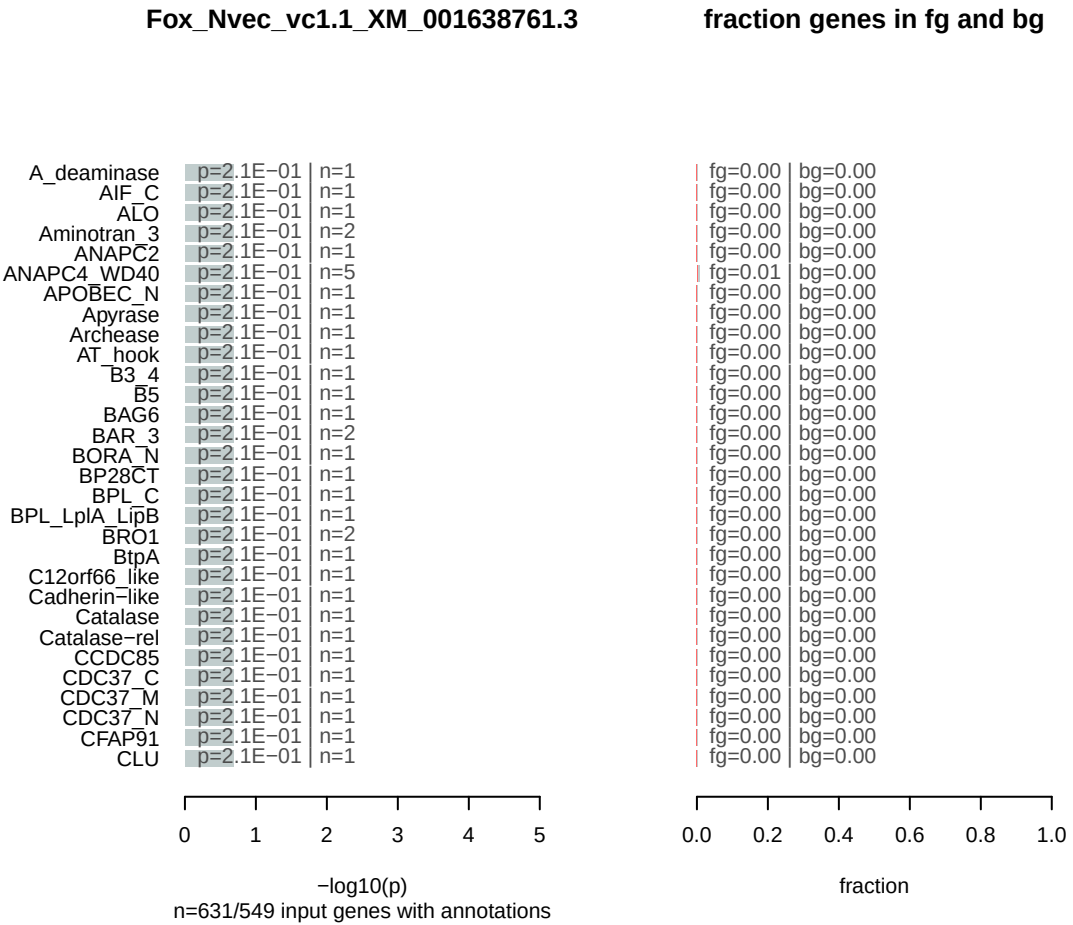
Fox_Nvec_vc1.1_XM_001638761.3
fraction genes in fg and bg
A_deaminase p=2.1E−01 n=1 fg=0.00 bg=0.00
AIF_C p=2.1E−01 n=1 fg=0.00 bg=0.00
ALO p=2.1E−01 n=1 fg=0.00 bg=0.00
Aminotran_3 p=2.1E−01 n=2 fg=0.00 bg=0.00
ANAPC2 p=2.1E−01 n=1 fg=0.00 bg=0.00
ANAPC4_WD40 p=2.1E−01 n=5 fg=0.01 bg=0.00
APOBEC_N p=2.1E−01 n=1 fg=0.00 bg=0.00
Apyrase p=2.1E−01 n=1 fg=0.00 bg=0.00
Archease p=2.1E−01 n=1 fg=0.00 bg=0.00
AT_hook p=2.1E−01 n=1 fg=0.00 bg=0.00
B3_4 p=2.1E−01 n=1 fg=0.00 bg=0.00
B5 p=2.1E−01 n=1 fg=0.00 bg=0.00
BAG6 p=2.1E−01 n=1 fg=0.00 bg=0.00
BAR_3 p=2.1E−01 n=2 fg=0.00 bg=0.00
BORA_N p=2.1E−01 n=1 fg=0.00 bg=0.00
BP28CT p=2.1E−01 n=1 fg=0.00 bg=0.00
BPL_C p=2.1E−01 n=1 fg=0.00 bg=0.00
BPL_LplA_LipB p=2.1E−01 n=1 fg=0.00 bg=0.00
BRO1 p=2.1E−01 n=2 fg=0.00 bg=0.00
BtpA p=2.1E−01 n=1 fg=0.00 bg=0.00
C12orf66_like p=2.1E−01 n=1 fg=0.00 bg=0.00
Cadherin−like p=2.1E−01 n=1 fg=0.00 bg=0.00
Catalase p=2.1E−01 n=1 fg=0.00 bg=0.00
Catalase−rel p=2.1E−01 n=1 fg=0.00 bg=0.00
CCDC85 p=2.1E−01 n=1 fg=0.00 bg=0.00
CDC37_C p=2.1E−01 n=1 fg=0.00 bg=0.00
CDC37_M p=2.1E−01 n=1 fg=0.00 bg=0.00
CDC37_N p=2.1E−01 n=1 fg=0.00 bg=0.00
CFAP91 p=2.1E−01 n=1 fg=0.00 bg=0.00
CLU p=2.1E−01 n=1 fg=0.00 bg=0.00
0 1 2 3 4 5
−log10(p)
n=631/549 input genes with annotations
0.0 0.2 0.4 0.6 0.8 1.0
fraction