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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, see Authors & Referees and the Editorial Policy Checklist.

Sta	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 san	nple 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.					
\boxtimes	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. <i>F</i> , <i>t</i> , <i>r</i>) with confidence intervals, effect sizes, degrees of freedom and <i>P</i> value noted Give <i>P</i> values as exact values whenever suitable.					
\boxtimes	For Bayesian	analysis, information on the choice of priors and Markov chain Monte Carlo settings				
\boxtimes	For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes					
\boxtimes	Estimates of e	effect 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.				
So	ftware and o	code				
Poli	cy information abo	out <u>availability of computer code</u>				
D	ata collection	Gen5 Secure				
Data analysis		Graphpad Prism 6.01, Microsoft Excel (2016), Zen software				
	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/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.					
Da	Data					
Policy information about <u>availability of data</u> All manuscripts must include a <u>data availability statement</u> . 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 authors declare that all data supporting the findings of this study are available within the article and its supplementary information or from the corresponding author upon reasonable request.						
Fi	Field-specific reporting					
Plea	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.					
\times	Life sciences Behavioural & social sciences Ecological, evolutionary & environmental sciences					

Life sciences study design

all studies must disclose on these points even when the disclosure is negative.					
Sample size	Details about chosen sample size are indicated in figure legends and/or methods. At least 3 biological repeats were performed for all experiments.				
Data exclusions	No data were excluded from the analysis.				
Replication	At least 3 biological repeats were performed for all experiments. Details about chosen sample size are indicated in figure legends and/or methods.				
Randomization	All experiments started from random clones.				
Blinding	NA				

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.

Ma	terials & experimental systems	Methods		
n/a	Involved in the study	n/a	Involved in the study	
\boxtimes	Antibodies	\boxtimes	ChIP-seq	
\boxtimes	Eukaryotic cell lines	\boxtimes	Flow cytometry	
\boxtimes	Palaeontology	\boxtimes	MRI-based neuroimaging	
\boxtimes	Animals and other organisms			
\boxtimes	Human research participants			
\boxtimes	Clinical data			