



Erratum to: Comparative performance of the BGISEQ-500 vs Illumina HiSeq2500 sequencing platforms for palaeogenomic sequencing

Mak, Sarah Siu Tze; Gopalakrishnan, Shyam; Carøe, Christian; Geng, Chunyu; Liu, Shanlin; Sinding, Mikkel-Holger S.; Kuderna, Lukas F. K.; Zhang, Wenwei; Fu, Shujin; Vieira, Filipe G.

Published in:
GigaScience

Link to article, DOI:
[10.1093/gigascience/giy151](https://doi.org/10.1093/gigascience/giy151)

Publication date:
2018

Document Version
Publisher's PDF, also known as Version of record

[Link back to DTU Orbit](#)

Citation (APA):

Mak, S. S. T., Gopalakrishnan, S., Carøe, C., Geng, C., Liu, S., Sinding, M-H. S., ... Gilbert, M. T. P. (2018). Erratum to: Comparative performance of the BGISEQ-500 vs Illumina HiSeq2500 sequencing platforms for palaeogenomic sequencing. *GigaScience*, 7(12). <https://doi.org/10.1093/gigascience/giy151>

General rights

Copyright and moral rights for the publications made accessible in the public portal are retained by the authors and/or other copyright owners and it is a condition of accessing publications that users recognise and abide by the legal requirements associated with these rights.

- Users may download and print one copy of any publication from the public portal for the purpose of private study or research.
- You may not further distribute the material or use it for any profit-making activity or commercial gain
- You may freely distribute the URL identifying the publication in the public portal

If you believe that this document breaches copyright please contact us providing details, and we will remove access to the work immediately and investigate your claim.

ERRATUM

Erratum to: Comparative performance of the BGISEQ-500 vs Illumina HiSeq2500 sequencing platforms for palaeogenomic sequencing

Sarah Siu Tze Mak^{1,†}, Shyam Gopalakrishnan^{1,†}, Christian Carøe^{1,2,†}, Chunyu Geng^{3,†}, Shanlin Liu^{1,4}, Mikkel-Holger S. Sinding^{1,5,6}, Lukas F.K. Kuderna^{7,8}, Wenwei Zhang³, Shujin Fu³, Filipe G. Vieira¹, Mietje Germonpré⁹, Hervé Bocherens^{10,11}, Sergey Fedorov¹², Bent Petersen², Thomas Sicheritz-Pontén², Tomas Marques-Bonet^{7,8,13}, Guojie Zhang^{4,14}, Hui Jiang^{3,*} and M. Thomas P. Gilbert^{1,15,16,*}

¹Centre for GeoGenetics, Natural History Museum of Denmark, University of Copenhagen, Øster Voldgade 5–7, 1350 Copenhagen, Denmark, ²DTU Bioinformatics, Department of Bio and Health Informatics, Technical University of Denmark, Building 208, DK-2800 Lyngby, Denmark, ³BGI-Shenzhen, Shenzhen 518083, China, ⁴China National GeneBank, BGI-Shenzhen, Shenzhen 518083, China, ⁵Natural History Museum, University of Oslo, PO Box 1172 Blindern, N-0318 Oslo, Norway, ⁶The Qimmeq Project, University of Greenland, Manutooq 1, PO Box 1061, 3905 Nuussuaq, Greenland, ⁷Institute of Evolutionary Biology (UPF-CSIC), PRBB, Dr. Aiguader 88, 08003 Barcelona, Spain, ⁸CNAG-CRG, Centre for Genomic Regulation (CRG), Barcelona Institute of Science and Technology (BIST), Baldri i Reixac 4, 08028 Barcelona, Spain, ⁹OD Earth and History of Life, Royal Belgian Institute of Natural Sciences, Vautierstraat 29, 1000 Brussels, Belgium, ¹⁰Department of Geosciences, Palaeobiology, University of Tübingen, Tübingen, Germany, ¹¹Senckenberg Centre for Human Evolution and Palaeoenvironment, University of Tübingen, Tübingen, Germany, ¹²Mammoth Museum, Institute of Applied Ecology of the North of the North-Eastern Federal University, ul. Kulakovskogo 48, 677980 Yakutsk, Russia, ¹³Catalan Institution of Research and Advanced Studies (ICREA), Passeig de Lluís Companys, 23, 08010, Barcelona, Spain, ¹⁴Centre for Social Evolution, Department of Biology, Universitetsparken 15, University of Copenhagen, Copenhagen DK-2100, Denmark, ¹⁵Trace and Environmental DNA Laboratory, Department of Environment and Agriculture, Curtin University, 6102 Perth, Australia and ¹⁶Norwegian University of Science and Technology, University Museum, 7491 Trondheim, Norway

*Correspondence address. BGI-Shenzhen, Shenzhen 518083, China. Tel: +86-0755-25273620; E-mail: jianghui@genomics.cn; Centre for GeoGenetics, Natural History Museum of Denmark, University of Copenhagen, Øster Voldgade 5–7, 1350 Copenhagen, Denmark. Tel: +45 23 71 25 19; E-mail: tgilbert@snm.ku.dk

[†]Equal contribution

Correction

In the article, 'Comparative performance of the BGISEQ-500 vs Illumina HiSeq2500 sequencing platforms for palaeogenomic sequencing, *GigaScience* 2017 6:8' by Mak *et al.* [1], the first author's names appear incorrectly in References 104 and 105.

The correct names in References 104 and 105 are:

104. Mak SST, Gopalakrishnan S, Carøe C *et al.* Supporting data for 'Comparative performance of the BGISEQ-500 vs Illumina sequencing platforms for palaeogenomic sequencing.' *GigaScience Database* 2017. <http://dx.doi.org/10.5524/100303> (accessed 1 March 2017)

105. Mak SST, Gopalakrishnan S, Carøe C *et al.* Protocols from 'Comparative performance of the BGISEQ-500 vs Illumina sequencing platforms for palaeogenomic sequencing.' *protocols.io* 2017. <http://dx.doi.org/10.17504/protocols.io.h99b996> (accessed 1 March 2017)

There is an error in numbering of Supplemental Table S6 in Supplemental File F1. The correct numbering should be Supplemental Table S7 and its corresponding text in the last sentence of first paragraph in Supplemental File F1 should be: Supplemental Table S7 gives an overview of the different conditions, amount of enzyme and addition of Reaction Enhancer.

References

1. Mak SST, Gopalakrishnan S, Carøe C *et al.* Comparative performance of the BGISEQ-500 vs Illumina sequencing platforms for palaeogenomic sequencing. *GigaScience* 2017;6(8):1–13 <https://doi.org/10.1093/gigascience/gix049>