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Last updated by author(s):	Nov 25, 2020

## **Reporting Summary**

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Statistics	
For all statistical analys	ses, 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 statistica Only common t	l test(s) used AND whether they are one- or two-sided rests 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 descript AND variation	cion of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient) in (e.g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals)
For null hypo Give P values a	thesis testing, the test statistic (e.g. $F$ , $t$ , $r$ ) with confidence intervals, effect sizes, degrees of freedom and $P$ value noted is exact values whenever suitable.
For Bayesian	analysis, information on the choice of priors and Markov chain Monte Carlo settings
For hierarchic	cal 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 o	code
Policy information abo	ut availability of computer code
Data collection	NA
Data analysis	NA
	com 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, ur</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: iique identifiers, or web links for publicly available datasets have associated raw data v restrictions on data availability
The data generated and the corresponding author	analyzed during the current study are included in this published article and its supplementary information file. Datasets are available from or on request.
Field-spec	ific reporting
Please select the one k	below that is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.
<b>x</b> Life sciences	Behavioural & social sciences Ecological, evolutionary & environmental sciences

## Life sciences study design

NA

Wild animals

THE SCIENCES S	tudy design	
All studies must disclose on the	ese points even when the disclosure is negative.	
	Sample size for each experiment is indicated in the figure legend for each experiment. The sample size was chosen based on previous experience for each experiment to yield high power to detect specific effects. No statistical methods were used to predetermine sample size.	
Data exclusions No data we	No data were exclued from the analysis.	
Replication All experime	All experimental findings were reliably reproducible.	
Randomization Each sample	ach samples were randomly allocated into experimental groups.	
Blinding For all expe	riments, the investigators were blind to the experimental groups during the analyses.	
We require information from auth	specific materials, systems and methods  ors about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material, 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 & experimenta	al systems Methods	
n/a Involved in the study	n/a Involved in the study    ChIP-seq     Flow cytometry     MRI-based neuroimaging	
Antibodies used	The antibodies used: anti-Phospho-CaMKII (Thr286) (D21E4, Cell Signaling Technology; Danvers, MA, USA); anti-CaMKIIα (6G9; Cell Signaling Technology); anti-CaMKIIβ (ab34703; Abcam); anti-GFP (M048-3, MBL; Woburn, MA, USA); anti-Cdc42 (11A11; Cell Signaling Technology); anti-β-Actin (8H10D10; Cell Signaling Technology); HRP-anti-mouse and -rabbit (Jackson Laboratory; Bar Harbor, ME, USA); anti-RFP (1G9, MBL; Woburn, MA, USA).	
Validation	All antibodies used in our manuscript were validated by the manufactures or validated in the manuscript.	
Eukaryotic cell lines		
Policy information about <u>cell li</u>	<u>nes</u>	
Cell line source(s)	HeLa, HEK293	
Authentication	None of the cell lines used were authenticated.	
Mycoplasma contamination	The cell lines were not tested for mycoplasma contamination.	
Commonly misidentified line (See <u>ICLAC</u> register)	S NA	
Animals and other o	organisms	
Policy information about studion	es involving animals; ARRIVE guidelines recommended for reporting animal research	
Laboratory animals	All dissociated cultures were prepared from C57BL/6N mice (SLC). All slice cultures were prepared from Wistar rats (SLC or Charles River). This study used dissociated and slice cultures from both male and female pups. For in vivo imaging, the experiments were performed using male and female adult (2 months) C57BL/6N mice (Charles River).	

Field-collected samples

NA

Ethics oversight

All animal procedures were approved by the National Institutes of Natural Sciences Animal Care and Use Committee and were performed under the relevant guidelines and regulations.

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