

## Corrigenda

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### ***Genome Research 25:* 27–40 (2015)**

#### **Corrigendum: H3K4me1 marks DNA regions hypomethylated during aging in human stem and differentiated cells**

Agustín F. Fernández, Gustavo F. Bayón, Rocío G. Urdinguio, Estela G. Toraño, María G. García, Antonella Carella, Sandra Petrus-Reurer, Cecilia Ferrero, Pablo Martinez-Camblor, Isabel Cubillo, Javier García-Castro, Jesús Delgado-Calle, Flor M. Pérez-Campo, José A. Riancho, Clara Bueno, Pablo Menéndez, Anouk Mentink, Katia Mareschi, Fabian Claire, Corrado Fagnani, Emanuela Medda, Virgilia Toccaceli, Sonia Brescianini, Sebastián Moran, Manel Esteller, Alexandra Stolzing, Jan de Boer, Lorenza Nisticò, Maria A. Stazi, and Mario F. Fraga

The authors would like to correct errors in sample labeling in Supplemental Table S1, which were inadvertently mislabeled in the initial publication of this article. The labels in the Sample ID column have now been corrected online in the Revised Supplemental Material.

The authors apologize for any confusion this might have caused.

doi: 10.1101/gr.249417.119

### ***Genome Research 27:* 362–373 (2017)**

#### **Corrigendum: Initiation of mtDNA transcription is followed by pausing, and diverges across human cell types and during evolution**

Amit Blumberg, Edward J. Rice, Anshul Kundaje, Charles G. Danko, and Dan Mishmar

The authors would like to correct an error in the labeling of the last two columns of Table 1 (Heavy strand and Light strand), which were inadvertently reversed in the initial publication of this article. The paper has been updated online to reflect the corrected table.

The authors apologize for any confusion this might have caused.

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## Corrigendum: H3K4me1 marks DNA regions hypomethylated during aging in human stem and differentiated cells

Agustín F. Fernández, Gustavo F. Bayón, Rocío G. Urduño, et al.

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[Genome Res. January , 2015 25: 27-40](#)

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