



Erratum: DeconPeaker, a Deconvolution Model to Identify Cell Types Based on Chromatin Accessibility in ATAC-Seq Data of Mixture Samples

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An Erratum on

DeconPeaker, a Deconvolution Model to Identify Cell Types Based on Chromatin Accessibility in ATAC-Seq Data of Mixture Samples

by Li, H., Sharma, A., Luo, K., Qin, Z. S., Sun, X., and Liu, H. (2020). *Front. Genet.* 11:392. doi: 10.3389/fgene.2020.00392

Due to a typesetting error, many of the references were cited in the wrong place. In addition the citations in the legends for Figures 2F and 7C were incorrect. In the former, “Jia et al.” should be “Jia et al. (2018)”. In the latter, “Corces et al.” should be “Corces et al. (2016)”. The publisher apologizes for this mistake. The original version of this article has been updated.

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