

ERRATUM

Open Access



# Erratum to: gespeR: a statistical model for deconvoluting off-target-confounded RNA interference screens

Fabian Schmich<sup>1,2</sup>, Ewa Szczurek<sup>1,2</sup>, Saskia Kreibich<sup>3</sup>, Sabrina Dilling<sup>3</sup>, Daniel Andritschke<sup>3</sup>, Alain Casanova<sup>4</sup>, Shyan Huey Low<sup>4</sup>, Simone Eicher<sup>4</sup>, Simone Muntwiler<sup>4</sup>, Mario Emmenlauer<sup>4</sup>, Pauli Rämö<sup>4</sup>, Raquel Conde-Alvarez<sup>5</sup>, Christian von Mering<sup>2,6</sup>, Wolf-Dietrich Hardt<sup>3</sup>, Christoph Dehio<sup>4</sup> and Niko Beerenwinkel<sup>1,2\*</sup>

After the publication of this work [1], some content was missing from the Acknowledgements section. The corrected Acknowledgements section is given below:

## Acknowledgements

This work has been funded by SystemsX.ch, the Swiss Initiative in Systems Biology, under IPhD grant 2009/025 and RTD grants 51RT-0\_126008 (InfectX) and 51RTPO\_151029 (TargetInfectX), evaluated by the Swiss National Science Foundation. E.S. was supported by the ETH Zurich Postdoctoral Fellowship Program and the Marie Curie Actions for People COFUND program (grant FEL-13 12-1). We are grateful to Mihaela Zavolan for insightful discussions. The genome sequence used in this research was derived from a HeLa cell line [dpGAP:phs000640.v1.p1]. Henrietta Lacks, and the HeLa cell line that was established from her tumour cells without her knowledge or consent in 1951, have made significant contributions to scientific progress and advances in human health. We are grateful to Henrietta Lacks, now deceased, and to her surviving family members for their contributions to biomedical research. This study was reviewed by the NIH HeLa Genome Data Access Working Group.

## Author details

<sup>1</sup>Department of Biosystems Science and Engineering, ETH, Zurich, Switzerland. <sup>2</sup>SIB Swiss Institute of Bioinformatics, Basel, Switzerland. <sup>3</sup>Department of Biology, ETH, Zurich, Switzerland. <sup>4</sup>Biozentrum, University of Basel, Basel, Switzerland. <sup>5</sup>Institute for Tropical Health and Departamento de Microbiología y Parasitología, Universidad de Navarra, Pamplona, Spain. <sup>6</sup>Institute of Molecular Life Sciences, University of Zurich, Zurich, Switzerland.

Received: 16 October 2015 Accepted: 16 October 2015

Published online: 21 October 2015

## Reference

1. Schmich F, Szczurek E, Kreibich S, Dilling S, Andritschke D, Casanova A, et al. gespeR: a statistical model for deconvoluting off-target-confounded RNA interference screens. *Genome Biol.* 2015;16:220.

\* Correspondence: niko.beerenwinkel@bsse.ethz.ch

<sup>1</sup>Department of Biosystems Science and Engineering, ETH, Zurich, Switzerland

<sup>2</sup>SIB Swiss Institute of Bioinformatics, Basel, Switzerland

Full list of author information is available at the end of the article

Submit your next manuscript to BioMed Central and take full advantage of:

- Convenient online submission
- Thorough peer review
- No space constraints or color figure charges
- Immediate publication on acceptance
- Inclusion in PubMed, CAS, Scopus and Google Scholar
- Research which is freely available for redistribution

Submit your manuscript at  
[www.biomedcentral.com/submit](http://www.biomedcentral.com/submit)

