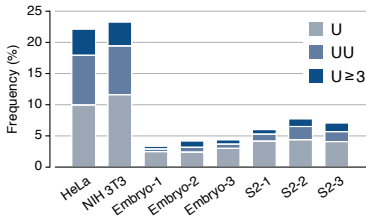
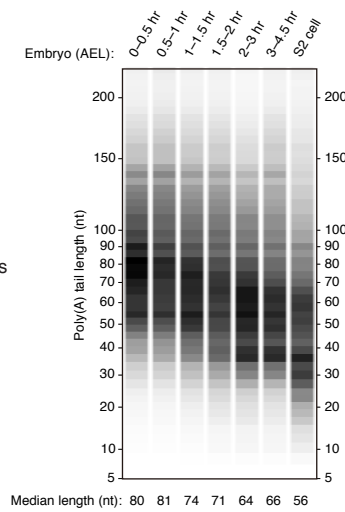


Lim_FigS2

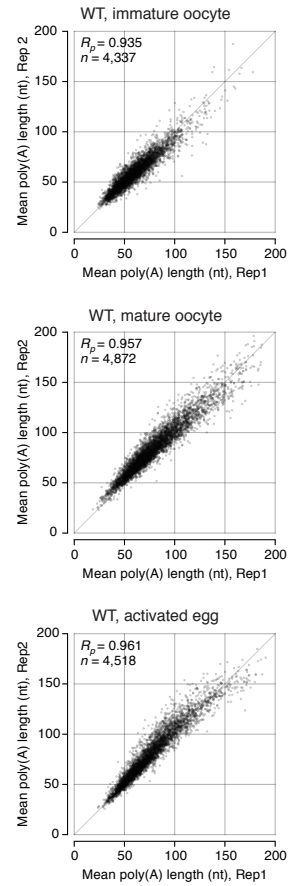
A 3' end uridylation frequency (by TAIL-seq)



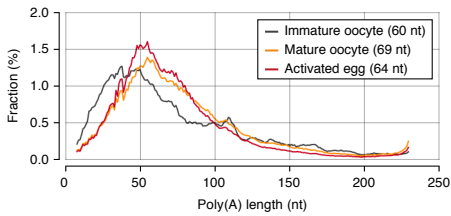
B Virtual gel image of poly(A) tail length distribution



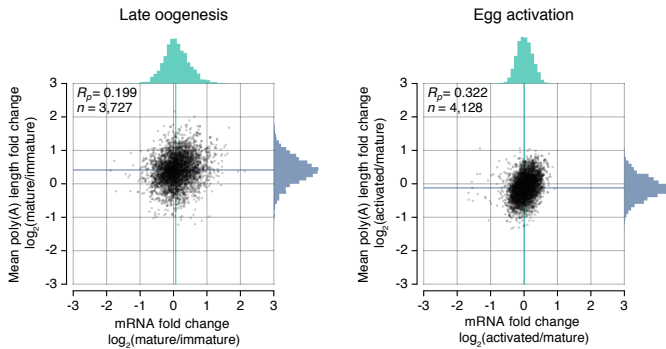
C Reproducibility of mTAIL-seq libraries



D Global measurement of poly(A) tail length in replicates



E Comparison of poly(A) length changes to mRNA abundance changes



F Hire-PAT assay

