

CORRECTION

Correction: Dissecting cancer heterogeneity based on dimension reduction of transcriptomic profiles using extreme learning machines

Kejun Wang, Xin Duan, Feng Gao, Wei Wang, Liangliang Liu, Xin Wang

An affiliation for the 6th author is not included. Xin Wang is affiliated with both #2 and #3, Shenzhen Research Institute, City University of Hong Kong, Shenzhen, China.

The following information is missing from the Funding section: The authors acknowledge the support received from Shenzhen Research Institute, City University of Hong Kong.

Reference

1. Wang K, Duan X, Gao F, Wang W, Liu L, Wang X (2018) Dissecting cancer heterogeneity based on dimension reduction of transcriptomic profiles using extreme learning machines. PLoS ONE 13(9): e0203824. <https://doi.org/10.1371/journal.pone.0203824> PMID: 30216380



OPEN ACCESS

Citation: Wang K, Duan X, Gao F, Wang W, Liu L, Wang X (2018) Correction: Dissecting cancer heterogeneity based on dimension reduction of transcriptomic profiles using extreme learning machines. PLoS ONE 13(10): e0205548. <https://doi.org/10.1371/journal.pone.0205548>

Published: October 5, 2018

Copyright: © 2018 Wang et al. This is an open access article distributed under the terms of the [Creative Commons Attribution License](#), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.