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Last updated by author(s): Mar 5, 2019

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 <u>Authors & Referees</u> and the <u>Editorial Policy Checklist</u>.

Statistics

For	all st	atistical analyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.
n/a	Cor	firmed
		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.
	\square	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 <i>Give P values as exact values whenever suitable</i> .
\boxtimes		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
	\square	Estimates of effect sizes (e.g. Cohen's d, Pearson's r), indicating how they were calculated
	1	Our web collection on <u>statistics for biologists</u> contains articles on many of the points above.

Software and code

Policy information about availability of computer code					
Data collection	(N/A				
Data analysis	N/A				

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.

Data

Policy information about availability of data

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

All relevant data related to this manuscript are available on request from the authors on reasonable request. The accession number for the RNA-sequencing data described in this study is GSE126888. Original un-cropped western blots are provided in source data. The source data underlying Figs. 1-7 and Supplementary Figs. 2-5 are provided.

Field-specific reporting

K Life sciences

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.

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	FOR MOST EXPERIMENTS, AT LEAST 3 EXPERIMENTAL SAMPLES WERE USED				
Data exclusions	NO DATA IS EXCLUDED				
Replication	ALL EXPERIEMENTS WERE REPEATED AT LEAST THREE TIMES				
Randomization	N/A				
Blinding	ALL ANIMAL EXPERIEMENTS WERE PERFORMED DOUBLE BLINDED				

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

Antibodies

Antibodies used	Listed in Supplementary table 6
Validation	Following manufactures' instructions

Eukaryotic cell lines

Policy information about <u>cell lines</u>			
Cell line source(s)	N/A		
Authentication	Describe the authentication procedures for each cell line used OR declare that none of the cell lines used were authenticated.		
Mycoplasma contamination	Confirm that all cell lines tested negative for mycoplasma contamination OR describe the results of the testing for mycoplasma contamination OR declare that the cell lines were not tested for mycoplasma contamination.		
Commonly misidentified lines (See ICLAC register)	Name any commonly misidentified cell lines used in the study and provide a rationale for their use.		

Animals and other organisms

Policy information about studies involving animals; ARRIVE guidelines recommended for reporting animal research

Laboratory animals The use of animals in this study conformed to the Public Health Service Guide for Care and Use of Laboratory Animals and was

Laboratory animals	approved by the Institutional Animal Care and Use Committee (IACUC) of Boston Children's Hospital and by the Institutional Animal Care and Use Committee of Zhejiang University.
Wild animals	N/A
Field-collected samples	N/A
Ethics oversight	Institutional Animal Care and Use Committee (IACUC) of Boston Children's Hospital and by the Institutional Animal Care and Use Committee of Zhejiang University

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