













Author Correction: Genome-wide association study of intracranial aneurysms identifies 17 risk loci and genetic overlap with clinical risk factors

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Correction to: *Nature Genetics* <https://doi.org/10.1038/s41588-020-00725-7>, published online 16 November 2020.

In the version of this article initially published, the following statement was missing from the Acknowledgements: “We are grateful to the GenoBiRD core facility (Biogenouest), the Clinical Investigation Center (INSERM CIC1413) and the Center of Biological Resources in Nantes (BB-0033-00040; CHU Nantes, France) for their assistance in managing and genotyping the ICAN and PREGO biobanks. R.R. was supported by the French Regional Council of Pays-de-la-Loire (VaCaRMe program) and the Agence Nationale de la Recherche (ANR-15-CE17-0008-01 to G.L.). H.D. and R.B. were supported by the French Ministry of Health (clinical trial NCT02848495 to H.D.), the Genavie Foundation, the Société Française de Radiologie and the Société Française de Neuroradiologie.” The error has been corrected in the HTML and PDF versions of the article.





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Author Correction: Exploring the structural distribution of genetic variation in SARS-CoV-2 with the COVID-3D online resource

Stephanie Portelli , Moshe Olshansky, Carlos H. M. Rodrigues , Elston N. D’Souza , Yoochan Myung, Michael Silk, Azadeh Alavi, Douglas E. V. Pires and David B. Ascher 

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The version of this article initially published contained statements that data obtained through GISAID were used. No data used in the COVID-3D online resource were accessed through GISAID. Accordingly, mentions of GISAID in Supplementary Fig. 2 and the Supplementary Methods have been removed; the reference to the GISAID resource has been removed from the main text; and “variants detected in more than 125,000 SARS-CoV-2 genomic sequences” has been changed to “more than 11,000 variants detected in circulating SARS-CoV-2 genomic sequences.” The errors have been corrected in the HTML and PDF versions of the article.

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