Manuscript Title

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Abstract

Background & Summary

We scanned a *lot* of different mouse strains and so-called hybrids, both male and female. One scan for each strain, each hybrid and each sex gives already 90 samples.

Methods

Sample preparation

- · Mice were anesthesized
- Mice were decapitated
- Mouse skulls were stored in ...

Tomographic imaging

- SkyScan 1272 with sample changer
- All log files available here: https://github.unibe.ch/david-haberthuer/InMice/tree/main/logfiles
- Report scaning settings and reconstruction parameter data according to¹

Image processing

• *Jupyter*² notebooks, available here: https://github.unibe.ch/david-haberthuer/InMice/, for reproducible research.

Data Records

Technical Validation

Usage Notes

Code Availability

• Jupyter notebooks

References

- 1. Stock, S. R. & De Carlo, F. <u>Meta-data for absorption tomography measurements</u>. *Tomography of Materials and Structures* **3**, 100015 (2023).
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- 3. ANSI/NISO Z39.104-2022, CRediT, Contributor Roles Taxonomy. doi:10.3789/ansi.niso.z39.104-2022.
- 4. Himmelstein, D. S. *et al.* Open collaborative writing with Manubot. *PLoS Comput Biol* **15**, e1007128 (2019).

Author Contributions

Contributor Roles Taxonomy (CRediT), as defined in³:

- <u>Data curation</u>: David Haberthür, Larisa Petra Kaija
- Formal analysis: David Haberthür
- Investigation: David Haberthür, Larisa Petra Kaija
- Methodology: David Haberthür, Larisa Petra Kaija
- Project administration: David Haberthür
- Software: David Haberthür
- Validation: David Haberthür, Larisa Petra Kaija
- Visualization: David Haberthür
- Writing original draft: David Haberthür
- Writing review & editing: David Haberthür, Larisa Petra Kaija

Competing Interests

Author	Competing Interests	Last Reviewed
David Haberthür	None	2025-06-27
Larisa Petra Kaija	None	

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