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Correction to: *Nature Genetics* <https://doi.org/10.1038/s41588-017-0011-x>, published online 22 December 2017.

In the HTML version of this article initially published, the author groups ‘CHD Exome+ Consortium’, ‘EPIC-CVD Consortium’, ‘ExomeBP Consortium’, ‘Global Lipids Genetic Consortium’, ‘GoT2D Genes Consortium’, ‘EPIC InterAct Consortium’, ‘INTERVAL Study’, ‘ReproGen Consortium’, ‘T2D-Genes Consortium’, ‘The MAGIC Investigators’ and ‘Understanding Society Scientific Group’ appeared at the end of the author list but should have appeared earlier in the list, after author Krina T. Zondervan. The errors have been corrected in the HTML version of the article.

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<https://doi.org/10.1038/s41588-019-0447-2>

Publisher Correction: BRD4 interacts with NIPBL and BRD4 is mutated in a Cornelia de Lange-like syndrome

Gabrielle Olley, Morad Ansari, Hemant Bengani, Graeme R. Grimes, James Rhodes, Alex von Kriegsheim, Ana Blatnik, Fiona J. Stewart, Emma Wakeling, Nicola Carroll, Alison Ross, Soo-Mi Park, Deciphering Developmental Disorders Study, Wendy A. Bickmore , Madapura M. Pradeepa  and David R. FitzPatrick 

Correction to: *Nature Genetics* <https://doi.org/10.1038/s41588-018-0042-y>, published online 29 January 2018.

In the HTML version of this article initially published, the author group ‘Deciphering Developmental Disorders Study’ appeared at the end of the author list but should have appeared in the fourth-to-last position. The error has been corrected in the HTML version of the article.

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Publisher Correction: Multiancestry genome-wide association study of 520,000 subjects identifies 32 loci associated with stroke and stroke subtypes

Rainer Malik, Ganesh Chauhan, Matthew Traylor , Muralidharan Sargurupremraj, Yukinori Okada , Aniket Mishra, Loes Rutten-Jacobs , Anne-Katrin Giese, Sander W. van der Laan , Solveig Gretarsdottir, Christopher D. Anderson, Michael Chong, Hieab H. H. Adams , Tetsuro Ago, Peter Almgren , Philippe Amouyel , Hakan Ay, Traci M. Bartz, Oscar R. Benavente, Steve Bevan, Giorgio B. Boncoraglio, Robert D. Brown Jr, Adam S. Butterworth, Caty Carrera, Cara L. Carty, Daniel I. Chasman, Wei-Min Chen, John W. Cole, Adolfo Correa , Ioana Cotlarciuc, Carlos Cruchaga , John Danesh, Paul I. W. de Bakker, Anita L. DeStefano, Marcel den Hoed , Qing Duan, Stefan T. Engelter, Guido J. Falcone, Rebecca F. Gottesman, Raji P. Grewal, Vilmundur Gudnason, Stefan Gustafsson , Jeffrey Haessler, Tamara B. Harris, Ahamad Hassan, Aki S. Havulinna , Susan R. Heckbert, Elizabeth G. Holliday, George Howard, Fang-Chi Hsu, Hyacinth I. Hyacinth, M. Arfan Ikram , Erik Ingelsson, Marguerite R. Irvin, Xueqiu Jian, Jordi Jiménez-Conde, Julie A. Johnson, J. Wouter Jukema , Masahiro Kanai , Keith L. Keene, Brett M. Kissela, Dawn O. Kleindorfer, Charles Kooperberg, Michiaki Kubo, Leslie A. Lange, Carl D. Langefeld, Claudia Langenberg, Lenore J. Launer, Jin-Moo Lee, Robin Lemmens, Didier Leys, Cathryn M. Lewis , Wei-Yu Lin 

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Correction to: *Nature Genetics* <https://doi.org/10.1038/s41588-018-0058-3>, published online 12 March 2018.

In the HTML version of this article initially published, the author groups 'AFGen Consortium', 'Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) Consortium', 'International Genomics of Blood Pressure (iGEN-BP) Consortium', 'INVENT Consortium', 'STARNET', 'BioBank Japan Cooperative Hospital Group', 'COMPASS Consortium', 'EPIC-CVD Consortium', 'EPIC-InterAct Consortium', 'International Stroke Genetics Consortium (ISGC)', 'METASTROKE Consortium', 'Neurology Working Group of the CHARGE Consortium', 'NINDS Stroke Genetics Network (SiGN)', 'UK Young Lacunar DNA Study' and 'MEGASTROKE Consortium' appeared at the end of the author list but should have appeared earlier in the list. In addition, the author group 'MEGASTROKE Consortium' was duplicated, and its members were not displayed in the 'Author information' section. The errors have been corrected in the HTML version of the article.

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Publisher Correction: Common schizophrenia alleles are enriched in mutation-intolerant genes and in regions under strong background selection

Antonio F. Pardiñas , Peter Holmans, Andrew J. Pocklington, Valentina Escott-Price, Stephan Ripke , Noa Carrera, Sophie E. Legge , Sophie Bishop, Darren Cameron , Marian L. Hamshere, Jun Han, Leon Hubbard, Amy Lynham , Kiran Mantripragada, Elliott Rees, James H. MacCabe , Steven A. McCarroll, Bernhard T. Baune , Gerome Breen , Enda M. Byrne , Udo Dannlowski, Thalia C. Eley , Caroline Hayward , Nicholas G. Martin, Andrew M. McIntosh , Robert Plomin, David J. Porteous , Naomi R. Wray , Armando Caballero , Daniel H. Geschwind, Laura M. Huckins , Douglas M. Ruderfer, Enrique Santiago , Pamela Sklar, Eli A. Stahl, Hyejung Won , Esben Agerbo , Thomas D. Als , Ole A. Andreassen, Marie Bækvad-Hansen, Preben Bo Mortensen, Carsten Bøcker Pedersen, Anders D. Børjglum, Jonas Bybjerg-Grauholm , Srdjan Djurovic, Naser Durmishi, Marianne Giørtz Pedersen, Vera Golimbet, Jakob Grove , David M. Hougaard , Manuel Mattheisen, Espen Molden , Ole Mors, Merete Nordentoft, Milica Pejovic-Milovancevic, Engilbert Sigurdsson , Teimuraz Silagadze, Christine Søholm Hansen , Kari Stefansson, Hreinn Stefansson, Stacy Steinberg, Sarah Tosato , Thomas Werge, GERAD1 Consortium, CRESTAR Consortium, David A. Collier, Dan Rujescu, George Kirov, Michael J. Owen , Michael C. O'Donovan  and James T. R. Walters 

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