



Erratum: Comparison of Normalization Methods for Analysis of TempO-Seq Targeted RNA Sequencing Data

Frontiers Production Office*

An Erratum on

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Comparison of Normalization Methods for Analysis of TempO-Seq Targeted RNA Sequencing Data

by Bushel, P. R., Ferguson, S. S., Ramaiahgari, S. C., Paules, R. S., and Auerbach, S. S. (2020). Front. Genet. 11:594. doi: 10.3389/fgene.2020.00594

Due to a production error, the equation for "Overall Accuracy" was typeset erroneously. The correct equation follows below:

 $Accuracy = \frac{(True Positives + True Negatives)}{(True Positives + True Negatives)}$

 $\frac{1}{(\text{True Positives} + \text{True Negatives} + \text{False Positives} + \text{False Negatives})}$

The publisher apologizes for this mistake. The original article has been updated.

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