

Corrigendum: Genome-wide association study identifies multiple susceptibility loci for pulmonary fibrosis

Tasha E Fingerlin, Elissa Murphy, Weiming Zhang, Anna L Peljto, Kevin K Brown, Mark P Steele, James E Loyd, Gregory P Cosgrove, David Lynch, Steve Groshong, Harold R Collard, Paul J Wolters, Williamson Z Bradford, Karl Kossen, Scott D Seiwert, Roland M du Bois, Christine Kim Garcia, Megan S Devine, Gunnar Gudmundsson, Helgi J Isaksson, Naftali Kaminski, Yingze Zhang, Kevin F Gibson, Lisa H Lancaster, Joy D Cogan, Wendi R Mason, Toby M Maher, Philip L Molyneaux, Athol U Wells, Miriam F Moffatt, Moises Selman, Annie Pardo, Dong Soon Kim, James D Crapo, Barry J Make, Elizabeth A Regan, Dinesha S Walek, Jerry J Daniel, Yoichiro Kamatani, Diana Zelenika, Keith Smith, David McKean, Brent S Pedersen, Janet Talbert, Raven N Kidd, Cheryl R Markin, Kenneth B Beckman, Mark Lathrop, Marvin I Schwarz & David A Schwartz

Nat. Genet. 45, 613–620 (2013); published online 14 April 2013; corrected after print 1 October 2013

In the version of this article initially published, the minor alleles for two SNPs (rs2076295 and rs7934606) were listed incorrectly in Table 1. The correct minor allele for rs2076295 is G, and the correct minor allele for rs7934606 is T. As a result, there were two incorrect statements in the text regarding the relationship between the disease risk allele of rs2076295 and *DSP* expression. The text should have stated that the allele of rs2076295 associated with increased risk of pulmonary fibrosis is associated with decreased *DSP* expression rather than increased *DSP* expression. These errors have been corrected in the HTML and PDF versions of the article.

Corrigendum: Common variants at *SCN5A-SCN10A* and *HEY2* are associated with Brugada syndrome, a rare disease with high risk of sudden cardiac death

Connie R Bezzina, Julien Barc, Yuka Mizusawa, Carol Ann Remme, Jean-Baptiste Gourraud, Floriane Simonet, Arie O Verkerk, Peter J Schwartz, Lia Crotti, Federica Dagradi, Pascale Guicheney, Véronique Fressart, Antoine Leenhardt, Charles Antzelevitch, Susan Bartkowiak, Eric Schulze-Bahr, Sven Zumhagen, Elijah R Behr, Rachel Bastiaenen, Jacob Tfelt-Hansen, Morten Salling Olesen, Stefan Kääh, Britt M Beckmann, Peter Weeke, Hiroshi Watanabe, Naoto Endo, Tohru Minamino, Minoru Horie, Seiko Ohno, Kanae Hasegawa, Naomasa Makita, Akihiko Nogami, Wataru Shimizu, Takeshi Aiba, Philippe Froguel, Beverley Balkau, Olivier Lantieri, Margherita Torchio, Cornelia Wiese, David Weber, Rianne Wolswinkel, Ruben Coronel, Bas J Boukens, Stéphane Bézieau, Eric Charpentier, Stéphanie Chatel, Aurore Despres, Françoise Gros, Florence Kyndt, Simon Lecointe, Pierre Lindenbaum, Vincent Portero, Jade Violleau, Manfred Gessler, Hanno L Tan, Dan M Roden, Vincent M Christoffels, Hervé Le Marec, Arthur A Wilde, Vincent Probst, Jean-Jacques Schott, Christian Dina & Richard Redon

Nat. Genet. 45, 1044–1049 (2013); published online 21 July 2013; corrected after print 4 October 2013

In the version of this article initially published, Martin Borggrefe and Rainer Schimpf were inadvertently omitted from the author list. Both are affiliated with the First Department of Medicine (Cardiology), University Medical Center, Medical Faculty Mannheim, Heidelberg University, Mannheim, Germany, and with DZHK (German Centre for Cardiovascular Research), partner site Heidelberg/Mannheim, Mannheim, Germany. In addition, one of the study's funding sources (the Ministry of Health, Labour and Welfare of Japan, research grant for cardiovascular diseases, H24-033) was omitted from the Acknowledgments section. These errors have been corrected in the HTML and PDF versions of the article.