

Derotation: a Python package for correcting rotation-induced distortions in line-scanning microscopy

Laura Porta¹, Simon Weiler¹, Adam L. Tyson^{1, 2}, and Troy W. Margrie¹

1 Sainsbury Wellcome Centre, University College London, UK 2 Gatsby Computational Neuroscience Unit, University College London, London, United Kingdom

Summary

Line-scanning microscopy, including multiphoton calcium imaging, is a powerful technique for observing dynamic processes at cellular resolution. However, when the imaged sample rotates during acquisition, the sequential line-by-line scanning process introduces geometric distortions. These artifacts, which manifest as shearing or curving of features, can severely compromise downstream analyses such as motion registration, cell detection, and signal extraction. While several studies have developed custom solutions for this issue (Vélez-Fort et al., 2018) (Hennestad et al., 2021) (Sit & Goard, 2023) (Voigts & Harnett, 2020), a general-purpose, accessible software package has been lacking.

derotation is an open-source Python package that algorithmically reconstructs movies from data acquired during sample rotation. By leveraging recorded rotation angles and the microscope's line acquisition clock, the software applies a precise, line-by-line inverse transformation to restore the original geometry of the imaged plane. This correction enables reliable cell segmentation during rapid rotational movements, making it possible to study yaw motion without sacrificing image quality (Figure 1).

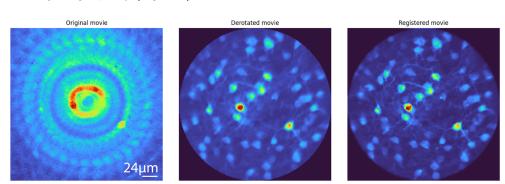


Figure 1: Example of derotation correction. On the left, the mean image of a 3-photon movie in which the sample was rotating. In the center, the mean image after derotation, and on the left the mean image of the derotated movie after suite2p registration (Pachitariu et al., 2016). As you can see, already after derotation the cells are visible and have well defined shapes. See the collection of 3-photon imaging data from head-fixed mice for more details.

Statement of Need

Any imaging modality that acquires data sequentially, such as line-scanning microscopy, is susceptible to motion artifacts if the sample moves during the acquisition of a single frame. When this motion is rotational, it produces characteristic "fan-like" distortions that corrupt the morphological features of the imaged structures (Figure 2). This significantly complicates,

DOI: DOIunavailable

Software

- Review 🗗
- Repository 🗗
- Archive ♂

Editor: Pending Editor ♂ Reviewers:

@Pending Reviewers

Submitted: N/A **Published:** N/A

License

Authors of papers retain copyright and release the work under a Creative Commons Attribution 4.0 International License (CC BY 4.0).



or even prevents, critical downstream processing steps like cell segmentation and automated region-of-interest tracking.

This problem is particularly acute in systems neuroscience, where researchers increasingly combine two-photon or three-photon calcium imaging with behavioral paradigms involving head rotation (Vélez-Fort et al., 2018) (Hennestad et al., 2021) (Sit & Goard, 2023) (Voigts & Harnett, 2020). In such experiments, where head-fixed animals may be passively or actively rotated, high-speed angular motion can render imaging data unusable without correction. The issue is even more acute in imaging modalities with lower frame rates, such as three-photon calcium imaging. While individual labs have implemented custom scripts to address this, there has been no validated, open-source, and easy-to-use Python tool available to the broader community.

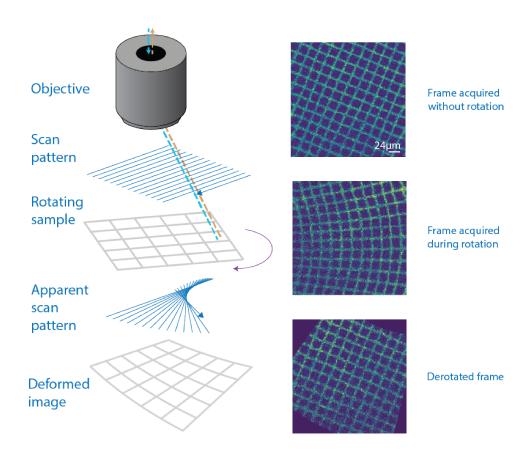


Figure 2: Schematic of line-scanning microscope distortion. Left: line scanning pattern plus sample rotation lead to fan-like artifacts when imaging a grid. Right: grid imaged while still (top), while rotating (middle), and after derotation (bottom), showing alignment restoration.

derotation directly fills this gap by providing a documented, tested, and modular solution for post hoc correction of imaging data acquired during rotation. It enables researchers to perform quantitative imaging during high-speed rotational movements. By providing a robust and accessible tool, derotation lowers the barrier for entry into complex behavioral experiments and improves the reproducibility of a key analysis step in a growing field of research.



Functionality

The core of the derotation package is a line-by-line affine transformation. It operates by first establishing a precise mapping between each scanned line in the movie and the rotation angle of the sample at that exact moment in time. It then applies an inverse rotation transform to each line around a specified or estimated center of rotation. Finally, the corrected lines are reassembled into frames, producing a movie that appears as if the sample had remained stationary.

Data Ingestion and Synchronization

The package accepts two types of input formats depending on the processing approach:

For pipeline workflows (FullPipeline and IncrementalPipeline): These pipelines are designed for experimental setups with synchronized rotation and imaging data. The required inputs are:

- An array of analog signals containing timing and rotation information, typically including the start of a new line and frame, when the rotation system is active, and the rotation position feedback;
- A **CSV** file describing speeds and directions.

For low-level core function: Advanced users can bypass the pipeline workflows and use the core transformation function directly by providing the original movie and an array of rotation angles for each line.

This modular design allows users with custom experimental setups to integrate the derotati on algorithm into their own analysis scripts while still benefiting from the core transformation logic.

Processing Pipelines

For ease of use, derotation provides two high-level processing workflows tailored to common experimental paradigms. These pipelines handle data loading, parameter validation, processing, and saving outputs, while also generating logs and debugging plots to ensure quality control.

- FullPipeline is engineered for experimental paradigms involving randomized, clockwise or counter-clockwise rotations. It assumes that there will be complete 360° rotations of the sample. As part of its workflow, it can optionally estimate the center of rotation automatically using Bayesian optimization, which minimizes residual motion in the corrected movie.
- IncrementalPipeline is optimized for stepwise, single-direction rotations. This rotation paradigm is useful for calibration of the luminance across rotation angles. It can also provide an alternative estimate of the center of rotation, fitting the trajectory of a cell across rotation angles.

Both pipelines are configurable via YAML files or Python dictionaries, promoting reproducible analysis by making it straightforward to document and re-apply the same parameters across multiple datasets.

Upon completion, a pipeline run generates a comprehensive set of outputs: the corrected movie, a CSV file with rotation angles and metadata for each frame, debugging plots, a text file containing the estimated optimal center of rotation, and log files with detailed processing information.



Validation

The package's effectiveness has been validated on three-photon recordings from head-fixed mice (Figure 3). The corrected images showed restored cellular morphology and were successfully processed by standard downstream analysis pipelines such as Suite2p (Pachitariu et al., 2016). In Figure 3, it is possible to compare the $\Delta F/F_0$ of two ROIs in the case of line-by-line derotation (as implemented in derotation) and frame-by-frame derotation (using scipy.ndimage.affine_transform). The line-by-line derotation restores the $\Delta F/F_0$ to its original value during rotation times, reducing the dips into negative values that are present in the frame-by-frame derotation.

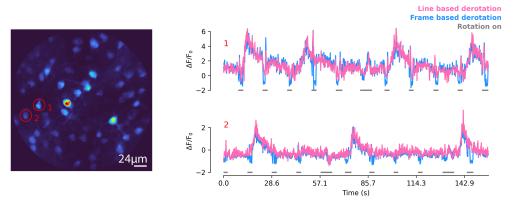


Figure 3: Example of derotation correction on 3-photon imaging data from head-fixed mice. Left: mean image of a 3-photon movie derotated line-by-line. Red circles represent the ROIs selected for the plots on the right. Right: Two segments of $\Delta F/F_0$ at different time points of two ROIs comparing frame-by-frame derotation, and line-by-line derotation. Gray line represents the rotation on times.

Synthetic Data Generation

For further development and testing, derotation includes a synthetic data generator that can create challenging synthetic datasets with misaligned centers of rotation and out-of-plane rotations. This feature is particularly useful for validating the robustness of the derotation algorithm and for developing new features.

The synthetic data can be generated using the following classes:

- Rotator class: Core class that applies line-by-line rotation to an image stack, simulating a rotating microscope.
- SyntheticData class: Creates fake cell images, assigns rotation angles, and generates synthetic stacks leveraging the Rotator class. It is a complete synthetic dataset generator.

Documentation and Installation

derotation is available on PyPI and can be installed with pip install derotation. It is distributed under a BSD-3-Clause license. Comprehensive documentation, tutorials, and example datasets are available at https://derotation.neuroinformatics.dev. Using Binder, users can run the software in a cloud-based environment with sample data without requiring any local installation.

Future Directions

Derotation is currently used to process 3-photon movies acquired during head rotation. Future directions can include further automated pipelines for specific motorised stages and experimen-



tal paradigms.

Methodological appendix

Collection of 3-photon imaging data from head-fixed mice

The package has been directly tested on 3-photon imaging data obtained from cortical layer 6 callosal-projecting neurons epxressing the calcium indicator GCaMP7f in the mouse visual cortex (see Figure 3). More specifically, wild type C57/B6 mice were injected with retro AAV-hSyn-Cre $(110^{14} \text{ units per ml})$ in the left and with AAV2/1.syn.FLEX.GCaMP7f $(1.810^{13} \text{ units per ml})$ in the right primary visual cortex. A cranial window was implanted over the right hemisphere, and then a headplate was cemented onto the skull of the animal. After 4 weeks of viral expression and recovery, animals were head-fixed on a rotation platform driven by a direct-drive motor (U-651, Physik Instrumente). 360 degree clockwise and counterclockwise rotations with different speed profiles (50, 100, 150, 200 deg/s) were performed while simultaneuolsy imaging awake neuronal activity using a 25x objective (XLPLN25XWMP2, NA 1.05, 25, Olympus). Imaging was conducted at 7 Hz with 256x256 pixels. For detailed description of the 3-photon power source and imaging please see (Cloves & Margrie, 2024).

Acknowledgements

We thank Eivind Hennestad for initial project discussion. We thank Mateo Vélez-Fort and Chryssanthi Tsitoura for their assistance in building and testing the the three-photon imaging and rotation setup as well as for feedback on the package. We also thank Igor Tatarnikov for contributing to the development of the package and the whole Neuroinformatics Unit. This package was inspired by previous work on derotation as described in (Voigts & Harnett, 2020). The authors are grateful to the support staff of the Neurobiological Research Facility at Sainsbury Wellcome Center (SWC). This research was funded by the Sainsbury Wellcome Centre core grant from the Gatsby Charitable Foundation (GAT3361) and Wellcome Trust (219627/Z/19/Z), a Wellcome Trust Investigator Award (214333/Z/18/Z) and Discovery Award (306384/Z/23/Z) to T.W.M. and by SWC core funding to the Neurobiological Research Facility. S.W. was funded by a Feodor-Lynen fellowship from the Alexander von Humboldt Foundation.

References

- Cloves, M., & Margrie, T. W. (2024). In vivo dual-plane 3-photon microscopy: Spanning the depth of the mouse neocortex. *Biomedical Optics Express*, *15*(12), 7022. https://doi.org/10.1364/BOE.544383
- Hennestad, E., Witoelar, A., Chambers, A. R., & Vervaeke, K. (2021). Mapping vestibular and visual contributions to angular head velocity tuning in the cortex. *Cell Reports*, *37*(12), 110134. https://doi.org/10.1016/j.celrep.2021.110134
- Pachitariu, M., Stringer, C., Dipoppa, M., Schröder, S., Rossi, L. F., Dalgleish, H., Carandini, M., & Harris, K. D. (2016). Suite2p: Beyond 10,000 neurons with standard two-photon microscopy. Neuroscience. https://doi.org/10.1101/061507
- Sit, K. K., & Goard, M. J. (2023). Coregistration of heading to visual cues in retrosplenial cortex. *Nature Communications*, *14*(1), 1992. https://doi.org/10.1038/s41467-023-37704-5
- Vélez-Fort, M., Bracey, E. F., Keshavarzi, S., Rousseau, C. V., Cossell, L., Lenzi, S. C., Strom, M., & Margrie, T. W. (2018). A Circuit for Integration of Head- and Visual-Motion Signals in Layer 6 of Mouse Primary Visual Cortex. *Neuron*, *98*(1), 179–191.e6. https://doi.org/10.1016/j.neuron.2018.02.023



Voigts, J., & Harnett, M. T. (2020). Somatic and Dendritic Encoding of Spatial Variables in Retrosplenial Cortex Differs during 2D Navigation. *Neuron*, *105*(2), 237–245.e4. https://doi.org/10.1016/j.neuron.2019.10.016