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Reporting Summary

Nature Research 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 Research policies, seeAuthors & Referees and theEditorial Policy Checklist.

Statistics				
For all statistical analyse	es, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.			
n/a Confirmed				
The exact sam	ple size (n) for each experimental group/condition, given as a discrete number and unit of measurement			
A statement o	n 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	X A description of all covariates tested			
A description	🗷 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 Give P values as exact values whenever suitable.				
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 e	ffect sizes (e.g. Cohen's d , Pearson's r), indicating how they were calculated			
·	Our web collection on <u>statistics for biologists</u> contains articles on many of the points above.			
Software and c	ode			
Policy information abou	ut <u>availability of computer code</u>			
Data collection	Bruker Topspin 3.2			
Data analysis	nmrPipe, nmrDraw, origin 8.0			
	om algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors/reviewers. deposition in a community repository (e.g. GitHub). See the Nature Research guidelines for submitting code & software for further information.			
Data				
- Accession codes, uni - A list of figures that	at availability of data nclude a data availability statement. This statement should provide the following information, where applicable: ique identifiers, or web links for publicly available datasets have associated raw data restrictions on data availability			
The datasets generated a	and/or analysed during the current study are included in the supporting information file.			
Field-speci	fic reporting			
Please select the one be	elow 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 do	ocument 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	No sample size calculation was performed. The various sample sizes for the different experiments was selected as no significant difference was observed.	
Data exclusions	No data were excluded in the analyses.	
Replication	All results were sucessfully replicated.	
Randomization	No randomization was performed because it is not relevant to the current study.	
Blinding	No blinding was performed because it is not relevant to the current study.	

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	
x	Eukaryotic cell lines	Flow cytometry	
x	Palaeontology	MRI-based neuroimaging	
x	Animals and other organisms	,	
X	Human research participants		
X	Clinical data		