

Corrigendum: Genome-wide association analyses identifies a susceptibility locus for tuberculosis on chromosome 18q11.2

Thorsten Thye, Fredrik O Vannberg, Sunny H Wong, Ellis Owusu-Dabo, Ivy Osei, John Gyapong, Giorgio Sirugo, Fatou Sisay-Joof, Anthony Enimil, Margaret A Chinbuah, Sian Floyd, David K Warndorff, Lifted Sichali, Simon Malema, Amelia C Crampin, Bagrey Ngwira, Yik Y Teo, Kerrin Small, Kirk Rockett, Dominic Kwiatkowski, Paul E Fine, Philip C Hill, Melanie Newport, Christian Lienhardt, Richard A Adegbola, Tumani Corrah, Andreas Ziegler, Andrew P Morris, Christian G Meyer, Rolf D Horstmann, Adrian V S Hill, African TB Genetics Consortium & The Wellcome Trust Case Control Consortium
Nat. Genet. 42, 739–741 (2010); published online 8 August 2010; corrected after print 22 September 2011

In the version of this article initially published, an acknowledgment was missing. The study was supported by funding from the NIHR Oxford Biomedical Research Centre programme. The error has been corrected in the HTML and PDF versions of the article.

Erratum: Common variants in *P2RY11* are associated with narcolepsy

Birgitte R Kornum, Minae Kawashima, Juliette Faraco, Ling Lin, Thomas J Rico, Stephanie Hesselson, Robert C Axtell, Hedwich Kuipers, Karin Weiner, Alexandra Hamacher, Matthias U Kassack, Fang Han, Stine Knudsen, Jing Li, Xiaosong Dong, Juliane Winkelmann, Giuseppe Plazzi, Sona Nevsimalova, Seung-Chul Hong, Yutaka Honda, Makoto Honda, Birgit Högl, Thanh G N Ton, Jacques Montplaisir, Patrice Bourgin, David Kemlink, Yu-Shu Huang, Simon Warby, Mali Einen, Jasmin L Eshragh, Taku Miyagawa, Alex Desautels, Elisabeth Ruppert, Per Egil Hesla, Francesca Poli, Fabio Pizza, Birgit Frauscher, Jong-Hyun Jeong, Sung-Pil Lee, Kingman P Strohl, William T Longstreth Jr, Mark Kvale, Marie Dobrovolna, Maurice M Ohayon, Gerald T Nepom, H-Erich Wichmann, Guy A Rouleau, Christian Gieger, Douglas F Levinson, Pablo V Gejman, Thomas Meitinger, Paul Peppard, Terry Young, Poul Jennum, Lawrence Steinman, Katsushi Tokunaga, Pui-Yan Kwok, Neil Risch, Joachim Hallmayer & Emmanuel Mignot
Nat. Genet. 43, 66–71 (2011); published online 19 December 2010; corrected after print 22 September 2011

In the version of this article initially published, the percentage reduction of P2RY11 expression in CD8⁺ T lymphocytes was incorrectly given as 339%, when it should have been 72%. The percentage was not given for NK cells and should have been 70%. The errors have been corrected in the HTML and PDF versions of the article.

Corrigendum: Identification of an imprinted master *trans* regulator at the *KLF14* locus related to multiple metabolic phenotypes

Kerrin S Small, Åsa K Hedman, Elin Grundberg, Alexandra C Nica, Gudmar Thorleifsson, Augustine Kong, Unnur Thorsteindottir, So-Youn Shin, Hannah B Richards, the GIANT Consortium, the MAGIC Investigators, the DIAGRAM Consortium, Nicole Soranzo, Kourosh R Ahmadi, Cecilia M Lindgren, Kari Stefansson, Emmanouil T Dermizakis, Panos Deloukas, Timothy D Spector & Mark I McCarthy for the MuTHER Consortium
Nat. Genet. 43, 561–564 (2011); published online 15 May 2011; corrected after print 22 September 2011

In the version of this article initially published, there were several errors in the *P* values reported in the Adiponectin and HOMA-IR columns of Table 3. These errors have been corrected in the HTML and PDF versions of the article.