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Erratum: An iteration normalization and test method for differential expression analysis of RNA-seq data

Yan Zhou^{1*}, Nan Lin² and Baoxue Zhang³

* Correspondence: zhouy1016@163.com

¹Institute for Genomic Biology, University of Illinois at Urbana-Champaign, Champaign, USA Full list of author information is available at the end of the article

Erratum

Following publication of our paper in *BioData Mining* [1], it came to light that numerous sections of text were similar or identical to a previous publication by Robinson and Oshlack [2]. In extending the method proposed by Robinson and Oshlack, we inadvertently copied text from their article [2]. We acknowledge that these areas of text should have been clearly indicated as quotations or excluded entirely. We sincerely apologize for this oversight.

Author details

¹Institute for Genomic Biology, University of Illinois at Urbana-Champaign, Champaign, USA. ²Department of Mathematics, Washington University in Saint Louis, St Louis, USA. ³Key Laboratory for Applied Statistics of MOE and School of Mathematics and Statistics, Northeast Normal University, Changchun, P. R. China.

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