Corrigendum: Exome sequencing in sporadic autism spectrum disorders identifies severe *de novo* mutations

Brian J O'Roak, Pelagia Deriziotis, Choli Lee, Laura Vives, Jerrod J Schwartz, Santhosh Girirajan, Emre Karakoc, Alexandra P MacKenzie, Sarah B Ng, Carl Baker, Mark J Rieder, Deborah A Nickerson, Raphael Bernier, Simon E Fisher, Jay Shendure & Evan E Eichler
***Nat. Genet.* 43, 585–589 (2011); published online 15 May 2011; corrected after print 9 March 2012**

In the version of this article initially published, the missense variant in CNTNAP2 identified in proband 12817 was incorrectly listed as p.His275Ala in the main text and Figure 1. The correct notation for this variant is p.His275Arg. In addition, the raw sequence reads have now been deposited in the National Database for Autism Research under accession number NDARCOL0001878. These errors have been corrected in the HTML and PDF versions of the article.