natureresearch

Corresponding author(s):	Weisheng Guo, Yang Luo and Xing-Jie Liang
Last updated by author(s):	Aug 26, 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>.

5	ta	t	IS	tI	C	S

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 (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. <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
\times	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 <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

Digital Micrograph3.7, ZEN 2010, Supcre G9

Data analysis

Statistical analyses were performed on Graphpad Prism or SPSS 17.0.

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

The data that support the findings of this study are available within the paper and its Supplementary Information files, or are available from the corresponding authors upon reasonable request. The source data underlying Figs. 2c-e, 3a, 3c-f, 4b, c, 5b, c, 6d, e, 6g, h, 6j, 7a, b as well as Supplementary Figures 1, 3-8, 11-12, 14, 16 are provided as a Source Data file.

Field-specific reporting							
<u>.</u>		he best fit for your research. If you are not sure, read the appropriate sections before making your selection.					
X Life sciences	☐ Beł	navioural & social sciences Ecological, evolutionary & environmental sciences					
For a reference copy of t	he document with all	sections, see <u>nature.com/documents/nr-reporting-summary-flat.pdf</u>					
Life scier	ices stu	dy design					
All studies must dis	close on these po	oints even when the disclosure is negative.					
Sample size		fect size was predetermined, but sample sizes employed in this study are consistent with previously published works. For example, in studies were repeated at least three times independently and in the in vivo experiments with 3-4 mice per group were performed.					
Data exclusions	No animals and/o	r data were excluded.					
Replication	All experiments w	Il experiments were repeated and experimental findings were reproducible.					
Randomization	The dosing group	the dosing groups were filled by randomly selecting from the same pool of animals for in vivo experiments.					
Blinding	All the investigato	ors were blinded to group allocation during data collection and analysis.					
We require informatic system or method list Materials & exp. n/a Involved in th Antibodies Eukaryotic Palaeontok Animals and Human reso	on from authors abed is relevant to your perimental system of the study cell lines by dother organisms earch participants	ecific materials, systems and methods out some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material, our study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response. Methods					
Antibodies							
Antibodies used	by s	The following primary antibodies were used for enzyme-linked immunosorbent assay. They are listed as antibody first, followed by supplier, catalognumber and clone/lot number as applicable. 1). Anti-PCT, Andy Gene, cat. no. M0218					
Validation	All antibodies were verified by the supplier and each lot has been quality tested.						
Eukaryotic co	ell lines						
Policy information a							
Cell line source(s)	(3	BT3 and HUVEC cells were purchased from the Institute of Basic Medical Sciences, Chinese Academy of Medical Sciences.					
Authentication	Į.	A short tandem repeat DNA profiling method was used to authenticate the cell lines and the results were compared with					

reference database. Moreover, no mycoplasma contamination was detected in the above cell lines.

Mycoplasma contamination

All cell lines were tested for mycoplasma contamination. No mycoplasma contamination was found.

Commonly misidentified lines (See ICLAC register)

No. 3T3 and HUVEC cell lines are not listed in the database.

Animals and other organisms

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

Laboratory animals Balb/c mice (female, 6-8 weeks) were purchased from Beijing Charles River Company (Beijing, China).

Wild animals
No wild animals were used in this study.

Field-collected samples The study did not involve samples collected from field.

Ethics oversight

All animal experiment protocols were reviewed and approved by the Institutional Animal Care and Use Committee of National Center for Nanoscience and Technology and complied with all relevant ethical regulations.

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