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## Reporting Summary

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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.   |
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| 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. 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. F, t, r) with confidence intervals, effect sizes, degrees of freedom and P value noted Give P values as exact values whenever suitable.   |
| 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  |
|     | 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

Provide a description of all commercial, open source and custom code used to collect the data in this study, specifying the version used OR state that no software was used.

Data analysis

Provide a description of all commercial, open source and custom code used to analyse the data in this study, specifying the version used OR state that no software was used.

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 Research 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 list of figures that have associated raw data
- A description of any restrictions on data availability

The data shown in Figure 4 is available from the supporting primary research paper previously published by Perez-Burillo et al.17.

- The data presented in Figs. 2&3 were generated for this protocol.
- The source data underlying Figs. 2&3 are provided as Source Data files with this protocol.

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| Field-specific reporting   |   |  |  |  |  |  |  |
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| Life sciences study design   |   |  |  |  |  |  |  |
| All studies must disclose on these points even when the disclosure is negative.  |   |  |  |  |  |  |  |
| Sample size  | Not needed for a protocol paper   |  |  |  |  |  |  |
| Data exclusions  | Not needed for a protocol paper   |  |  |  |  |  |  |
| Replication  | Not needed for a protocol paper   |  |  |  |  |  |  |
| Randomization  | Not needed for a protocol paper   |  |  |  |  |  |  |
| Blinding   | Not needed for a protocol paper   |  |  |  |  |  |  |
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| Reportin   | g for specific i  | materials, systems and methods   |  |  |  |  |  |
|  |   | s of materials, experimental systems and methods used in many studies. Here, indicate whether each material, are not sure if a list item applies to your research, read the appropriate section before selecting a response. |  |  |  |  |  |
|  | perimental systems  | Methods  |  |  |  |  |  |
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| n/a Involved in the study Antibodies   |   | ChiP-seq   |  |  |  |  |  |
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| Palaeontology and archaeology  |   | MRI-based neuroimaging   |  |  |  |  |  |
| Animals and other organisms  |   |  |  |  |  |  |  |
| Human research participants  |   |  |  |  |  |  |  |
| Clinical data  |   |  |  |  |  |  |  |
| Dual use research of concern   |   |  |  |  |  |  |  |
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| Human research participants  |   |  |  |  |  |  |  |

Policy information about studies involving human research participants

Population characteristics

Not needed for a protocol paper

Recruitment

Describe how participants were recruited. Outline any potential self-selection bias or other biases that may be present and how these are likely to impact results.

Ethics oversight

Not needed for a protocol paper

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

