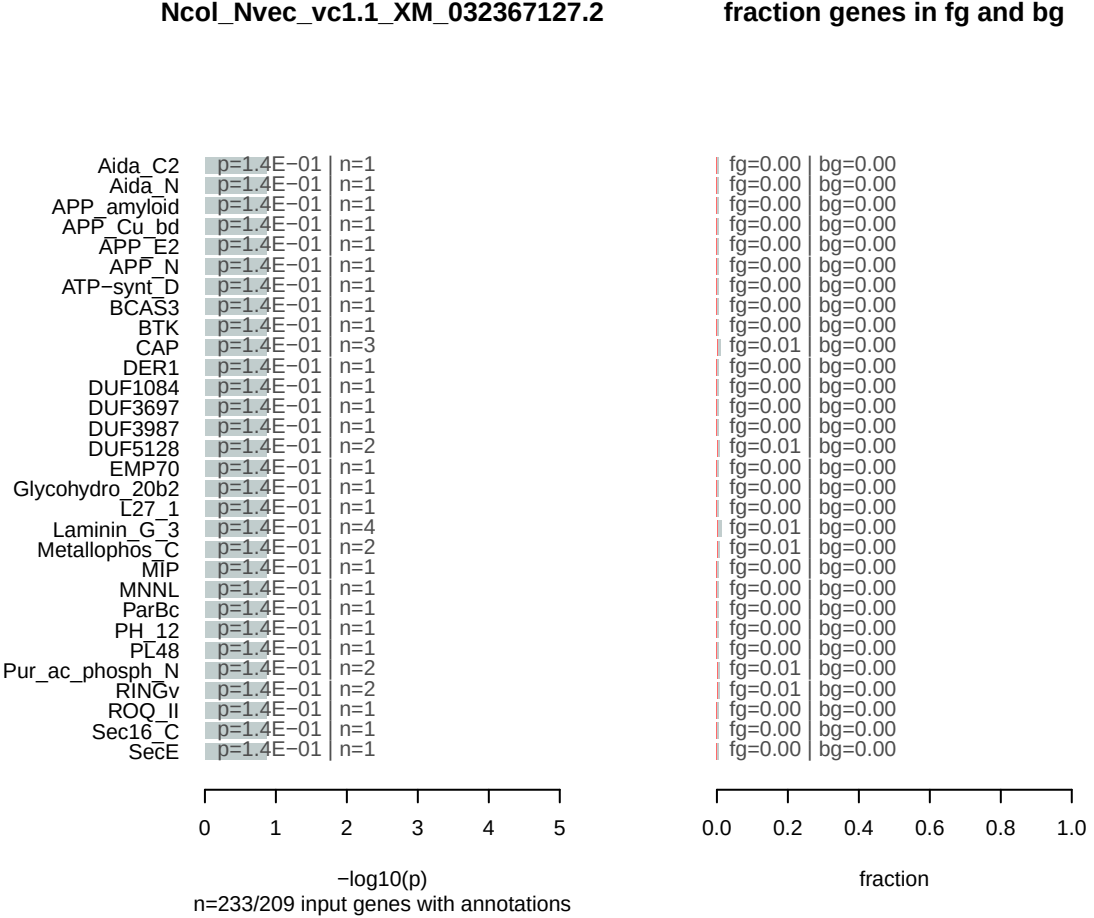
Ncol_Nvec_vc1.1_XM_032367127.2
fraction genes in fg and bg
Aida_C2 p=1.4E−01 n=1 fg=0.00 bg=0.00
Aida_N p=1.4E−01 n=1 fg=0.00 bg=0.00
APP_amyloid p=1.4E−01 n=1 fg=0.00 bg=0.00
APP_Cu_bd p=1.4E−01 n=1 fg=0.00 bg=0.00
APP_E2 p=1.4E−01 n=1 fg=0.00 bg=0.00
APP_N p=1.4E−01 n=1 fg=0.00 bg=0.00
ATP−synt_D p=1.4E−01 n=1 fg=0.00 bg=0.00
BCAS3 p=1.4E−01 n=1 fg=0.00 bg=0.00
BTK p=1.4E−01 n=1 fg=0.00 bg=0.00
CAP p=1.4E−01 n=3 fg=0.01 bg=0.00
DER1 p=1.4E−01 n=1 fg=0.00 bg=0.00
DUF1084 p=1.4E−01 n=1 fg=0.00 bg=0.00
DUF3697 p=1.4E−01 n=1 fg=0.00 bg=0.00
DUF3987 p=1.4E−01 n=1 fg=0.00 bg=0.00
DUF5128 p=1.4E−01 n=2 fg=0.01 bg=0.00
EMP70 p=1.4E−01 n=1 fg=0.00 bg=0.00
Glycohydro_20b2 p=1.4E−01 n=1 fg=0.00 bg=0.00
L27_1 p=1.4E−01 n=1 fg=0.00 bg=0.00
Laminin_G_3 p=1.4E−01 n=4 fg=0.01 bg=0.00
Metallophos_C p=1.4E−01 n=2 fg=0.01 bg=0.00
MIP p=1.4E−01 n=1 fg=0.00 bg=0.00
MNNL p=1.4E−01 n=1 fg=0.00 bg=0.00
ParBc p=1.4E−01 n=1 fg=0.00 bg=0.00
PH_12 p=1.4E−01 n=1 fg=0.00 bg=0.00
PL48 p=1.4E−01 n=1 fg=0.00 bg=0.00
Pur_ac_phosph_N p=1.4E−01 n=2 fg=0.01 bg=0.00
RINGv p=1.4E−01 n=2 fg=0.01 bg=0.00
ROQ_II p=1.4E−01 n=1 fg=0.00 bg=0.00
Sec16_C p=1.4E−01 n=1 fg=0.00 bg=0.00
SecE p=1.4E−01 n=1 fg=0.00 bg=0.00
0 1 2 3 4 5
−log10(p)
n=233/209 input genes with annotations
0.0 0.2 0.4 0.6 0.8 1.0
fraction