nature portfolio

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Reporting Summary

Nature Portfolio 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 Portfolio policies, see our <u>Editorial Policies</u> and the <u>Editorial Policy Checklist</u>.

Statistics				
For all statistical analyse	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	ple size (n) for each experimental group/condition, given as a discrete number and unit of measurement			
A statement o	statement on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly			
The statistical Only common te	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	of all covariates tested			
A description of	of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons			
A full descripti	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 Give <i>P</i> values as exact values whenever suitable.				
For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings				
For hierarchical	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 c	ode			
Policy information abou	ut availability of computer code			
Data collection P	collection PER data were collected using visual observations.			
Data analysis FIJ	II freely accessible software for image quantifications and Graphpad Prism8 for statistical analyses were used.			
	om algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors and irage code deposition in a community repository (e.g. GitHub). See the Nature Portfolio 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 description of any restrictions on data availability
- For clinical datasets or third party data, please ensure that the statement adheres to our policy

Raw data, genotypes, full statistical analyses and p values are available in the source data file: https://doi.org/10.6084/m9.figshare.24972729.v1

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Policy information about studies with <u>human participants or human data</u>. See also policy information about <u>sex, gender (identity/presentation)</u>, <u>and sexual orientation</u> and <u>race</u>, <u>ethnicity</u> and <u>racism</u>.

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Reporting on sex	and gender	Not applicable		
Reporting on race	e, ethnicity, or	Not applicable		
other socially relevant				
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Population charac	characteristics Not applicable			
Recruitment	Not applicable			
Ethics oversight	Not applicable			
Note that full informa	tion on the appro	oval of the study protocol must also be provided in the manuscript.		
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Field-spe	cific re	porting		
Please select the or	ne below that is	the best fit for your research. If you are not sure, read the appropriate sections before making your selection.		
X Life sciences	iences Behavioural & social sciences Ecological, evolutionary & environmental sciences			
For a reference copy of t	he document with a	all sections, see <u>nature.com/documents/nr-reporting-summary-flat.pdf</u>		
Lifo scion	ococ cti	ıdy design		
All studies must disc	sclose on these points even when the disclosure is negative.			
Sample size	The sample size for each experiment is indicated in the source data file.			
Data exclusions	No data were excluded.			
Replication	The number of independent replicates is indicated in the source data file.			
Randomization	Flies used in this study were collected from different vials and generations.			
Blinding	Not applicable.			

Reporting for specific materials, systems and 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 & experim	ental systems Me	ethods		
n/a Involved in the study	y n/a	Involved in the study		
Antibodies	\boxtimes	ChIP-seq		
Eukaryotic cell line	es 🔀	Flow cytometry		
Palaeontology and	l archaeology 🔀	MRI-based neuroimaging		
Animals and other	organisms			
Clinical data				
Dual use research	of concern			
Plants				
'				
Animals and other	er research organism	NS STATE OF THE PROPERTY OF TH		
Policy information about <u>studies involving animals</u> ; <u>ARRIVE guidelines</u> recommended for reporting animal research, and <u>Sex and Gender in</u> <u>Research</u>				
Laboratory animals	Drosophila melanogaster. 5-7 days old adults for PER and GCaMP assyas were used.			
Wild animals	Not applicable			

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

Not applicable

CNRS

Reporting on sex

Ethics oversight

Field-collected samples

All experiments were performed with adult females.