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Corrigendum to 'the anti-fecundity effect of 5-azacytidine (5-AzaC) on *Schistosoma mansoni* is linked to dis-regulated transcription, translation and stem cell activities'

Geyer, Kathrin K.; Munshi, Sabrina E.; Vickers, Martin; Squance, Michael; Wilkinson, Toby; Berrar, Daniel; Chaparro, Cristian ; Swain, Martin; Hoffmann, Karl

Published in:

International Journal for Parasitology: Drugs and Drug Resistance

DOI:

[10.1016/j.ijpddr.2018.07.002](https://doi.org/10.1016/j.ijpddr.2018.07.002)

Publication date:

2018

Citation for published version (APA):

Geyer, K. K., Munshi, S. E., Vickers, M., Squance, M., Wilkinson, T., Berrar, D., Chaparro, C., Swain, M., & Hoffmann, K. (2018). Corrigendum to 'the anti-fecundity effect of 5-azacytidine (5-AzaC) on *Schistosoma mansoni* is linked to dis-regulated transcription, translation and stem cell activities': [Int. J. Parasitol. Drugs and Drug Resist. 8 (2018) 213–222]. *International Journal for Parasitology: Drugs and Drug Resistance*, 8(3), 493. <https://doi.org/10.1016/j.ijpddr.2018.07.002>

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Contents lists available at ScienceDirect

IJP: Drugs and Drug Resistance

journal homepage: www.elsevier.com/locate/ijpddr

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Kathrin K. Geyer^{a,1}, Sabrina E. Munshi^{a,1}, Martin Vickers^b, Michael Squance^a, Toby J. Wilkinson^c, Daniel Berrar^d, Cristian Chaparro^e, Martin T. Swain^a, Karl F. Hoffmann^{a,*}

^a Institute of Biological, Environmental and Rural Sciences (IBERS), Edward Lhwyd Building, Aberystwyth University, Aberystwyth, SY23 3DA, United Kingdom

^b Department of Cell and Developmental Biology, John Innes Centre, Norwich Research Park, Norwich, NR4 7UH, United Kingdom

^c The Roslin Institute, The University of Edinburgh, Easter Bush Campus, Midlothian, EH25 9RG, United Kingdom

^d Data Science Laboratory, Tokyo Institute of Technology, Tokyo, 152-8550, Japan

^e University of Perpignan Via Domitia, 58 Avenue Paul Alduy, Bat R, F-66860, Perpignan Cedex, France

The authors regret that the text describing the filtering of the RNA-Seq data contains an inaccuracy. We would like to correct this information in two places within the manuscript.

In the Materials and Methods section 2.5, 6th sentence, it should read: “Here, 4115 differentially expressed transcripts mapping to 4036 unique Smp gene identifiers were classified.”

In the Materials and methods section 2.6, 10th sentence, it should read: “To illustrate an overview of the genome locations and differential expression levels of the 4057 Smps (4115 transcripts) in the filtered RNA-Seq dataset, a Circos plot (Krzywinski et al., 2009) was generated using the *S. mansoni* genome v 5.2.”

The authors would like to apologise for any inconvenience caused.

DOI of original article: <https://doi.org/10.1016/j.ijpddr.2018.03.006>

* Corresponding author.

¹ Contributed equally to this work.

E-mail addresses: kkg6@aber.ac.uk (K.K. Geyer), sabrina.munshi@hotmail.co.uk (S.E. Munshi), martinj.vickers@gmail.com (M. Vickers), michaels@terravesta.com (M. Squance), toby.wilkinson@ed.ac.uk (T.J. Wilkinson), daniel.berrar@ict.e.titech.ac.jp (D. Berrar), Cristian.chaparro@univ-perp.fr (C. Chaparro), mts11@aber.ac.uk (M.T. Swain), krh@aber.ac.uk (K.F. Hoffmann).

<https://doi.org/10.1016/j.ijpddr.2018.07.002>

Available online 17 July 2018

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