

ERRATUM

Open Access



# Erratum to: MSRE-HTPrimer: a high-throughput and genome-wide primer design pipeline optimized for epigenetic research

Ram Vinay Pandey<sup>1,2</sup>, Walter Pulverer<sup>1</sup>, Rainer Kallmeyer<sup>1</sup>, Gabriel Beikircher<sup>1</sup>, Stephan Pabinger<sup>1</sup>, Albert Kriegner<sup>1</sup> and Andreas Weinhäusel<sup>1\*</sup>

## Erratum

Unfortunately, the original version of this article [1] contained an error. The name of one of the authors was included incorrectly and it read “Pulverer Walter” instead of “Walter Pulverer”.

The name has been updated in the original article and is correctly included in full in this erratum.

## Author details

<sup>1</sup>Health & Environment Department, Molecular Diagnostics, AIT—Austrian Institute of Technology GmbH, Vienna, Austria. <sup>2</sup>Institut für Populationsgenetik, Vetmeduni Vienna, Veterinärplatz 1, 1210 Vienna, Austria.

Received: 29 July 2016 Accepted: 29 July 2016

Published online: 04 August 2016

## References

1. Pandey RV, Pulverer W, Kallmeyer R, Beikircher G, Pabinger S, Kriegner A, et al. MSRE-HTPrimer: a high-throughput and genome-wide primer design pipeline optimized for epigenetic research. *Clin Epigenetics*. 2016;8:26. doi: 10.1186/s13148-016-0190-9.

\* Correspondence: andreas.weinhaeusel@ait.ac.at

<sup>1</sup>Health & Environment Department, Molecular Diagnostics, AIT—Austrian Institute of Technology GmbH, Vienna, Austria

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

Submit your next manuscript to BioMed Central and we will help you at every step:

- We accept pre-submission inquiries
- Our selector tool helps you to find the most relevant journal
- We provide round the clock customer support
- Convenient online submission
- Thorough peer review
- Inclusion in PubMed and all major indexing services
- Maximum visibility for your research

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

