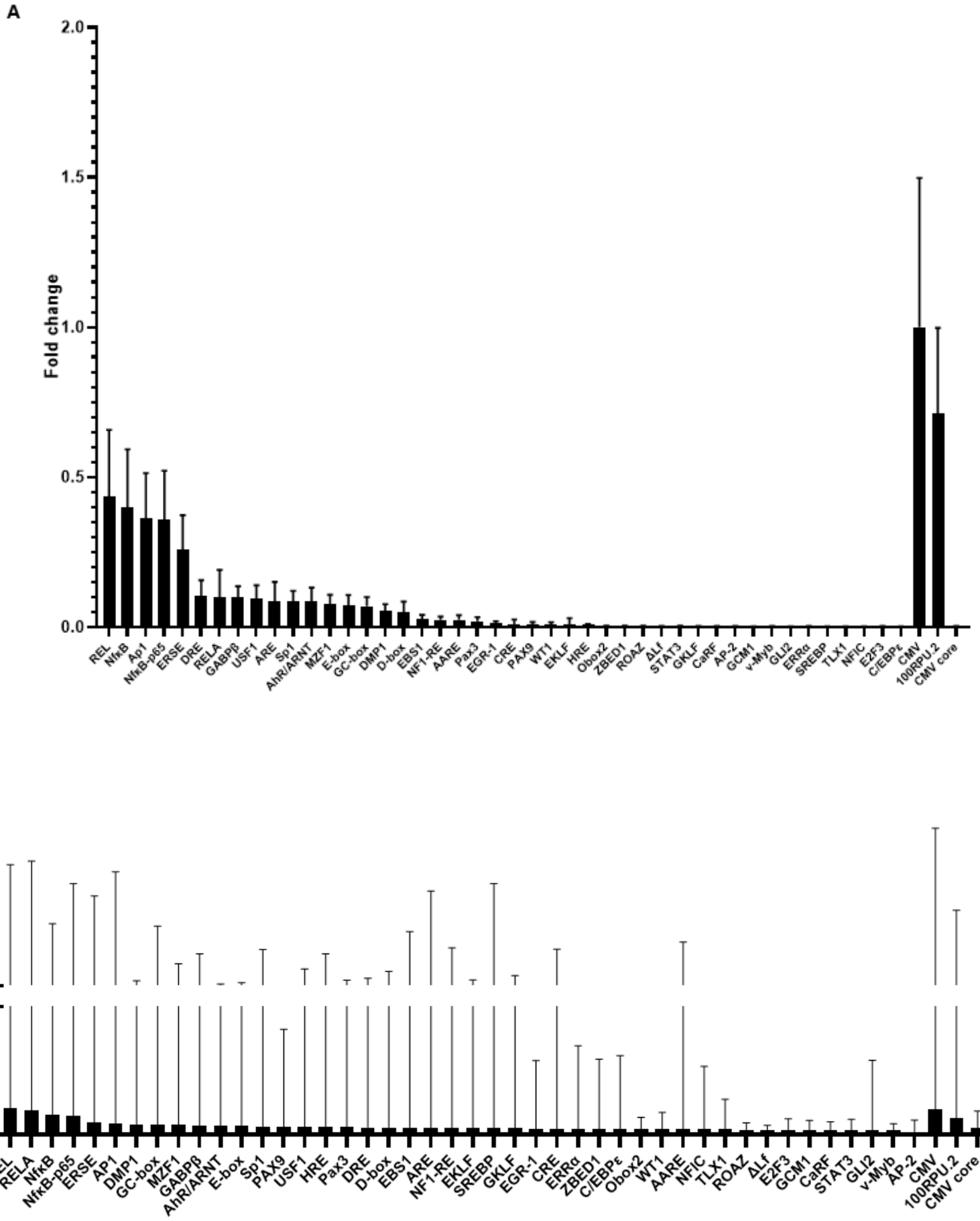Fig. S3: Screening of homotypic promoters (A) Eight copies of each TFRE were positioned upstream of a CMV core promoter in a reporter vector encoding mCherry. The fold change was calculated by normalizing with MFI of the full CMV promoter. (B) MFIs of mCherry driven by different homotypic promoters in transient condition. Fold change was calculated by normalizing with MFI of full CMV promoter.