

CORRECTION

Correction: Exome Sequencing in 53 Sporadic Cases of Schizophrenia Identifies 18 Putative Candidate Genes

Michel Guipponi, Federico A. Santoni, Vincent Setola, Corinne Gehrig, Maud Rotharmel, Macarena Cuenca, Olivier Guillin, Dimitris Dikeos, Georgios Georgantopoulos, George Papadimitriou, Logos Curtis, Alexandre Méary, Franck Schürhoff, Stéphane Jamain, Dimitri Avramopoulos, Marion Leboyer, Dan Rujescu, Ann Pulver, Dominique Campion, David P. Siderovski, Stylianos E. Antonarakis

Notice of Republication

Table S1 was published in error. The error was corrected in the HTML and PDF versions of this article on October 12, 2015, and the corrected table is available in the Supporting Information of this article as [S1 Table](#).

Supporting Information

S1 Table. Clinical data for each SCZ trio.
(DOCX)

Reference

1. Guipponi M, Santoni FA, Setola V, Gehrig C, Rotharmel M, Cuenca M, et al. (2014) Exome Sequencing in 53 Sporadic Cases of Schizophrenia Identifies 18 Putative Candidate Genes. PLoS ONE 9(11): e112745. doi:[10.1371/journal.pone.0112745](https://doi.org/10.1371/journal.pone.0112745) PMID: [25420024](https://pubmed.ncbi.nlm.nih.gov/25420024/)



OPEN ACCESS

Citation: Guipponi M, Santoni FA, Setola V, Gehrig C, Rotharmel M, Cuenca M, et al. (2015) Correction: Exome Sequencing in 53 Sporadic Cases of Schizophrenia Identifies 18 Putative Candidate Genes. PLoS ONE 10(10): e0141630. doi:[10.1371/journal.pone.0141630](https://doi.org/10.1371/journal.pone.0141630)

Published: October 21, 2015

Copyright: © 2015 Guipponi et al. This is an open access article distributed under the terms of the [Creative Commons Attribution License](https://creativecommons.org/licenses/by/4.0/), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.