

CORRECTION

Correction: Cryptococcus neoformans resists to drastic conditions by switching to viable but non-culturable cell phenotype

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The Data Availability statement for this paper is incorrect. The correct statement is: All RNA sequence data from this study have been submitted to NCBI (<https://www.ncbi.nlm.nih.gov/geo>) under accession number GSE118549. The mass spectrometry proteomics data have been deposited to the ProteomeXchange Consortium via the PRIDE partner repository with the dataset identifier PXD012570 <http://proteomecentral.proteomexchange.org/cgi/GetDataset>.

Additionally, the authors inadvertently omit Dr. Caitlin Pepperell in their Acknowledgements section.

The correct Acknowledgements section should read: We would like to acknowledge Christina A. Cuomo, Laurent Châtre, Guilhem Janbon, Jean-Yves Coppée, Pierre Rocheteau, Pierre Henri Commere, Christine Schmitt, Olivier Gorgetto, Jacomine Krijnse-Locker, Caroline Proux, Rachel Legendre for their comments on this work and help at different steps of the study.

Reference

1. Hommel B, Sturny-Leclère A, Volant S, Veluppillai N, Duchateau M, Yu C-H, et al. (2019) *Cryptococcus neoformans* resists to drastic conditions by switching to viable but non-culturable cell phenotype. PLoS Pathog 15(7): e1007945. <https://doi.org/10.1371/journal.ppat.1007945> PMID: 31356623



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