nature portfolio

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Reporting Summary

Nature Portfolio wishes to improve the reproducibility of the work that we publish. This form provides structure for consistency and transparency in reporting. For further information on Nature Portfolio policies, see our <u>Editorial Policies</u> and the <u>Editorial Policy Checklist</u>.

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For	all statistical analyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.		
n/a	Confirmed		
	The exact sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement		
	A statement on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly		
	The statistical test(s) used AND whether they are one- or two-sided Only common tests should be described solely by name; describe more complex techniques in the Methods section.		
X	A description of all covariates tested		
\times	A description of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons		
	A full description of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient AND variation (e.g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals)		
	For null hypothesis testing, the test statistic (e.g. <i>F</i> , <i>t</i> , <i>r</i>) with confidence intervals, effect sizes, degrees of freedom and <i>P</i> value noted <i>Give P values as exact values whenever suitable.</i>		
X	For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings		
X	For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes		
X	Estimates of effect sizes (e.g. Cohen's <i>d</i> , Pearson's <i>r</i>), indicating how they were calculated		
	Our web collection on <u>statistics for biologists</u> contains articles on many of the points above.		
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Software and code

Policy information about <u>availability of computer code</u>

Data collection Micro manager 1.44

Data analysis Matlab R2017a

For manuscripts utilizing custom algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors and reviewers. We strongly encourage code deposition in a community repository (e.g. GitHub). See the Nature Portfolio guidelines for submitting code & software for further information.

Data

Policy information about availability of data

All manuscripts must include a data availability statement. This statement should provide the following information, where applicable:

- Accession codes, unique identifiers, or web links for publicly available datasets
- A description of any restrictions on data availability
- For clinical datasets or third party data, please ensure that the statement adheres to our policy

Source data are provided with this paper. Light-sheet microscopy datasets exceed multiple gigabyte, and will be made available upon reasonable request.

Policy information about studies involving human research participants and Sex and Gender in Research. Reporting on sex and gender NA Population characteristics NA Recruitment NA Ethics oversight NA Note that full information on the approval of the study protocol must also be provided in the manuscript.
Population characteristics NA Recruitment NA Ethics oversight NA
Recruitment NA Ethics oversight NA
Ethics oversight NA
Note that full information on the approval of the study protocol must also be provided in the manuscript.
Field-specific reporting
Please select the one below that is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.
✓ Life sciences ☐ Behavioural & social sciences ☐ Ecological, evolutionary & environmental sciences
For a reference copy of the document with all sections, see nature.com/documents/nr-reporting-summary-flat.pdf
Life sciences study design
All studies must disclose on these points even when the disclosure is negative.
Sample size Under the assumption that biological variability is normal distributed, the sample size was chosen as five or more biological repeats per condition
Data exclusions No data was excluded
Replication All attempts of replication where successful The general strategy consists of time aligning datasets, and then computing correlation functions.
Randomization This is a hypothesis driven experimental work. These hypotheses are tested using specific perturbation methods, designed by the experimentalists. It is unclear how randomization applies in such a situation.
Blinding This is a hypothesis driven experimental work. These hypotheses are tested using specific perturbation methods, designed by the experimentalists. It is unclear how blinding applies in such a situation.
Reporting for specific materials, systems and methods
We require information from authors about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material,
system or method listed is relevant to your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.
Materials & experimental systems Methods
n/a Involved in the study n/a Involved in the study
Antibodies ChIP-seq
Eukaryotic cell lines Flow cytometry
Palaeontology and archaeology MRI-based neuroimaging Animals and other organisms
Clinical data
Dual use research of concern
Animals and other research organisms
Policy information about <u>studies involving animals</u> ; <u>ARRIVE guidelines</u> recommended for reporting animal research, and <u>Sex and Gender in Research</u>

Laboratory animals

2 hours old Drosophila melanogaster embryos

Wild animals	No wild animals where used for this study.
Reporting on sex	NA
Field-collected samples	No field collected samples where used in this study.
Ethics oversight	This study did not require an ethics oversight committee.

Note that full information on the approval of the study protocol must also be provided in the manuscript.