

**FAST-EM array tomography:
a workflow for multibeam volume electron microscopy**

Kievits, A.J.; Duinkerken, B.H. Peter ; Lane, R.; de Heus, Cecilia ; van Beijeren Bergen en Henegouwen, Daan ; Höppener, T.R.; Wolters, Anouk H.G.; Liv, Nalan; Giepmans, Ben N.G.; Hoogenboom, J.P.

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Corrigendum

Arent J. Kievits*, B. H. Peter Duinkerken, Ryan Lane, Cecilia de Heus, Daan van Beijeren Bergen en Henegouwen, Tibbe Höppener, Anouk H. G. Wolters, Nalan Liv, Ben N. G. Giepmans and Jacob P. Hoogenboom*

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Corrigendum to: Arent J. Kievits*, B. H. Peter Duinkerken, Ryan Lane, Cecilia de Heus, Daan van Beijeren Bergen en Henegouwen, Tibbe Höppener, Anouk H. G. Wolters, Nalan Liv, Ben N. G. Giepmans and Jacob P. Hoogenboom. FAST-EM array tomography: a workflow for multibeam volume electron microscopy. *Methods in Microscopy (MIM)*. 2024; Volume 1, Issue 1, pp. 49–64 (DOI: <https://doi.org/10.1515/mim-2024-0005>).

***Corresponding authors: Arent J. Kievits and Jacob P. Hoogenboom**, Department of Imaging Physics, Delft University of Technology, Delft, The Netherlands, E-mail: A.J.Kievits@tudelft.nl (A. J. Kievits), J.P.hoogenboom@tudelft.nl (J. P. Hoogenboom). <https://orcid.org/0000-0003-4457-9627> (A. J. Kievits). <https://orcid.org/0000-0003-4539-8772> (J. P. Hoogenboom)

B. H. Peter Duinkerken, Daan van Beijeren Bergen en Henegouwen, Anouk H. G. Wolters and Ben N. G. Giepmans, Department of Biomedical Sciences, University Medical Center Groningen, Groningen, The Netherlands. <https://orcid.org/0000-0003-0699-0001> (B.H.P. Duinkerken). <https://orcid.org/0000-0001-5105-5915> (B.N.G. Giepmans)

Ryan Lane and Tibbe Höppener, Department of Imaging Physics, Delft University of Technology, Delft, The Netherlands. <https://orcid.org/0000-0002-5887-2069> (R. Lane)

Cecilia de Heus and Nalan Liv, Center for Molecular Medicine, University Medical Center Utrecht, Utrecht, The Netherlands. <https://orcid.org/0000-0001-8618-8451> (C. de Heus). <https://orcid.org/0000-0003-2654-9117> (N. Liv)

The authors have noted that some information is missing in the published article:

- **Data availability:** The full 3D datasets as presented in this article can be viewed through Nanotomomy.org (<http://www.nanotomomy.org/OA/Kievits2024MIM/index.html>). The full raw datasets are available through the 4TU repository (doi: <https://doi.org/10.4121/bf3f2b23-2328-4d81-a0f4-05fdb33117d7>) and EMPIAR (entries EMPIAR-12174, EMPIAR-12190 and EMPIAR-12193).
- **Software availability statement:** The code to generate the 3D reconstructions presented in this article and a sample dataset are available through the 4TU repository (doi: <https://doi.org/10.4121/bf3f2b23-2328-4d81-a0f4-05fdb33117d7>). The scripts used to analyze the MitoNet results (Table 2) and generate the views in Figure 6 and S5 are available on Github: <https://github.com/hoogenboom-group/Kievits-FASTEM-array-tomography-2024>.
- The reference footnote to the online data of Figure 6 is not working. The link to the data is: <http://www.nanotomomy.org/OA/Kievits2024MIM/index.html>
- Table 3 contains a minor copy-editing mistake. In the row that indicates the dwell time, the number indicated for “bd-TEM” should be the dwell time for “ATUM-MultiSEM”. The dwell time for bd-TEM is equal to the FoV acquisition time.