



Corrigendum: Evaluation of strategies for evidence-driven genome annotation using long-read RNA-seq

Alejandro Paniagua, Cristina Agustín-García, Francisco J. Pardo-Palacios, et al.

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Corrigenda

Genome Research 36: 1108–1124 (2026)

Corrigendum: Characterizing cytosine methylation of polymorphic transposable element insertions using the human pangenome resources

Xiaoyu Zhuo, Chad Tomlinson, Edward A. Belter Jr, Prashant Kumar Kuntala, Wesley N. Saintilnord, Juan Jiang, Tina Lindsay, Juan F. Macias-Velasco, Robert S. Fulton, Human Pangenome Reference Consortium, and Ting Wang

The name of coauthor Juan F. Macias-Velasco was incomplete in the original author line of this article.

The author list has been corrected in both the PDF and full-text HTML files online.

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Corrigendum: Evaluation of strategies for evidence-driven genome annotation using long-read RNA-seq

Alejandro Paniagua, Cristina Agustín-García, Francisco J. Pardo-Palacios, Thomas Brown, Maite De Maria, Nancy D. Denslow, Camila J. Mazzoni, and Ana Conesa

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Genome Research 31: 890–899 (2021)

Corrigendum: Systematic evaluation of the effect of polyadenylation signal variants on the expression of disease-associated genes

Meng Chen, Ran Wei, Gang Wei, Mingqing Xu, Zhixi Su, Chen Zhao, and Ting Ni

The authors would like to correct an error in the Supplemental Materials in the initial publication of this article. The authors mistakenly used a duplicated image for the semiquantitative RT-PCR results for Supplemental Figures S3E and S4E. The authors have now replaced the panel in Supplemental Figure S3E with the correct image from the raw data. This can be found in the Revised Supplemental Material.

The authors confirm that this correction does not impact the overall conclusions or interpretation of the results presented in the article.

The authors apologize for any confusion this may have caused.

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