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 Editorial Policies and the Editorial Policy Checklist.

Statistics

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.
☐ A description of all covariates tested
☐ 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. mean), variability (e.g. standard deviation), and sample size (n)
☐ For null hypothesis testing, the test statistic (e.g. F, t, r) with confidence intervals, effect sizes, degrees of freedom and P value noted. Give P values as exact values whenever possible.
☐ For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings
☐ For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes
☐ Estimates of effect sizes (e.g. Cohen’s d, Pearson’s r), indicating how they were calculated

Our web collection on statistics for biologists contains articles on many of the points above.

Software and code

Policy information about availability of computer code

Data collection
- WSM4.1.3; NI-DAQmx and BNC-2110 LabVIEW; AR120804+1821 + Igor Pro 6.22A, PKPM Data Processing, LEaP (Amber); SDA-flex 7.1; Amber 14
- MATLAB, LabVIEW, OriginPro 8.5.0 SR1; Origin 2018, Origin 2019b; MultiSeq VMD 13.1; Matplotlib, SDA-flex 7.1; UCSF Chimera 1.15rc; ABPS1.5, volmap plugin 1.1, matplotlib 3.5.1, VMD 1.9.4ab5, Graphpad Prism 9

For manuscripts utilizing custom algorithms or software that are 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

All the data generated in this study are available within the main text and the Supplementary Information file; source data are provided in the Source Data file. Data is also available from the corresponding author upon request.
Human research participants

Policy information about studies involving human research participants and Sex and Gender in Research.

| Reporting on sex and gender | n/a |
|-----------------------------|-----|
| Population characteristics  | n/a |
| Recruitment                 | n/a |
| Ethics oversight            | n/a |

Note: 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 | Sample size was determined according to what is considered sufficient in published reports on similar measurements |
|-------------|---------------------------------------------------------------------------------------------------|
| Data exclusions | No data exclusions |
| Replication | For the type of measurements, not much variability is expected. At least 2 independent measurements were taken. |
| Randomization | Randomisation was not applicable, since in all cases the size of the sample was small (N = 2 or 3 independent experiments), which we understand that could not guarantee the equivalence between groups. |
| Blinding | Since all the measurements were objective, we estimated the blinding unnecessary. |

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 |
| x Antibodies                     | x ChiP-seq |
| x Eukaryotic cell lines          | x Flow cytometry |
| x Palaeontology and archaeology  | x MRI-based neuroimaging |
| x Animals and other organisms    |         |
| x Clinical data                  |         |
| x Dual use research of concern   |         |