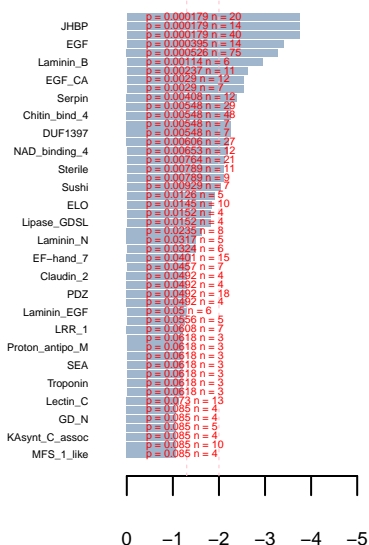


## ALL.RUN-RFE\_difpos domain



n = 1936 genes | p adjust BH TRUE