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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'*

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*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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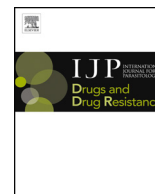
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Contents lists available at ScienceDirect

## IJP: Drugs and Drug Resistance

journal homepage: [www.elsevier.com/locate/ijpddr](http://www.elsevier.com/locate/ijpddr)

## 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]



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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>

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

Available online 17 July 2018

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