Check for updates

OPEN ACCESS

EDITED AND REVIEWED BY Qi Zhao, University of Science and Technology Liaoning, China

*CORRESPONDENCE Yong Liang, yongliangresearch@gmail.com

SPECIALTY SECTION This article was submitted to Bioprocess Engineering, a section of the journal Frontiers in Bioengineering and Biotechnology

RECEIVED 29 July 2022 ACCEPTED 08 August 2022 PUBLISHED 30 August 2022

CITATION

Ouyang D, Miao R, Wang J, Liu X, Xie S, Ai N, Dang Q and Liang Y (2022), Corrigendum: Predicting multiple types of associations between miRNAs and diseases based on graph regularized weighted tensor decomposition. *Front. Bioeng. Biotechnol.* 10:1006237. doi: 10.3389/fbioe.2022.1006237

COPYRIGHT

© 2022 Ouyang, Miao, Wang, Liu, Xie, Ai, Dang and Liang. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or

reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.

Corrigendum: Predicting multiple types of associations between miRNAs and diseases based on graph regularized weighted tensor decomposition

Dong Ouyang¹, Rui Miao¹, Jianjun Wang², Xiaoying Liu³, Shengli Xie⁴, Ning Ai¹, Qi Dang¹ and Yong Liang⁵*

¹Faculty of Information Technology, Macau University of Science and Technology, Macau, China, ²School of Mathematics and Statistics, Southwest University, Chongqing, China, ³Computer Engineering Technical College, Guangdong Polytechnic of Science and Technology, Zhuhai, China, ⁴Institute of Intelligent Information Processing, Guangdong University of Technology, Guangzhou, China, ⁵Peng Cheng Laboratory, Shenzhen, China

KEYWORDS

multiple types of miRNA-disease associations, weighted tensor decomposition, graph Laplacian regularization, L2, 1 norm, multi-view biological similarity network

A Corrigendum on

Predicting multiple types of associations between miRNAs and diseases based on graph regularized weighted tensor decomposition

by Ouyang D, Miao R, Wang J, Liu X, Xie S, Ai N, Dang Q and Liang Y (2022). Front. Bioeng. Biotechnol. 10:911769. doi: 10.3389/fbioe.2022.911769

In the published article, there was an error in affiliation(s) **1**. Instead of "Faculty of Information Technology, Macau University of Science and Technology, Taipa, China," it should be "Faculty of Information Technology, Macau University of Science and Technology, Macau, China."

In the published article, there was an error. Mathematical symbols are inconsistent.

A correction has been made to **3 Methods**, "3.1 CP decomposition," Paragraph Number 5.

This sentence previously stated:

"CANDECOMP/PARAFAC (CP) decomposition is one of the most common tensor decomposition forms (Kolda and Bader, 2009). Given the miRNA-disease-type tensor $\mathcal{X} \in \mathbb{R}^{|m| \times |n| \times |t|}$, the CP decomposition model can be represented as follows:

$$\mathcal{X} \approx \sum_{s=1}^{S} m_s \circ d_s \circ t_s \equiv \llbracket M, D, T \rrbracket$$
(1)

where the symbol \circ represents the vector outer product, S is a positive integer and $\mathbf{m}_s \in \mathbb{R}^{|m| \times |1|}$, $\mathbf{d}_s \in \mathbb{R}^{|m| \times |1|}$ and $\mathbf{t}_s \in \mathbb{R}^{|t| \times |1|}$. $M = [\mathbf{m}_1 \mathbf{m}_2 \cdots \mathbf{m}_S], D = [\mathbf{d}_1 \mathbf{d}_2 \cdots \mathbf{d}_S]$, and $T = [\mathbf{t}_1 \mathbf{t}_2 \cdots \mathbf{t}_S]$ are the factor matrices with respect to different dimensions."

The corrected sentence appears below:

"CANDECOMP/PARAFAC (CP) decomposition is one of the most common tensor decomposition forms (Kolda and Bader, 2009). Given the miRNA-disease-type tensor $\mathcal{X} \in \mathbb{R}^{|m| \times |n| \times |t|}$, the CP decomposition model can be represented as follows:

$$\mathcal{X} \approx \sum_{s=1}^{S} \boldsymbol{m}_{s} \circ \boldsymbol{d}_{s} \circ \boldsymbol{t}_{s} \equiv [[M, D, T]]$$
(2)

where the symbol \circ represents the vector outer product, *S* is a positive integer and $m_s \in \mathbb{R}^{|m| \times 1}$, $d_s \in \mathbb{R}^{|n| \times 1}$ and $t_s \in \mathbb{R}^{|t| \times 1}$. $M = [m_1 \ m_2 \ \cdots \ m_S]$, $D = [d_1 \ d_2 \ \cdots \ d_S]$, and $T = [t_1 \ t_2 \ \cdots \ t_S]$ are the factor matrices with respect to different dimensions."

Reference

Kolda, T. G., and Bader, B. W. (2009). Tensor decompositions and applications. *SIAM Rev.* 51, 455–500. doi:10.1137/07070111X Note that mathematic symbols are bolded to represent vectors. Also, " $m_s \in \mathbb{R}^{|m| \times |1|}$, $d_s \in \mathbb{R}^{|n| \times |1|}$ and $t_s \in \mathbb{R}^{|t| \times |1|}$ " should be changed to " $m_s \in \mathbb{R}^{|m| \times 1}$, $d_s \in \mathbb{R}^{|n| \times 1}$ and $t_s \in \mathbb{R}^{|t| \times 1}$."

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.