SCIENTIFIC REPORTS

Published online: 04 December 2017

OPEN Author Correction: Antagonistic evolution of an antibiotic and its molecular chaperone: how to maintain a vital ectosymbiosis in a highly fluctuating habitat

Claire Papot¹, François Massol¹, Didier Jollivet² & Aurélie Tasiemski¹

Scientific Reports 7:1454; doi:10.1038/s41598-017-01626-2; Article published online 03 May 2017

Funding sources and the scientific contributions of B Shillito, J Ravaux and G Hamel were not acknowledged in this Article. The authors apologise for these accidental omissions. The Acknowledgements section,

"We thank the captain and crew of the RV Atalante, the DSV Nautile group (IFREMER), along with N. Le Bris (UMR8222) and F. Lallier (UMR7144), chief scientists of the MESCAL cruises. We thank A.S. Leport (UMR7144) for her technical assistance and B. Shillito (BOREA) and S. Hourdez (UMR7144) for the use of BALIST and DESEARES, respectively. We also acknowledge I. Probert (MBRC, Roscoff) for the English proofreading. This work was funded by the Université de Lille (BQR), the CNRS, the GDR ECCHIS and the Région Nord Pas de Calais-FRB (VERMER project)".

should read:

"We thank the captain and crew of the RV Atalante, the DSV Nautile group (IFREMER), along with N. Le Bris (UMR8222) and F. Lallier (UMR7144), chief scientists of the MESCAL cruises. We thank A.S. Leport (UMR7144) for her technical assistance and B. Shillito (BOREA) and S. Hourdez (UMR7144) for the use of BALIST and DESEARES, respectively. We also acknowledge I. Probert (MBRC, Roscoff) for the English proofreading. This work was funded by the Université de Lille (BQR), the CNRS, the GDR ECCHIS and the Région Nord Pas de Calais-FRB (VERMER project). We thank B Shillito, J Ravaux (UMR 7208 BOREA) and G Hamel (UMR 7590 IMPMC), for conducting and performing the BALIST pressure experiments, and for providing access to the corresponding samples. The isobaric sampling and transfer equipments used in this study were funded by the EXOCET/D project (FP6 program, FP6-GOCE-CT-2003-505342), the BALIST project (ANR program, ANR-08-BLAN-0252), and by the University Pierre et Marie Curie (BQR)".

In addition, the Author Contributions Statement is incorrect.

"C.P. generated and analyzed the sequence dataset. D.J. designed and contributed to the genetic analysis and performed the sampling with A.T. A.T. conducted and performed the experiments under pressure and the RT-qPCR. F.M. and C.P. participated in writing the paper. A.T. and D.J. supervised the work and wrote the paper".

should read:

"C.P. generated and analyzed the sequence dataset. D.J. designed and contributed to the genetic analysis and performed the sampling with A.T. F.M. and C.P. participated in writing the paper. A.T. and D.J. supervised the work and wrote the paper".

¹University Lille, CNRS, UMR 8198 - Evo-Eco-Paleo, SPICI group, F-59000, Lille, France. ²AD2M, ABICE team, Université Pierre et Marie Curie-CNRS, UMR7144, Station Biologique de Roscoff, 29682, Roscoff, France. Didier Jollivet and Aurélie Tasiemski contributed equally to this work. Correspondence and requests for materials should be addressed to A.T. (email: aurelie.tasiemski@univ-lille1.fr)

Open Access This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons license, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons license, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons license and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this license, visit http://creativecommons.org/licenses/by/4.0/.

© The Author(s) 2017