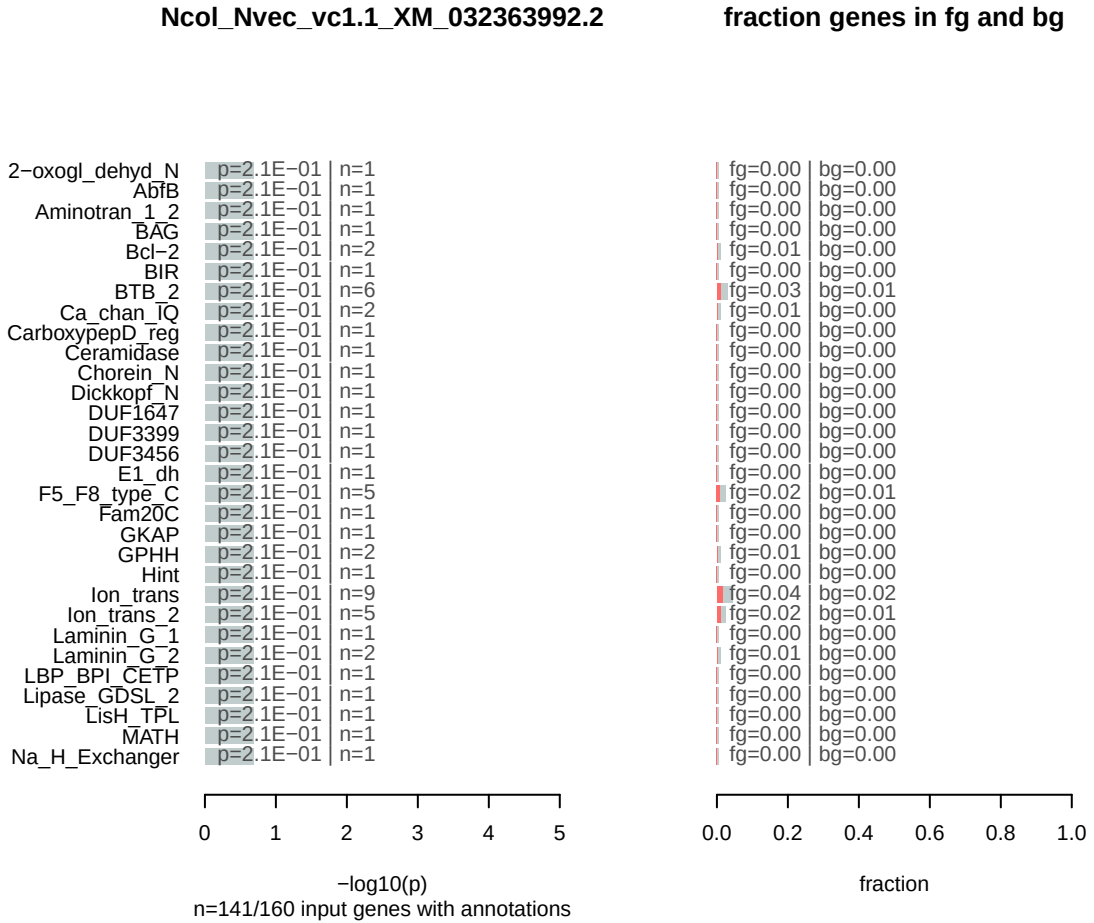
NcoI_Nvec_vc1.1_XM_032363992.2
fraction genes in fg and bg
2−oxogl_dehyd_N p=2.1E−01 n=1 fg=0.00 bg=0.00
AbfB p=2.1E−01 n=1 fg=0.00 bg=0.00
Aminotran_1_2 p=2.1E−01 n=1 fg=0.00 bg=0.00
BAG p=2.1E−01 n=1 fg=0.00 bg=0.00
Bcl−2 p=2.1E−01 n=2 fg=0.01 bg=0.00
BIR p=2.1E−01 n=1 fg=0.00 bg=0.00
BTB_2 p=2.1E−01 n=6 fg=0.03 bg=0.01
Ca_chan_IQ p=2.1E−01 n=2 fg=0.01 bg=0.00
CarboxypepD_reg p=2.1E−01 n=1 fg=0.00 bg=0.00
Ceramidase p=2.1E−01 n=1 fg=0.00 bg=0.00
Chorein_N p=2.1E−01 n=1 fg=0.00 bg=0.00
Dickkopf_N p=2.1E−01 n=1 fg=0.00 bg=0.00
DUF1647 p=2.1E−01 n=1 fg=0.00 bg=0.00
DUF3399 p=2.1E−01 n=1 fg=0.00 bg=0.00
DUF3456 p=2.1E−01 n=1 fg=0.00 bg=0.00
E1_dh p=2.1E−01 n=1 fg=0.00 bg=0.00
F5_F8_type_C p=2.1E−01 n=5 fg=0.02 bg=0.01
Fam20C p=2.1E−01 n=1 fg=0.00 bg=0.00
GKAP p=2.1E−01 n=1 fg=0.00 bg=0.00
GPHH p=2.1E−01 n=2 fg=0.01 bg=0.00
Hint p=2.1E−01 n=1 fg=0.00 bg=0.00
Ion_trans p=2.1E−01 n=9 fg=0.04 bg=0.02
Ion_trans_2 p=2.1E−01 n=5 fg=0.02 bg=0.01
Laminin_G_1 p=2.1E−01 n=1 fg=0.00 bg=0.00
Laminin_G_2 p=2.1E−01 n=2 fg=0.01 bg=0.00
LBP_BPI_CETP p=2.1E−01 n=1 fg=0.00 bg=0.00
Lipase_GDSL_2 p=2.1E−01 n=1 fg=0.00 bg=0.00
LisH_TPL p=2.1E−01 n=1 fg=0.00 bg=0.00
MATH p=2.1E−01 n=1 fg=0.00 bg=0.00
Na_H_Exchanger p=2.1E−01 n=1 fg=0.00 bg=0.00
0 1 2 3 4 5
−log10(p)
n=141/160 input genes with annotations
0.0 0.2 0.4 0.6 0.8 1.0
fraction