# 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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FOI	all statistical analyses, confirm that the following items are present in the figure regend, table regend, main text, of interhoos 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.
$\times$	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>
$\boxtimes$	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 <u>availability of computer code</u>

Data collection

Metamorph

Data analysis

ImageJ, MATLAB, Google Colab, Jupyter Notebooks, CellPose, https://codeocean.com/capsule/9982751/tree, https://github.com/sarahwaity/ProteiML, https://github.com/BerndtLab, https://doi.org/10.1101/2023.04.13.536801

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

All of the datasets generated within this study are available on figshare(ref 48). The files include data used for main figures, supplemental figures, and supplementary tables. We included the Chen (ref 4) and Dana (ref 5) datasets used to run our model and an amino acid property matrix derived from AAINDEX (ref

32) in the Suppleme (RCSB: 3SG3) and in		GCaMP crystal structure used in this manuscript is accessible online (https://www.rcsb.org/structure/3sg3), GCaMP3 D380Y ary Data.				
Human rese	earch part	icipants				
Policy information	about <u>studies i</u>	involving human research participants and Sex and Gender in Research.				
Reporting on sex	and gender	N/A				
Population chara	acteristics	N/A				
Recruitment		N/A				
Ethics oversight		N/A				
- ield-sp∈	ecific re	eporting				
Please select the o	ne below that i	is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.				
Life sciences	E	Behavioural & social sciences Ecological, evolutionary & environmental sciences				
or a reference copy of	the document with	n all sections, see <u>nature.com/documents/nr-reporting-summary-flat.pdf</u>				
ife scier	nces sti	udy design				
all studies must dis	sclose on these	e points even when the disclosure is negative.				
Sample size	Samples were each average.	Samples were taken at least until we reached statistical significance, and typically beyond. Most data sets include hundreds of samples for each average.				
Data exclusions		for automated cell segmentation: For cell segmentation, average sizes for cells were calculated. Regions of interest that were SD mean for the size of cells were excluded.				
Replication		ata point contains samples from at least 3 biological replicates, that is cell cultures wells. No technical replicates were takes and noved. The in vivo data contains data from 2 biological replicates (mice) under repeated time-locked stimulation, which is e field.				
Randomization	preparation of	The study was conducted with large numbers of samples and variants by individual researchers who required complete oversight from the preparation of experiments, to conduct, and analysis. We minimized research bias by including large sample sizes and an unbiased, automated image analysis algorithm. No outliers were removed and data was analyzed independently.				
Blinding	Test were not blinded because we had only one person taken measurments. However, data analysis was unbiased by an automated image analysis pipeline including a cell segmentation process (CellPose) which automatically identified all sensor expressing cells and extracted change in fluorescence from them.					
We require informati	ion from authors ited is relevant to perimental s	pecific materials, systems and methods s about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each materials of your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.  Systems  Methods  n/a Involved in the study				
Antibodies	5	ChIP-seq				
Eukaryotic	Eukaryotic cell lines Flow cytometry					

Materials & experimental 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 organisms	•		
Clinical data			
Dual use research of concern			
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Antibodies used	anti-GFP-AlexaFluor488 (Life Technology cat #A-21311)	
Validation	Manufacturer validated AB using western blots against positive and negative controls.	

# Eukaryotic cell lines

Policy information about cell lines and Sex and Gender in Research

Cell line source(s) HEK293, ATCC: CRL-1573

Authentication HEK293 cell were authenticated by ATCC prior to shipping with STR profiling following ISO 9001 and ISO/IEC 17025 quality standards. We discard cells after 25 passages and start new cultures from P2 or P3 frozen stocks.

Mycoplasma contamination HEK293 cells were tested negative for mycoplasma contamination

Commonly misidentified lines (See ICLAC register)

Commonly misidentified lines were not used

## Animals and other research organisms

Policy information about <u>studies involving animals</u>; <u>ARRIVE guidelines</u> recommended for reporting animal research, and <u>Sex and Gender in Research</u>

Laboratory animals	6–7-week-old male and female C57BL/6J mice (JAX) were used for in vivo studies. Cultured neurons were dissected from P0 rat pubs (female and male, Sprague Dawley, Envigo).
Wild animals	N/A
Reporting on sex	Male and female mice were used.
Field-collected samples	N/A
Ethics oversight	All experiments were conducted in accordance with UC Davis's and University of Washington's Institutional Animal Care and Use Committees.

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

# Dual use research of concern

Policy information about <u>dual use research of concern</u>

#### Hazards

Could the accidental, deliberate or reckless misuse of agents or technologies generated in the work, or the application of information presented in the manuscript, pose a threat to:

Vo	Yes
X	Public health
X	National security
X	Crops and/or livestock
X	Ecosystems
$\boxtimes$	Any other significant area

# Experiments of concern

Doe	s the work involve any of these experiments of concern:
No	Yes
$\boxtimes$	Demonstrate how to render a vaccine ineffective
$\boxtimes$	Confer resistance to therapeutically useful antibiotics or antiviral agents
$\boxtimes$	Enhance the virulence of a pathogen or render a nonpathogen virulent
$\boxtimes$	Increase transmissibility of a pathogen
$\boxtimes$	Alter the host range of a pathogen
$\boxtimes$	Enable evasion of diagnostic/detection modalities
$\boxtimes$	Enable the weaponization of a biological agent or toxin
$\boxtimes$	Any other potentially harmful combination of experiments and agents