

## Corrigenda

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**Genome Research 25:** 655–666 (2015)

**Corrigendum: Extensive microRNA-mediated crosstalk between lncRNAs and mRNAs in mouse embryonic stem cells**

Jennifer Y. Tan, Tamara Sirey, Frantisek Honti, Bryony Graham, Allison Piovesan, Matthias Merckenschlager, Caleb Webber, Chris P. Ponting, and Ana C. Marques

In the above-mentioned article, one result reported in the third paragraph of the Results subsection “ceRNAs of individual lncRNAs tend to be functionally related” was in error. This paragraph should now read: “On average, mESC-expressed lncRNAs have 20.2 predicted MREs per kb of transcript that are specific to 12 different mESC-expressed miRNAs. This MRE density is similar to the density within 3′ UTRs of ceRNAs (20.4 MREs predicted per kb;  $P=0.34$ , two-tailed Mann-Whitney  $U$  test) (Supplemental Fig. S9; Supplemental Table S8). A single lncRNA might, therefore, be as likely as an mRNA to regulate post-transcriptionally the transcript abundance of many mRNAs via crosstalk with many miRNAs.” Corrected versions of Supplemental Figure S9 and Supplemental Table S8 are also now provided online.

This correction does not affect any of the conclusions of the paper. The authors apologize for any confusion caused by this error.

doi: 10.1101/gr.196568.115

**Genome Research 25:** 1018–1029 (2015)

**Corrigendum: Unraveling determinants of transcription factor binding outside the core binding site**

Michal Levo, Einat Zalckvar, Eilon Sharon, Ana Carolina Dantas Machado, Yael Kalma, Maya Lotan-Pompan, Adina Weinberger, Zohar Yakhini, Remo Rohs, and Eran Segal

The name of the sixth author was originally misspelled in the author line of this article. Please note the correct spelling as Maya Lotan-Pompan. The file has already been corrected online.

doi: 10.1101/gr.196725.115



## Corrigendum: Extensive microRNA-mediated crosstalk between lncRNAs and mRNAs in mouse embryonic stem cells

Jennifer Y. Tan, Tamara Sirey, Frantisek Honti, et al.

*Genome Res.* 2015 25: 1410.1

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Jennifer Y. Tan, Tamara Sirey, Frantisek Honti, et al.  
[Genome Res. May , 2015 25: 655-666](#)

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