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>.

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For	all statistical analyses, 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 statement on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly
	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.
X	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 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 <i>Give P values as exact values whenever suitable.</i>
X	For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings
X	For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes
X	Estimates of effect sizes (e.g. Cohen's <i>d</i> , Pearson's <i>r</i>), 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

Histological data were acquired using Zeiss Zen software (3.8). In vivo calcium imaging data were acquired using ThorCam Software (3.7.0). In vivo electrophysiology data were acquired using Intan Technologies Recording Controller (2.07), and the measured stimuli were synchronized and measured using Matlab (version R2019a or R2021a). Sexual reflexes of male mice were video recorded by SpinView (1.1.0.43). Mouse mating behaviors were recorded using IC capture (2.5).

Data analysis

Data were analyzed using Matlab version R2019a and R2021a, ImageJ (version 1.53t), Jupyter notebook (v. 6.5.4), Boris (v. 8.20.3), GraphPad Prism (Version 10), JRCLUST (version 3.2.2). We used a previously developed pipeline to analyze MEA data (https://github.com/ajemanuel/analyzeMEA). Custom scripts used in this study are posted on Github (https://github.com/NeuronQi/Krause_corpuscle).

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 and reviewers. We strongly encourage 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

The data generated in this study are available from the corresponding author upon request. Source data are provided with the paper.

Research involving human participants, their data, or biological material

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>.

Reporting on sex and gender	N/A
Reporting on race, ethnicity, or other socially relevant groupings	N/A
Population characteristics	N/A
Recruitment	N/A
Ethics oversight	N/A

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

Field-specific reporting

Please select the one below that is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.

	-			
\times	Life sciences	Behavioural & social sciences	Ecological, evolutionar	v & environmental science:

For a reference copy of the document with all sections, see $\underline{\mathsf{nature.com/documents/nr-reporting-summary-flat.pdf}}$

Life sciences study design

All studies must disclose on these points even when the disclosure is negative.

Sample size

No sample size calculation was performed. Sample sizes were based on previous studies from our lab and others (e.g., Neubarth et al., 2020 Science).

Data exclusions

For male sexual reflex testing, males showing excessive spontaneous erections after externalizing the glans penis were excluded. For female sexual reflex testing, females showing frequent whole-body movements that strongly disrupted the vaginal pressure measurements were excluded. During mating behavior, intromission bouts lasting less than two seconds were excluded from the analysis due to the lack of rhythmicity of movements and the possibility that penetration did not fully occur. For mating assay of naturally-cycled females, the females that did not show proestrus or estrus in 10 days were excluded.

Replication

We performed experiments with multiple animals to confirm reproducibility. All attempts at replication were successful. The number of replications is noted in the figure legends.

Randomization

Stimuli of different vibration frequencies during in vivo calcium imaging or electrophysiology were randomized. The allocation of mice of different genotypes into experimental groups was randomized.

Blinding

For histological experiments, images were collected and analyzed by investigators who were blinded to genotype whenever possible. The mating behaviors were scored by an investigator blinded to the genotype of the animals, and the genotype was only revealed after quantification and summary of the entire behavioral dataset. The experimenter was not blinded to the genotype of animals during in vivo eletrophysiological recordings, in vivo calcium imaging or sexual reflex tests, because only the animals of the desired genotypes were used in the experiments.

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.

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Materials & experime	ntal systems	Methods
n/a Involved in the study		n/a Involved in the study
Antibodies		ChIP-seq
Eukaryotic cell lines		Flow cytometry
Palaeontology and archaeology		MRI-based neuroimaging
Animals and other o	rganisms	
Clinical data		
Dual use research o	f concern	
Plants		
a call to		
<u>Antibodies</u>		
Antibodies used	chicken anti-GFP (Aves Labs	•
	goat anti-GFP (US Biological, goat anti-mCherry (CedarLa	•
	rabbit anti-CGRP (Immunost	
	chicken anti-NF200 (Aves La rabbit anti-NF200 (Sigma, N	, , , , , , , , , , , , , , , , , , , ,
	· ·	ch, 15146-1-AP, 1:200-1:500),
	rat anti-TROMA-1 (DSHB, AE goat anti-CD31 (R&D System	
) (custom-made, Handler et al., 2023)
	IB4 (Alexa 647 conjugated) (Invitrogen, 132450, 1:500).
Validation		s, GFP-1020), goat anti-GFP (US Biological,G8965-01E), goat anti-mCherry (CedarLane, AB0040-200),
	'	tar, 24112), chicken anti-NF200 (Aves Labs, NFH), rabbit anti-S100 (ProteinTech, 15146-1-AP) and guinea d in Handler et al., 2023 (PMID: 37725982);
	, , ,	I41422ML) and IB4 (Alexa 647 conjugated) (Invitrogen, I32450) were validated in Qi et al., 2024 (PMID:
	38442711); Rat anti-TROMA-1 (DSHB, Al	B 531826) was validated in Emanuel et al., 2021 (PMID: 34789880);
	goat anti-CD31 (R&D System	ns, AF3628) were validated in Ben-Zvi et al., 2014 (PMID: 24828040).
Animals and othe	r research organ	isms
Policy information about st	udies involving animals; A	RRIVE guidelines recommended for reporting animal research, and Sex and Gender in
Research		
Laboratory animals	All mice used in this study h	ave been previously described, including: TrkBCreER (JAX 027214), RetCreER (MGI 4437245), AdvillinCre
	(JAX 032536), AdvillinFlpO (0	Choi et al., Nature, 2020), RetCFP (MGI 3777555), TrkBflox(Liu et al., Science, 2012), Brn3acKOAP (JAX
		ert et al., Cell, 2021), MrgprdGFP (Zylka et al., Neuron, 2005), Th2A-CreER (RRID:IMSR_JAX:025614), . Nature, 2013), PLPEGFP (JAX 033357), Piezo2smFP-FLAG (Handler et al., Neuron, 2023), R26FSF-LSL-
	ReaChR-mCitrine (JAX 02484	46), Ai80 (R26FSF-LSL-CatCh; JAX 025109), Ai14 (R26LSL-tdTomato; JAX 007914), Ai65 (R26FSF-LSL-
		6 (R26LSL-GCaMP6s; Jax 028866), and Ai148 (TIGRELSL-GCaMP6f-tTA2, 030328). All lines were kept on a dyillinGre and TrkBflox were bred from mixed background to C57BI/6 background for two generations for

Both male and female mice were used for anatomical, physiological and behavioral experiments in this study.

Animals were handled according to protocols approved by the Harvard Standing Committee on Animal Care following the NIH Guide

for the Care and Use of Laboratory Animals. Mice were housed in a temperature-controlled and humidity-controlled facility,

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

maintained on a 12h light/dark cycle, and given food and water ad libitum.

No field-collected samples were used.

mating behavior testing.

No wild animals were used.

Wild animals

Reporting on sex

Ethics oversight

Field-collected samples