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Corrigendum

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

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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.nanotomy.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.nanotomy.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.

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