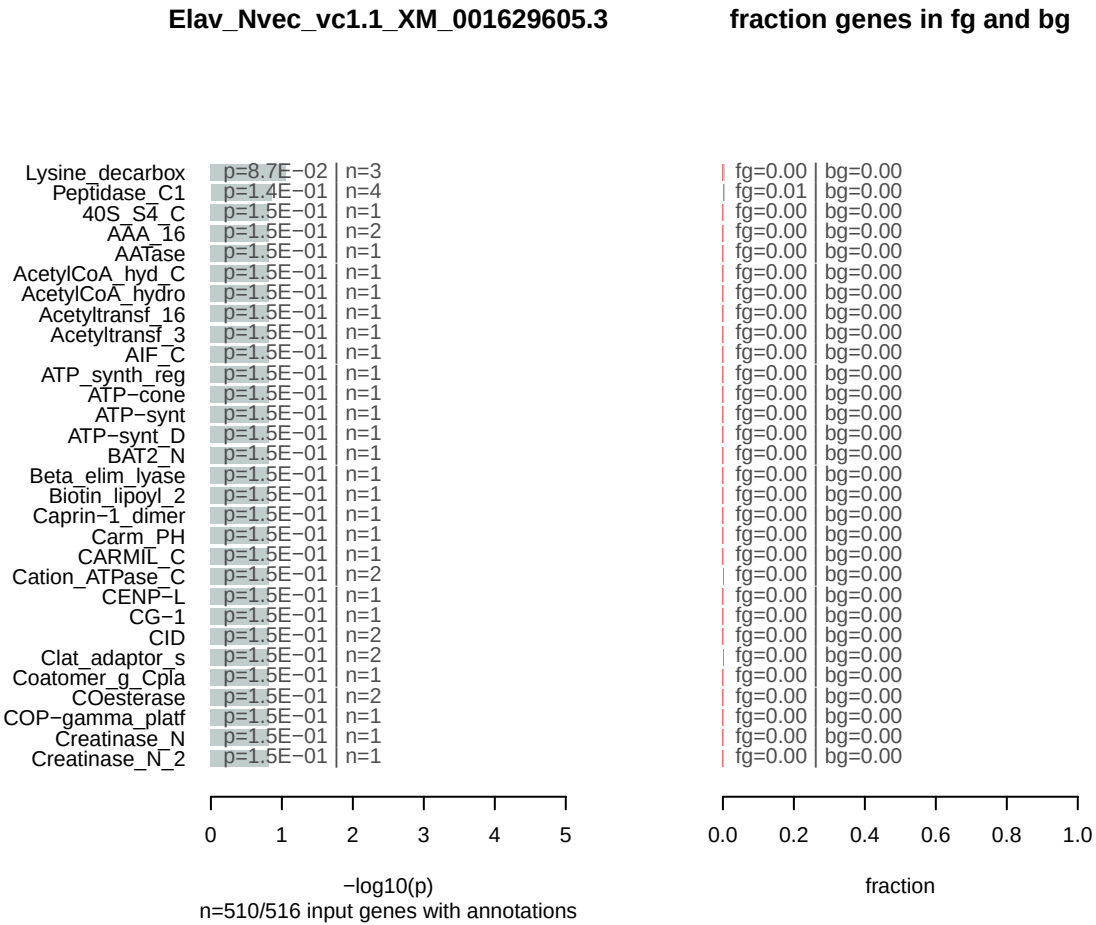
Elav_Nvec_vc1.1_XM_001629605.3
fraction genes in fg and bg
Lysine_decarbox p=8.7E−02 n=3 fg=0.00 bg=0.00
Peptidase_C1 p=1.4E−01 n=4 fg=0.01 bg=0.00
40S_S4_C p=1.5E−01 n=1 fg=0.00 bg=0.00
AAA_16 p=1.5E−01 n=2 fg=0.00 bg=0.00
AATase p=1.5E−01 n=1 fg=0.00 bg=0.00
AcetylCoA_hyd_C p=1.5E−01 n=1 fg=0.00 bg=0.00
AcetylCoA_hydro p=1.5E−01 n=1 fg=0.00 bg=0.00
Acetyltransf_16 p=1.5E−01 n=1 fg=0.00 bg=0.00
Acetyltransf_3 p=1.5E−01 n=1 fg=0.00 bg=0.00
AIF_C p=1.5E−01 n=1 fg=0.00 bg=0.00
ATP_synth_reg p=1.5E−01 n=1 fg=0.00 bg=0.00
ATP−cone p=1.5E−01 n=1 fg=0.00 bg=0.00
ATP−synt p=1.5E−01 n=1 fg=0.00 bg=0.00
ATP−synt_D 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
Beta_elim_lyase p=1.5E−01 n=1 fg=0.00 bg=0.00
Biotin_lipoyl_2 p=1.5E−01 n=1 fg=0.00 bg=0.00
Caprin−1_dimer p=1.5E−01 n=1 fg=0.00 bg=0.00
Carm_PH p=1.5E−01 n=1 fg=0.00 bg=0.00
CARMIL_C p=1.5E−01 n=1 fg=0.00 bg=0.00
Cation_ATPase_C p=1.5E−01 n=2 fg=0.00 bg=0.00
CENP−L p=1.5E−01 n=1 fg=0.00 bg=0.00
CG−1 p=1.5E−01 n=1 fg=0.00 bg=0.00
CID p=1.5E−01 n=2 fg=0.00 bg=0.00
Clat_adaptor_s p=1.5E−01 n=2 fg=0.00 bg=0.00
Coatomer_g_Cpla 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
COP−gamma_platf p=1.5E−01 n=1 fg=0.00 bg=0.00
Creatinase_N p=1.5E−01 n=1 fg=0.00 bg=0.00
Creatinase_N_2 p=1.5E−01 n=1 fg=0.00 bg=0.00
0 1 2 3 4 5
−log10(p)
n=510/516 input genes with annotations
0.0 0.2 0.4 0.6 0.8 1.0
fraction