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# Author Correction: Using *de novo* transcriptome assembly and analysis to study RNAi in *Phenacoccus solenopsis* Tinsley (Hemiptera: Pseudococcidae)

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Correction to: *Scientific Reports* <https://doi.org/10.1038/s41598-019-49997-y>, published online 23 September 2019

This Article contains errors.

Reference 17 was incorrectly given as:

Verma, M., Ghangal, R., Sharma, R., Sinha, A. K. & Jain, M. Transcriptome analysis of *Catharanthus roseus* for gene discovery and expression profiling. *PLoS One*. <https://doi.org/10.1371/journal.pone.0103583> (2014).

The correct reference is listed below as ref. 1.

Additionally, a citation to reference 17 should be included in the Results and Discussion, where the sentence,

“Assembly of the available transcriptomes of *P. solenopsis* discovered the presence of 55198 transcripts in second instar (SRR6782025), 55569 transcripts in adult (SRR6782023), 49193 transcripts in third instar (SRR6782022) and 64497 in the eggs (SRR6782024)”

should read:

“Assembly of the available transcriptomes of *P. solenopsis* discovered the presence of 55198 transcripts in second instar (SRR6782025), 55569 transcripts in adult (SRR6782023), 49193 transcripts in third instar (SRR6782022) and 64497 in the eggs (SRR6782024)<sup>17</sup>”

## Reference

1. Arya, S. K. *et al.* *De novo* characterization of *Phenacoccus solenopsis* transcriptome and analysis of gene expression profiling during development and hormone biosynthesis. *Sci. Rep.* **8**, 7573, <https://doi.org/10.1038/s41598-018-25845-3> (2018).



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