



Corrigendum: A systematic comparison reveals substantial differences in chromosomal versus episomal encoding of enhancer activity

Fumitaka Inoue, Martin Kircher, Beth Martin, et al.

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Corrigenda

Genome Research 28: 256–265 (2018)

Corrigendum: Detecting differential copy number variation between groups of samples

Craig B. Lowe, Nicelio Sanchez-Luege, Timothy R. Howes, Shannon D. Brady, Rhea R. Daugherty, Felicity C. Jones, Michael A. Bell, and David M. Kingsley

Due to a formatting error, Supplemental Table S4 in the above-mentioned article displayed incorrect gene identifiers as the closest transcription start site in column E. The revised Supplemental Table S4 has been posted online to reflect the correct gene identifiers. This error does not affect the analyses performed in this paper. The authors apologize for the inconvenience.

doi: 10.1101/gr.237370.118

Genome Research 28: 145–158 (2018)

Corrigendum: Evolutionary expansion of DNA hypomethylation in the mammalian germline genome

Jianghan Qu, Emily Hodges, Antoine Molaro, Pascal Gagneux, Matthew D. Dean, Gregory J. Hannon, and Andrew D. Smith

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“This research was partially supported by National Institutes of Health grant R01 HG005238 to A.D.S. and National Institutes of Health grant R01 GM098536 to M.D.D.”

The authors apologize for the inconvenience.

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Genome Research 27: 38–52 (2017)

Corrigendum: A systematic comparison reveals substantial differences in chromosomal versus episomal encoding of enhancer activity

Fumitaka Inoue, Martin Kircher, Beth Martin, Gregory M. Cooper, Daniela M. Witten, Michael T. McManus, Nadav Ahituv, and Jay Shendure

The authors would like to correct the omission of Supplemental Files 1, 2, and 3 in the initial publication of this article. Please note that the Supplemental files have now been added to Supplemental Material online. The authors apologize for any confusion this may have caused.

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