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Last updated by author(s):	YYYY-MM-DD	

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 Authors & Referees and the Editorial Policy Checklist.

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FOL	ali St	atistical analyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.
n/a	Cor	nfirmed
	\boxtimes	The exact sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement
	\boxtimes	A statement on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly
	\boxtimes	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.
\boxtimes		A description of all covariates tested
\boxtimes		A description of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons
	\boxtimes	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)
	\boxtimes	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
\boxtimes		For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes
	\boxtimes	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

Data are collected with a custom setup. We used nichrome electrode from California Fine Wire Company (California Fine Wire Co, Grover Beach, CA, USA). The signal go through a manufactured headstage and theter connected to Neuralynx amplifiers (Neuralynx, Bozeman, MT, USA). The signal is stored with DataWave Sciworks acquisition software (DataWave Technologies, Loveland, CO, USA).

Data analysis

Spike sorting were made using Offline Sorter software (Plexon)

Data analysis were made using personal code (Matlab R2017b software), with following toolboxs: Signal Processing, Image Processing, Statistic and Machine Learning, Curve Fitting,

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 $\underline{guidelines}$ for submitting \underline{code} & $\underline{software}$ 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 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 from the corresponding author upon reasonable request.

Field-specific reporting						
Please select the o	ne below tha	It 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				
For a reference copy of t	the document w	ith all sections, see <u>nature.com/documents/nr-reporting-summary-flat.pdf</u>				
Life sciences study design						
All studies must dis	sclose on the	se points even when the disclosure is negative.				
Sample size	We choose s	sample size based on literatures in the field				
Data exclusions	No data wer	ere excluded from analysis				
Replication	N/A					
Randomization	There is no g	s no group. Data from all animals were pooled together because our comparison were made between conditions and no groups.				
Blinding	N/A					
D	C					
Reportin	g tor s	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 & exp	perimenta	l systems Methods				
n/a Involved in th	ne study	n/a Involved in the study				
Antibodies ChIP-seq		ChiP-seq				
Eukaryotic	Eukaryotic cell lines					
Palaeontology MRI-based neuroimaging						
Animals and other organisms						
Human research participants						
Clinical data						
Animals and	other o	rganisms				
Policy information about studies involving animals; ARRIVE guidelines recommended for reporting animal research						
Laboratory anima	als	Long Evans male rats from Janvier Labs (Le Genest-Saint-Isle, France), weighted 250-300 gr at their arrival				
Wild animals	Id animals The study did not involve wild animals					
Field-collected sa	amples	The study did not involve samples collected from the field				

All procedures complied with both European (directive 2010/63/EU of the European Parliament and of the Council) and French (AGRG1238767A) institutional guidelines.

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

Ethics oversight