

Figure A. Most of the TOFU transcripts have longer UTRs than current annotation.

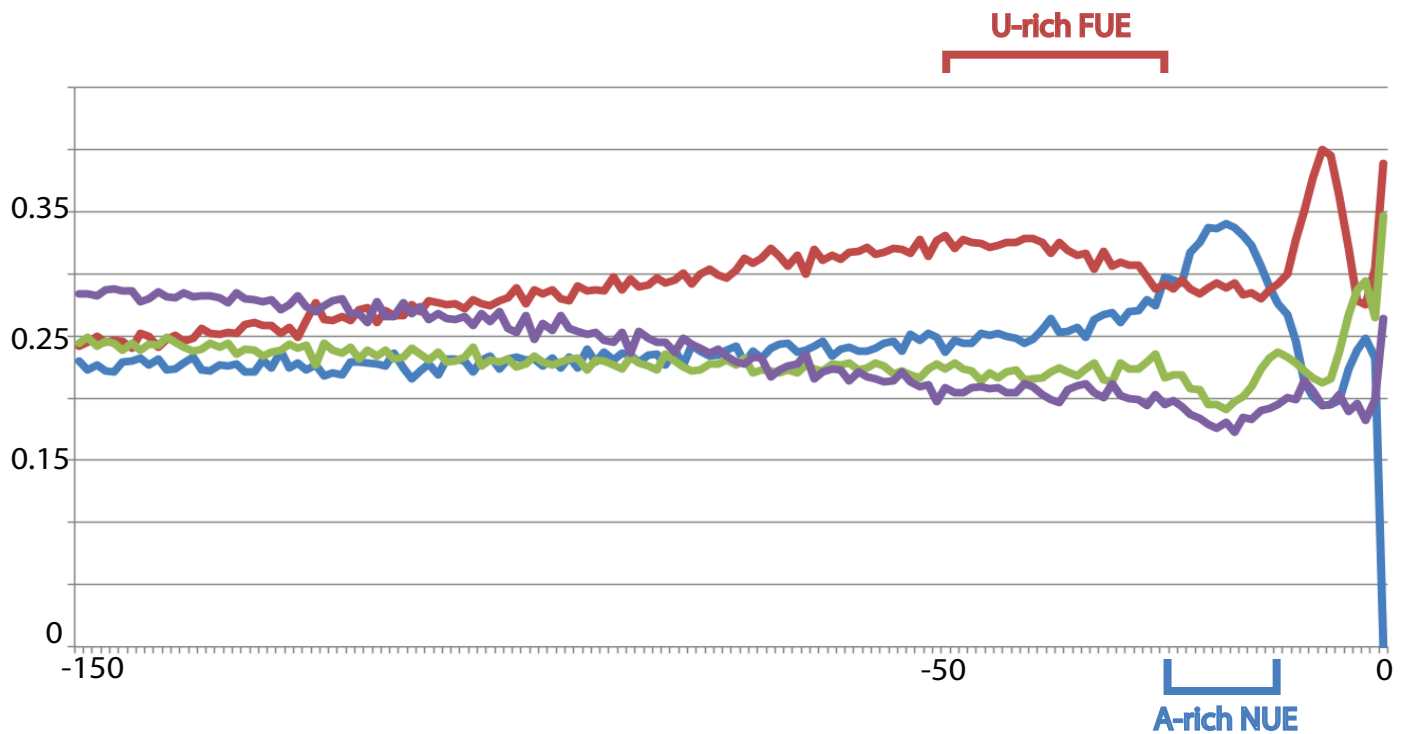


Figure B. Genome-wide analysis of the transcription termination signals in *P. crispa*. Average nucleotide composition was plotted for all non-polycistronic ToFU transcripts upstream of the poly-adenylation site. A-rich NUE and U-rich FUE elements are indicated. Green represents C-content and purple represent G-content, respectively. X-axis indicates the distance to polyA site.