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Reporting Summary

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Statistics	
For all statistical an	nalyses, 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 stateme	ent on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly
The statis Only comm	tical test(s) used AND whether they are one- or two-sided non tests should be described solely by name; describe more complex techniques in the Methods section.
A descript	tion of all covariates tested
A descript	tion of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons
	cription of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient) ition (e.g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals)
X	ypothesis testing, the test statistic (e.g. F , t , r) with confidence intervals, effect sizes, degrees of freedom and P value noted es as exact values whenever suitable.
For Bayes	ian analysis, information on the choice of priors and Markov chain Monte Carlo settings
For hierar	chical 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 an	d code
Policy information	about <u>availability of computer code</u>
Data collection	N/A
Data analysis	N/A
	g custom algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors and 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 <u>availability of data</u>
•	ust include a <u>data availability statement</u> . This statement should provide the following information, where applicable:
	s, unique identifiers, or web links for publicly available datasets that have associated raw data
- A description of	f any restrictions on data availability
The data that suppor	rt the findings of this study are available from the corresponding author upon reasonable request.

Field-spe	cific reporting	
\times Life sciences	ne 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 be document with all sections, see nature.com/documents/nr-reporting-summary-flat.pdf	
Life scier	nces study design	
All studies must dis	close on these points even when the disclosure is negative.	
Sample size	ample size was demonstrated in pilot experiments to achieve sufficient outcomes.	
Data exclusions	data was excluded.	
Replication	st all of the experiments in this study were performed at least three times independently and inconsistent results were not observed.	
Randomization	Experimental group were defined by specific treatment or control to exam the effects of treatments. Randomization was not relevant to this study. RNA-seq was conducted once with three biological samples but validated by RT-qPCR.	
Blinding	Analysis was objective and did not require blinding.	
We require informatis system or method lists and the system or method lists. Materials & ex n/a Involved in the Antibodies Eukaryotic Palaeontol Animals and Human res Clinical date Dual use res	cell lines cell lines mathematicipants ChIP-seq Flow cytometry MRI-based neuroimaging dother organisms earch participants	
Antibodies		
Antibodies used	Rabbit antibodies against GFP (Santa Cruz, sc-9996). Anti-rabbit IgG HRP conjugated (Thermo Scientific, cat. #31460).	
Validation	Shu, Y.N., et al. CKII-SIRT1-SM22α loop evokes a self-limited inflammatory response in vascular smooth muscle cells. Cardiovasc. Res. 113: 1198-1207 (2017). Marjanovic, M.P., et al. MacroH2A1.1 regulates mitochondrial respiration by limiting nuclear NAD+ consumption. Nat. Struct. Mol. Biol. 24: 902-910 (2017). Kang, J.W.M., et al. Resolving the contributions of anaesthesia, surgery, and nerve injury on brain derived neurotrophic factor expression in the medial prefrontal cortex of male rats in the CCI model of neuropathic pain. J. Neurosci. Res. 95: 2376-2390 (2017).	
Animals and	other organisms	
	about <u>studies involving animals</u> ; <u>ARRIVE guidelines</u> recommended for reporting animal research	
Laboratory	Elias ware obtained from the Drocephile stock center (https://hdcc.indiana.edu/) and used w1118 flies as the wild type central	

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

Laboratory animals

Flies were obtained from the Drosophila stock center (https://bdsc.indiana.edu/) and used w1118 flies as the wild-type control.

Wild animals

Wild animal was not involved in this study.

Field-collected samples

Sample collected from the field was not involved in this study.

Ethics oversight

No ethical approval.

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