RNA 21: 1219–1232 (2015)

Corrigendum: Conserved RNA secondary structures and long-range interactions in hepatitis C viruses

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In the above-noted article, the Acknowledgments section was incomplete and should instead read as follows:

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This article has been corrected in both the PDF and full-text HTML files online.

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Corrigendum: Long-range RNA pairings contribute to mutually exclusive splicing

YUAN YUE, YUN YANG, LANZHI DAI, GUOZHENG CAO, RAN CHEN, WEILING HONG, BAOPING LIU, YANG SHI, YIJUN MENG, FENG SHI, MU XIAO, and YONGFENG JIN

In the Discussion section (first paragraph, third sentence) on page 105 of the above-noted article, the sentence "The RNA architectures identified in our study were not predicted by in silico approaches (Pervouchine 2014), nor were they detected using recent high-throughput RNA structure sequencing techniques (Daines et al. 2011; Li et al. 2012)" was not clear and should instead read as follows: "The RNA bidirectional structural architectures identified in our study were not detected using recent high-throughput RNA structure sequencing techniques (Daines et al. 2011; Li et al. 2012), nor were they discovered by in silico approaches, although stem II of the *Drosophila srp* pre-mRNA was predicted (Raker et al. 2009; Pervouchine 2014)."

Accordingly, the following reference should be included in the References section:

Raker VA, Mironov AA, Gelfand MS, Pervouchine DD. 2009. Modulation of alternative splicing by long-range RNA structures in *Drosophila*. *Nucleic Acids Res* **37**: 4533–4544.

The authors apologize for any confusion this may have caused.

This article has been corrected in both the PDF and full-text HTML files online.

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