

## CORRIGENDUM

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### **Corrigendum: UAP56/DDX39B is a major cotranscriptional RNA–DNA helicase that unwinds harmful R loops genome-wide**

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In the above-mentioned article, during the preparation of Supplemental Figure S7D for the final version of this article, two of the DRIPc-seq profiles—those of the siC and siUAP56 samples untreated with RNH1—were unintentionally plotted using a coverage bigwig file with a bin size different from the others. The amended version of this figure, in which the RNH-sensitive regions have been marked, can be found in the Revised Supplemental Material online. The revised plots correspond to the same original data and do not alter the conclusions of the paper. The authors apologize for this error.

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