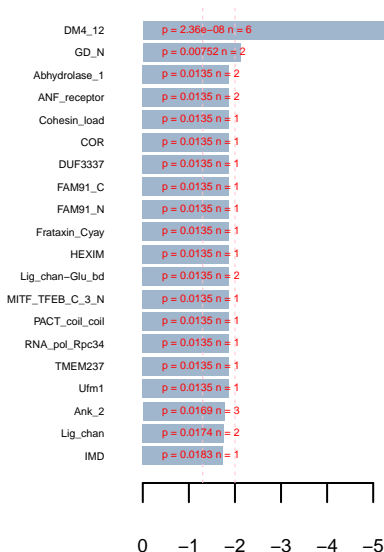


## PBStop\_genes domain



log(p)

n = 56 genes | p adjust BH TRUE