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

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 (<i>n</i>) 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				
\boxtimes	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				
\boxtimes	Estimates of effect sizes (e.g. Cohen's <i>d</i> , Pearson's <i>r</i>), indicating how they were calculated				
Our web collection on <u>statistics for biologists</u> contains articles on many of the points above.					

Software and code

Policy information about <u>availability of computer code</u>					
Data collection	Commercial software was used to collect data. No custom software was used.				
Data analysis	OriginPro 8, GraphPad Prism 6.0, GE Microview 2.2, VivoQuant 2.0, Clear Canvas Worksation 7.0, IVIS Living Imaging 3.0 and Vevo LAB were used for data analysis.				

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 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 main data supporting the findings of this study are available within the paper and its Supplementary Information. All data generated, both raw images and analysed datasets, for the figures in this study are available from figshare with the identifier https://doi.org/10.6084/m9.figshare.10732283.v2.

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 <u>nature.com/documents/nr-reporting-summary-flat.pdf</u>

Life sciences study design

All studies must disclose on these points even when the disclosure is negative.					
Sample size	Sample sizes were determined according to previous experimental experience.				
Data exclusions	No data were excluded.				
Replication	All attempts at replication were successful.				
Randomization	All samples were randomly allocated into experimental groups.				
Blinding	No blinding was applied.				

Reporting for specific materials, systems and methods

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

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

Eukaryotic cell lines

Policy information about <u>cell lines</u>

Cell line source(s)	Murine breast cancer cells (4T1) were obtained from American Type Culture Collection (ATCC), and human glioma cancer U87-luciferase cells were obtained from PerkinElmer.		
Authentication	No further authentication was done after the cells were obtained from the vendors.		
Mycoplasma contamination	All cell lines were tested for mycoplasma contamination. No mycoplasma contamination was found.		
Commonly misidentified lines (See <u>ICLAC</u> register)	No commonly misidentified cell lines were used.		

Animals and other organisms

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

Laboratory animals	Balb/c female white or nude mice (20 ± 2g) were obtained from Charles River (USA) or Hunan SJA Laboratory Animal Co. Ltd. (China).
Wild animals	The study did not involve wild animals.
Field-collected samples	The study did not involve samples collected from the field.
Ethics oversight	All animal experiments were carried out in compliance with the Guidelines for the Care and Use of Research Animals established

by the Stanford University Animal Studies Committee and by the Institutional Animal Care and Use Committee of Hunan University.

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