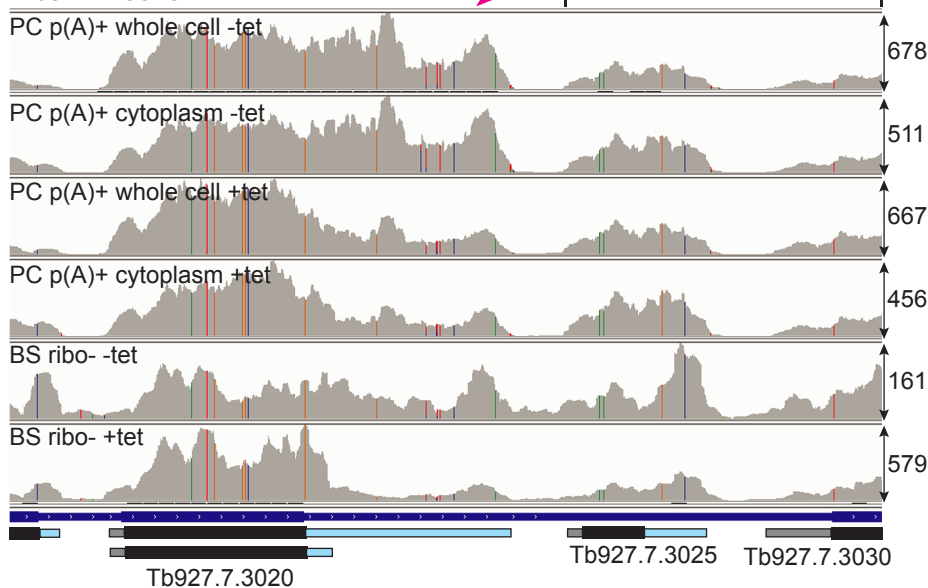
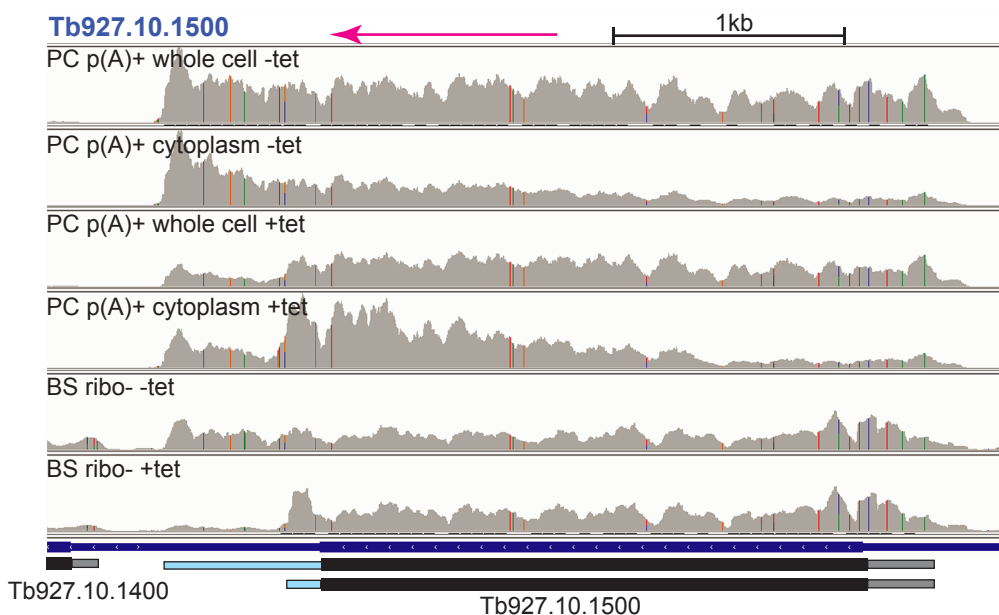


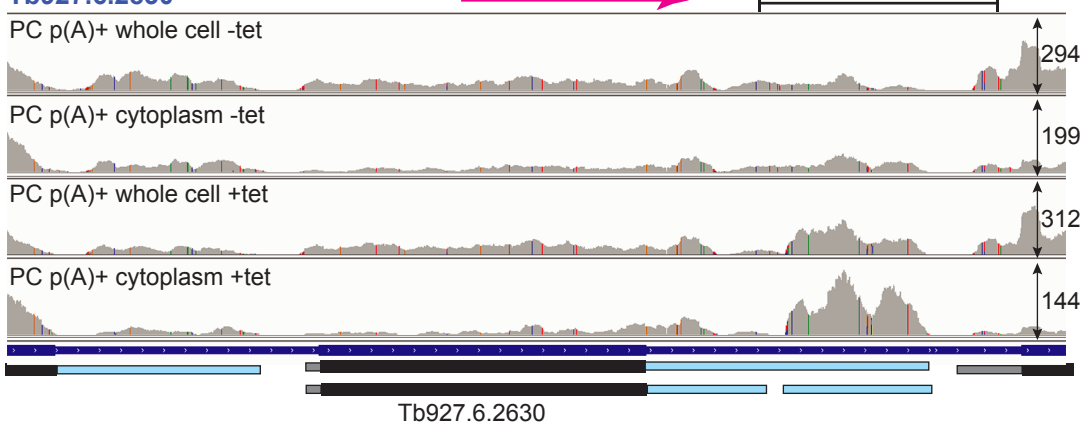
Tb927.7.3020



Tb927.10.1500



Tb927.6.2830



Supplementary Figure S7. Effects of DRBD18 depletion on mRNA processing: screen shots

The figure shows typical results for one of the three replicates for each type of sample, displayed on the Integrated Genomics Viewer. In the maps coding regions are black, 5'-UTRs are grey, and 3'-UTRs are pale cyan. The alternative truncated mRNAs are also indicated. The direction of transcription is indicated by a magenta arrow and the scale is indicated at the top. This Figure shows results for Tb927.7.3020, Tb927.10.1500 and Tb927.6.2630