1 fg=0.00 bg=0.00
DUF908 p=9.9E−02 n=1 fg=0.00 bg=0.00
DUF913 p=9.9E−02 n=1 fg=0.00 bg=0.00
EF−hand_5 p=9.9E−02 n=3 fg=0.01 bg=0.00
Elf1 p=9.9E−02 n=1 fg=0.00 bg=0.00
Elong_Iki1 p=9.9E−02 n=1 fg=0.00 bg=0.00
FAO_M p=9.9E−02 n=2 fg=0.01 bg=0.00
Formyl_trans_C p=9.9E−02 n=1 fg=0.00 bg=0.00
GCV_T p=9.9E−02 n=2 fg=0.01 bg=0.00
GCV_T_C p=9.9E−02 n=2 fg=0.01 bg=0.00
GGACT p=9.9E−02 n=1 fg=0.00 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=272/229 input genes with annotations