








## Author Correction: Predicting the clinical impact of human mutation with deep neural networks






Lakshman Sundaram , Hong Gao, Samskruthi Reddy Padigepati , Jeremy F. McRae , Yanjun Li , Jack A. Kosmicki, Nondas Fritzilas, Jörg Hakenberg , Anindita Dutta, John Shon, Jinbo Xu, Serafim Batzoglou, Xiaolin Li  and Kyle Kai-How Farh 

Correction to: *Nature Genetics* <https://doi.org/10.1038/s41588-018-0167-z>, published online 23 July 2018.

In the version of this article originally published, the name of author Serafim Batzoglou was misspelled. The error has been corrected in the HTML and PDF versions of the article.

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<https://doi.org/10.1038/s41588-018-0329-z>

## Author Correction: Assembly of a pan-genome from deep sequencing of 910 humans of African descent





Rachel M. Sherman , Juliet Forman, Valentin Antonescu, Daniela Puiu, Michelle Daya, Nicholas Rafaels, Meher Preethi Boorgula, Sameer Chavan, Candelaria Vergara , Victor E. Ortega, Albert M. Levin, Celeste Eng, Maria Yazdanbakhsh , James G. Wilson, Javier Marrugo, Leslie A. Lange, L. Keoki Williams, Harold Watson, Lorraine B. Ware, Christopher O. Olopade, Olufunmilayo Olopade, Ricardo R. Oliveira, Carole Ober, Dan L. Nicolae, Deborah A. Meyers, Alvaro Mayorga, Jennifer Knight-Madden, Tina Hartert, Nadia N. Hansel, Marilyn G. Foreman, Jean G. Ford, Mezbah U. Faruque, Georgia M. Dunston, Luis Caraballo, Esteban G. Burchard, Eugene R. Bleecker, Maria I. Araujo, Edwin F. Herrera-Paz , Monica Campbell, Cassandra Foster, Margaret A. Taub, Terri H. Beaty , Ingo Ruczinski, Rasika A. Mathias, Kathleen C. Barnes and Steven L. Salzberg 

Correction to: *Nature Genetics* <https://doi.org/10.1038/s41588-018-0273-y>, published online 19 November 2018.

In the version of this article initially published, the statement “there are no pan-genomes for any other animal or plant species” was incorrect. The statement has been corrected to “there are no reported pan-genomes for any other animal species, to our knowledge.” We thank David Edwards for bringing this error to our attention. The error has been corrected in the HTML and PDF versions of the article.

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## Publisher Correction: Immune genes are primed for robust transcription by proximal long noncoding RNAs located in nuclear compartments

Stephanie Fanucchi, Ezio T. Fok, Emiliano Dalla, Youtaro Shibayama, Kathleen Börner, Erin Y. Chang, Stoyan Stoychev, Maxim Imakaev , Dirk Grimm, Kevin C. Wang , Guoliang Li , Wing-Kin Sung and Musa M. Mhlanga 

Correction to: *Nature Genetics* <https://doi.org/10.1038/s41588-018-0298-2>, published online 10 December 2018

In the version of this article initially published, ‘+’ and ‘-’ labels were missing from the graph keys at the bottom of Fig. 8d. The error has been corrected in the HTML and PDF versions of the article.

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