Correction to: Inference of Gene Regulatory Network from Single-Cell Transcriptomic Data Using pySCENIC

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Correction to:

Chapter 10 in: Shahid Mukhtar (ed.), *Modeling Transcriptional Regulation: Methods and Protocols*, Methods in Molecular Biology, vol. 2238, https://doi.org/10.1007/978-1-0716-1534-8_10

In the original version of this book, chapter 10 was published with incomplete list of citations. This has now been rectified in this revised version of the book.

The original version of chapter 10 was published without citing “SCENIC: single-cell regulatory network inference and clustering” by Aiber et al., *Nature Methods* 2017, its Python implementation “A scalable SCENIC workflow for single-cell gene regulatory network analysis” by Van de Sande et al., *Nature Protocols* 2020, GitHub repository (<https://github.com/aertslab/pySCENIC>), and the scRNA-seq dataset (GSE115978).

The updated online version of this chapter can be found at
https://doi.org/10.1007/978-1-0716-1534-8_10

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