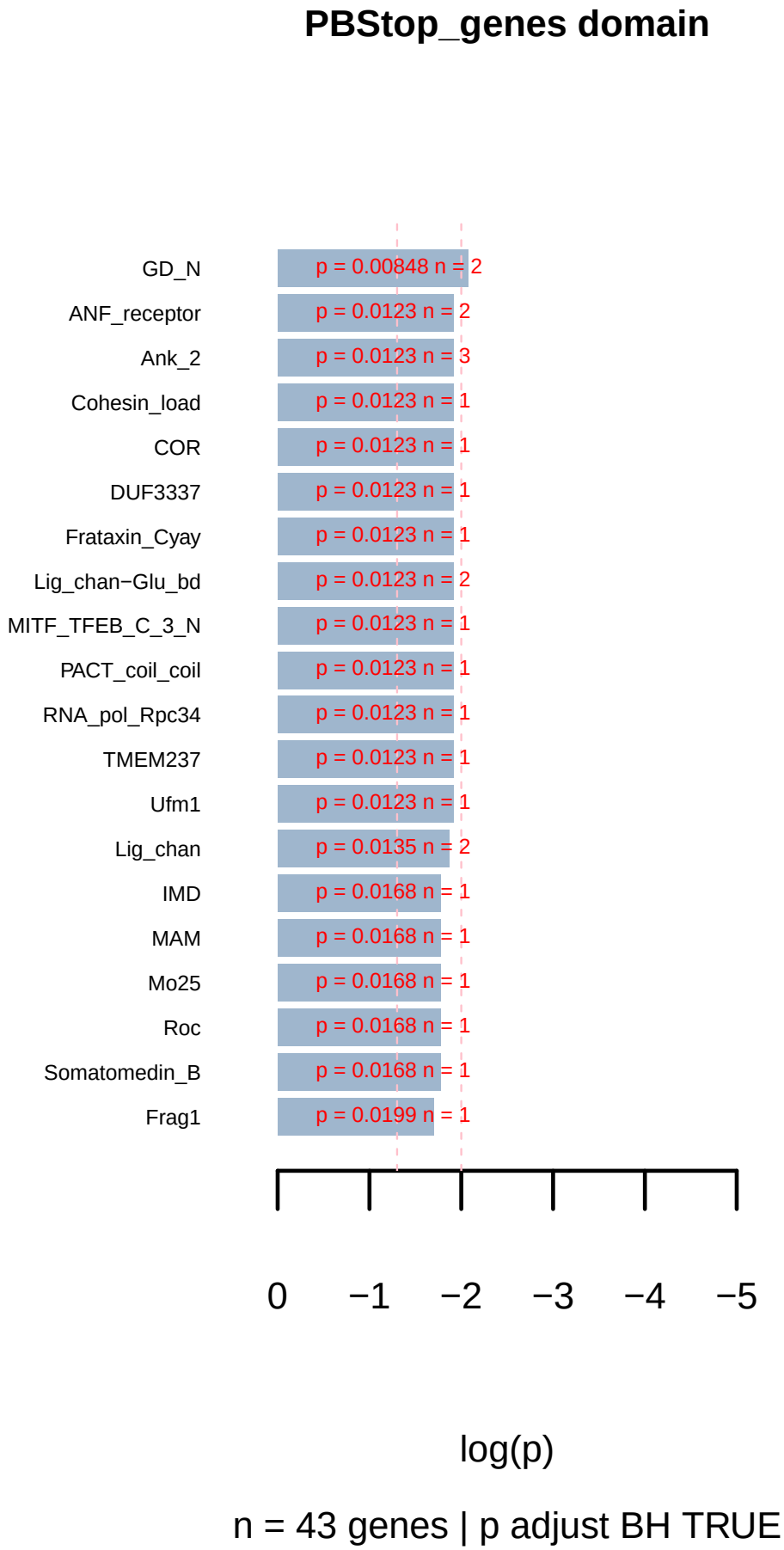
PBStop_genes domain
GD_N
ANF_receptor
Ank_2
Cohesin_load
COR
DUF3337
Frataxin_Cyay
Lig_chan−Glu_bd
MITF_TFEB_C_3_N
PACT_coil_coil
RNA_pol_Rpc34
TMEM237
Ufm1
Lig_chan
IMD
MAM
Mo25
Roc
Somatomedin_B
Frag1
p = 0.00848 n = 2
p = 0.0123 n = 2
p = 0.0123 n = 3
p = 0.0123 n = 1
p = 0.0123 n = 1
p = 0.0123 n = 1
p = 0.0123 n = 1
p = 0.0123 n = 2
p = 0.0123 n = 1
p = 0.0123 n = 1
p = 0.0123 n = 1
p = 0.0123 n = 1
p = 0.0123 n = 1
p = 0.0135 n = 2
p = 0.0168 n = 1
p = 0.0168 n = 1
p = 0.0168 n = 1
p = 0.0168 n = 1
p = 0.0168 n = 1
p = 0.0199 n = 1
0
−1
−2
−3
−4
−5
log(p)
n = 43 genes | p adjust BH TRUE