

CORRECTION

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# Correction to: Custom selected reference genes outperform pre-defined reference genes in transcriptomic analysis

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**Correction to: BMC Genomics 21, 35 (2020)**  
<https://doi.org/10.1186/s12864-019-6426-2>

Following publication of the original article [1], the author reported that although they used the word “covariance” throughout the manuscript, the actual statistics used for selection of reference genes was the “coefficient of variation”.

Note that this does not affect the method, nor the scripts present in GitHub (<https://github.com/KarenGoncalves/CustomSelection>) to process the datasets.

The original article has been corrected.

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## Reference

1. dos Santos KCG, Desgagné-Penix I, Germain H. Custom selected reference genes outperform pre-defined reference genes in transcriptomic analysis. *BMC Genomics*. 2020;21(1):35. <https://doi.org/10.1186/s12864-019-6426-2>.

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The original article can be found online at <https://doi.org/10.1186/s12864-019-6426-2>.

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