

Supplementary Information

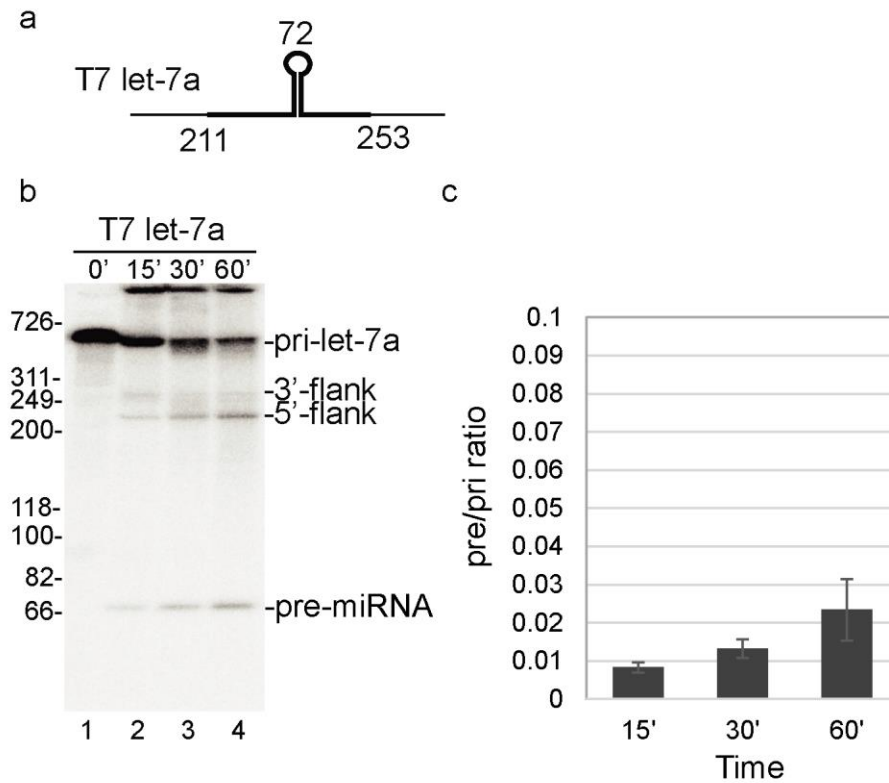
Title:

Primary microRNA processing is functionally coupled to RNAP II transcription in vitro

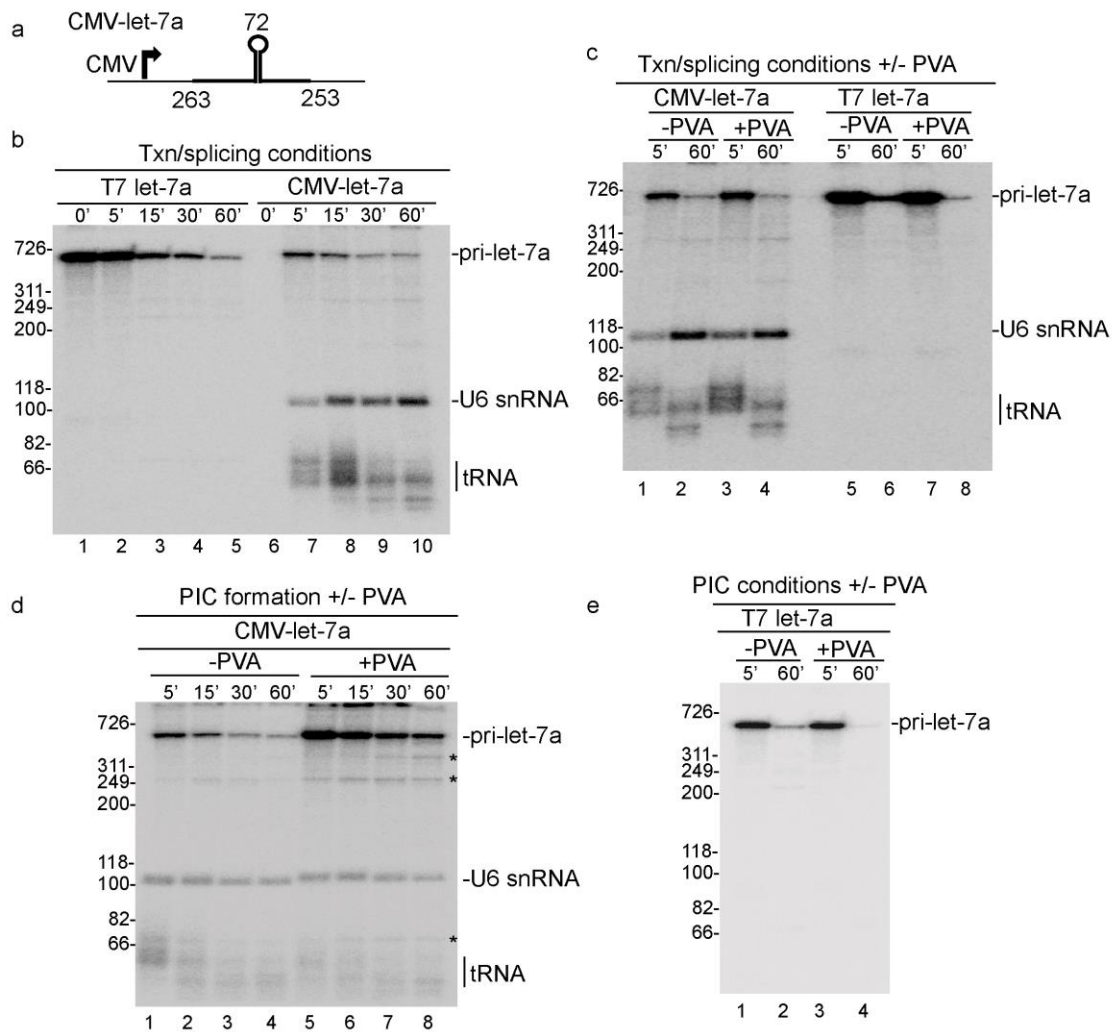
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Supplementary Information



Supplementary Figure S1. Pri-miRNA processing of naked T7 let-7a pri-miRNA in vitro. (a) Schematic showing the T7 let-7a pri-miRNA transcript used for in vitro processing. (b) T7 let-7a pri-miRNA was incubated with nuclear extract in the presence of 6.4 mM MgCl₂ for the indicated times. Total RNA was isolated, run on an 8% denaturing polyacrylamide gel, and bands detected by phosphorimager. (c) The percentage of T7 let-7a pri-miRNA processed at each time point in panel C was determined as the amount of pre-let-7a relative to pri-mRNA at the start of the reaction. RNAs were detected by phosphorimager and quantified using Quantity One software. Mean \pm S.D. of three biological replicates are shown.



Supplementary Figure S2. Optimization of RNAP II txn/pri-miRNA

processing conditions. (a) Structure of the DNA template encoding CMV-let-7a

pri-miRNA used to optimize RNAP II transcription and pri-miRNA processing.

The sizes of the 5' flank, pre-miRNA, and 3' flank are shown. (b) T7 let-7a pri-

miRNA or CMV DNA template encoding let-7a pri-miRNA was incubated with

nuclear extract under txn/splicing conditions (with 3.2 mM MgCl₂), for the

indicated times. Total RNA was isolated and run on an 8% denaturing

Supplementary Figure S3. Coupled txn/pri-miRNA processing using two different miRNAs. (a, b) CMV DNA templates encoding miR-21 pri-miRNA (a) or encoding miR-26 (b) pri-miRNA or the corresponding T7 transcripts were incubated in nuclear extracts in the txn/pri-miRNA processing system for the indicated times. Total RNA was isolated and fractionated on an 8% denaturing polyacrylamide gel. The RNA species are labeled and ori indicates the gel origin.