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Neuromantic (V1.6.3)

Matlab 2019b (MathWorks) Prism 6 (GraphPad) ImageJ (NIH)

Code: https://zenodo.org/record/4568820

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Reporting Summary

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Data analysis

For	all statistical anal	lyses, 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 statemen	t on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly				
		cal test(s) used AND whether they are one- or two-sided n tests should be described solely by name; describe more complex techniques in the Methods section.				
×	A description	on of all covariates tested				
	🗶 A descriptio	on of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons				
		ption of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient) on (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 Bayesia	n analysis, information on the choice of priors and Markov chain Monte Carlo settings				
X	For hierarch	nical and complex designs, identification of the appropriate level for tests and full reporting of outcomes				
	x Estimates o	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.				
So	ftware and	code				
Poli	cy information ab	pout <u>availability of computer code</u>				
Da	5	MED-PC IV (MedAssociates) Synapse s3 (TDT) pClamp 11 (Molecular Devices) Eclipse Ti software (Nikon)				

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 Research guidelines for submitting code & software for further information.

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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

Data availability statement in the manuscrip	t: "All	datasets are available up	oon reasonable request."
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x Life sciences	Behavioural & social sciences		Ecological, evolutionary & environmental sciences

For a reference copy of the document with all sections, see 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

Data

Sample sizes were determined based on power analysis, with expected variance and effect sizes based on literature (Chemogenetics/ behavior: e.g. Luchicchi et al., 2016, PMID 27630545; Koike et al., 2015, PMID 26224620 - Photometry: e.g. Kupferschmidt et al., 2017, PMID 29024667; Pisansky et al., 2019, PMID 31471038) and experience

Data exclusions

Animals with mistargeted viral expression (e.g. in case of fiber photometry, where there were no GCaMPm-expressing neurons under the fiber tip in the area where we would expect signal, as shown in Supplementary Figures or Kupferschmidt et al., 2017, PMID 29024667), without any virus expression, or with optic fiber implants not in the appropriate brain region were excluded in fiber photometry experiments. In the chemogenetics experiments, animals with unilateral or no DREADD expression were excluded. Prior perturbation studies in the 5-CSRTT have often chosen for bilateral manipulations (e.g Chudasama et al., 2003, PMID 14643464, or Koike et al., 2016, PMID 26224620), we chose to remain consistent with previous literature, especially in experiments where behavioral performance is the read-out. For neuroanatomical experiments, animals with mistargeted retrobead injections were excluded.

Replication

We validated the CombiCage so that behavioral performance would be stable and comparable to conventional 5-CSRTT (published in Bruinsma et al., 2019, PMID 30826849). Behavioral training and experiments (photometry, chemogenetics) were replicated by 3 different researchers with consistent outcomes. Slice electrophysiology experiments were performed by 2 different researchers with consistent outcomes. Virus injections for all experiments (photometry, chemogenetics, tracing, and ephys) were done by 4 different researchers yielded consistent outcomes in viral targeting and expression. Fiber implants were performed by two different researchers, targeting was consistent.

Randomization

In behavioral experiments, animals were randomly allocated to a 'projection target' group. CNO injection doses were varied randomly, as were variable delay and stimulus duration sessions. Within the sessions, delay and cue durations were varied pseudorandomly (so that the session would end up with a relatively similar number of trials in each condition).

Blinding

Blinding was not possible this study. Virus injections and implantations required highly targeted injections, precluding any blinding.

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 & experimental systems			Methods			
n/a	Involved in the study	n/a	Involved in the study			
	x Antibodies	×	ChIP-seq			
x	Eukaryotic cell lines	×	Flow cytometry			
x	Palaeontology and archaeology	x	MRI-based neuroimaging			
	X Animals and other organisms					
x	Human research participants					
x	Clinical data					
×	Dual use research of concern					

Antibodies

Antibodies used We used the following antibodies: mouse anti-NeuN (Abcam, Cat#104224, concentration 1:1000) with Alexa Fluor 647 donkey anti-

mouse (Cat#15980296, Thermo Fisher Scientific, 1:400), rabbit anti-RFP (Cat#600401379, Rockland, 1:1000) with Alexa Fluor 546 donkey anti-rabbit (Cat#10593125, Thermo Fisher Scientific, 1:400), mouse anti-GAD-67 (Cat#5406, Millipore, 1:1000) with Alexa Fluor 647 donkey anti-mouse (Cat#15980296, Thermo Fisher Scientific, 1:400), and rabbit anti-GFP (Abcam, 1:1000) with Alexa Fluor

488 donkey anti-rabbit (Cat#10424752, Thermo Fisher Scientific, 1:400)

Validation mouse anti-NeuN: 294 references (ex. PMID 31552908)

rabbit anti-RFP: >100 references (ex. PMID 32332079) mouse anti-GAD-67: 193 references (ex. PMID 24723034) rabbit anti-GFP: 894 references (ex. PMID 31859030)

Additionally, we included specific controls for each staining session (staining without either primary or secondary Ab)

Animals and other organisms

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

Laboratory animals Long Evans Rats, male, 6-32 weeks old

Wild animals No wild animals were used in this study.

Field-collected samples No field-collected samples were used in this study.

Ethics oversight All experimental procedures were in accordance with European and Dutch law and approved by the animal ethical care

committees of the VU University and VU University Medical Center, Amsterdam.

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