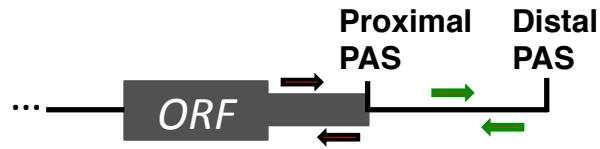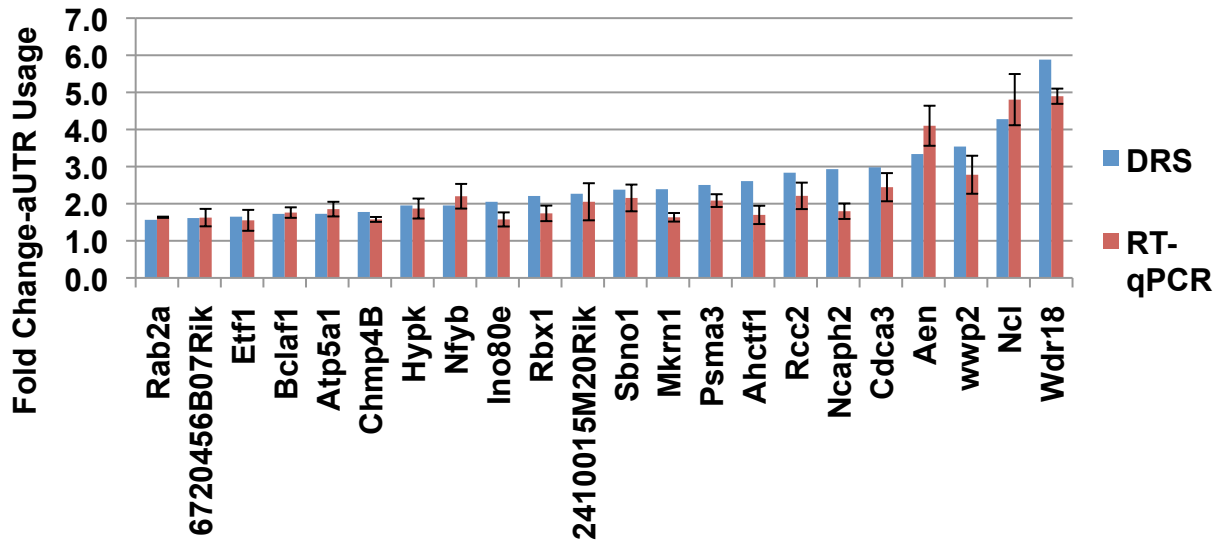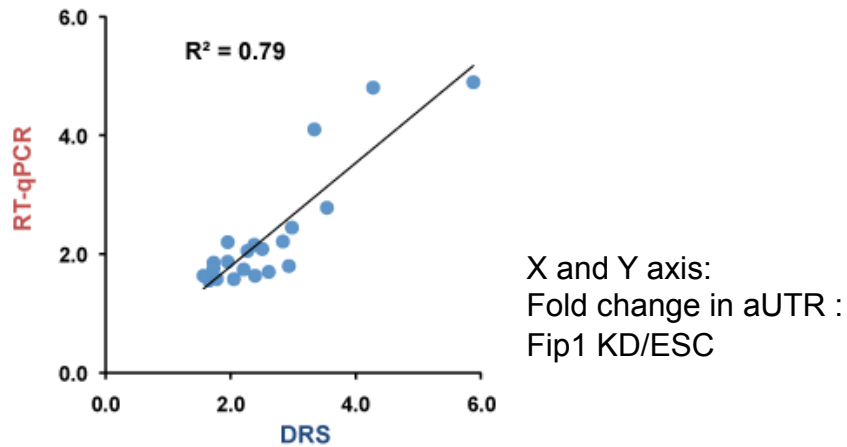**Figure S8. Validation of APA fold change by RT-qPCR.** (A) Primer pairs used to detect common region and the eUTRs. (B) Comparison of relative fold change of aUTR (aUTR/common region) in Mock- or siFip1-2- treated ESCs based on DRS (blue columns) and RT-qPCR (shown are mean ± SEM; n=3). (C) Correlation between fold change in aUTR usage based on DRS (X axis) and RT-qPCR as shown in B. R^2=0.79.