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## **Reporting Summary**

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Statistics					
For all statistical analys	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	$\square$ The exact sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement				
A statement of	on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly				
The statistical Only common to	test(s) used AND whether they are one- or two-sided ests should be described solely by name; describe more complex techniques in the Methods section.				
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 descript AND variation	ion of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient) (e.g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals)				
X	thesis testing, the test statistic (e.g. $F$ , $t$ , $r$ ) with confidence intervals, effect sizes, degrees of freedom and $P$ value noted a 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	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 c	ode				
Policy information abou	ut <u>availability of computer code</u>				
Data collection No software was used.					
Data analysis	GraphPad Prism 7 were used for statistical analysis.				
	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					
<ul><li>Accession codes, un</li><li>A list of figures that</li></ul>	ut <u>availability of data</u> include a <u>data availability statement</u> . 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 authors declare that the corresponding autho	the data supporting the findings of this study are available within the article and its Supplementary Information Files, or are available from rs on resonable request.				
Field-speci	fic reporting				
Please select the one b	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					

## Life sciences study design

All studies must dis	close on these	points even when the disclosure is negative.	
Sample size	Sample size was chosen depending on the experimental approaches used based on our experience and estimation. No statistical method was used to predetermine sample size.		
Data exclusions	No data was excluded.		
Replication	The shown results could successfully and reliably be replicated and reproduced.		
Randomization	All the cells and mice were randomly allocated into different groups.		
Blinding	No blinding wa	blinding was done.	
We require informatic system or method list  Materials & exp n/a Involved in th	perimental sy e study cell lines ogy d other organism	n/a Involved in the study  ChIP-seq  Flow cytometry  MRI-based neuroimaging	
Antibodies used	ph PC ph p- β-a	NF-κB2 (p52), Santa Cruz, cat#sc-7386; NF-κB2 (p52) for ChIP, Santa Cruz, cat#sc-7386X; phospho-NF-κB p65 (Ser536), Cell Signaling Technology, cat#3033S; NF-κB (p65), Cell Signaling Technology, cat#4764S; PDE4B, Santa Cruz, cat#sc-25812; phospho-PKA substrates, Cell Signaling Technology, cat#9621S; phospho-CREB (Ser133), Cell Signaling Technology, cat#9198S; CREB, Cell Signaling Technology, cat#9197S; p-NF-κB2 P100 (s866/870), Cell Signaling Technology, cat#4810P; NF-κB2 P100, Cell Signaling Technology, cat#4882T; β-actin, Cell Signaling Technology, cat#4970S; β-actin, Bioss, cat#bs-0061R; GAPDH, Cell Signaling Technology, cat#5174S; PCNA, Beyotime, cat#AF1363.	
Validation	All	antibodies used in the study are commercially available and validation statement is given on manufacturer's website.	
Eukaryotic c	ell lines		
Policy information a	about <u>cell lines</u>		
Cell line source(s)		HepG2 cell line was obtained from the American Type Culture Collection and 293T cell line was obtained from Stem Cell Bank, Chinese Academy of Sciences.	
Authentication		The cell lines were purchased from the authorized source, and none of them were authenticated thereafter.	
Mycoplasma cont	tamination	293T was negative for mycoplasma. HepG2 cell line used tested negative for mycoplasma contamination prior to this study.	
Commonly miside	entified lines	No commonly misidentified lines were used in this study.	

## Animals and other organisms

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

Laboratory animals

(See <u>ICLAC</u> register)

6-8-week-old male C57BL/6J mice were used in the study.

Wild animals The study did not involve wild animals.

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

The procedures for experiments and animal care were approved by the Institutional Animal Care and Use Committee of China Ethics oversight

Pharmaceutical University (Nanjing, China).

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