

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 [Editorial Policies](#) and the [Editorial Policy Checklist](#).

Please do not complete any field with "not applicable" or n/a. Refer to the help text for what text to use if an item is not relevant to your study.

For final submission: please carefully check your responses for accuracy; you will not be able to make changes later.

Statistics

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 |
|----------------------------------|---|
| <input checked="" type="radio"/> | <input checked="" type="radio"/> The exact sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement |
| <input checked="" type="radio"/> | <input checked="" type="radio"/> A statement on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly |
| <input checked="" type="radio"/> | <input checked="" type="radio"/> The statistical test(s) used AND whether they are one- or two-sided
<i>Only common tests should be described solely by name; describe more complex techniques in the Methods section.</i> |
| <input checked="" type="radio"/> | <input checked="" type="radio"/> A description of all covariates tested |
| <input checked="" type="radio"/> | <input checked="" type="radio"/> A description of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons |
| <input checked="" type="radio"/> | <input checked="" type="radio"/> 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) |
| <input checked="" type="radio"/> | <input checked="" type="radio"/> For null hypothesis testing, the test statistic (e.g. F , t , r) with confidence intervals, effect sizes, degrees of freedom and P value noted
<i>Give P values as exact values whenever suitable.</i> |
| <input checked="" type="radio"/> | <input checked="" type="radio"/> For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings |
| <input checked="" type="radio"/> | <input checked="" type="radio"/> For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes |
| <input checked="" type="radio"/> | <input checked="" type="radio"/> Estimates of effect sizes (e.g. Cohen's d , Pearson's r), indicating how they were calculated |

Our web collection on [statistics for biologists](#) contains articles on many of the points above.

Software and code

Policy information about [availability of computer code](#)

Data collection	Motic Image Plus 2.0, Leica application suite X version 3.5.6.21594 for SP8, Leica application suite X version 3.7.1.21655 for DMi8, ZEISS Efficient
Data analysis	Data was analyzed using: ImageJ (v1.8.0_112), Imaris x64 version 9.5.0, Adobe Photoshop (version 21.2.3)

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 data generated and processed are included in this article, with additional raw data available from the corresponding authors upon reasonable requests. There are no restrictions on data availability.

Human research participants

Policy information about studies involving human research participants and Sex and Gender in Research.

Reporting on sex and gender	Three female fibroblast cell lines were used in this study protocol and they were sourced commercially.
Population characteristics	NA
Recruitment	No human research participants recruitment were performed in this study.
Ethics oversight	The initial experiments using the three commercially sourced fibroblast cell lines were performed first under the approval of the

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.

☒ Life sciences ☐ Behavioural & social sciences ☐ Ecological, evolutionary & environmental sciences

Life sciences study design

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

Sample size	We did not involve statistical methods to pre-determine the sample size, this was determined based on previous experience and other similar
Data exclusions	No data were excluded.
Replication	For the data demonstrated in this study, we have reproduced them from 3 different biologically independent cell lines with experimental
Randomization	The experiments were not randomized as the cell lines utilized in this article were sourced commercially.
Blinding	The investigators were not blinded during data collection and analysis, as neither specific grouping were involved in this manuscript.

Behavioural & social sciences study design

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

Study description	
Research sample	
Sampling strategy	
Data collection	
Timing	
Data exclusions	
Non-participation	
Randomization	

Ecological, evolutionary & environmental sciences study design

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

Study description	
Research sample	
Sampling strategy	
Data collection	
Timing and spatial scale	
Data exclusions	
Reproducibility	
Randomization	

Blinding

Did the study involve field work? ☐ Yes ☐ No

Field work, collection and transport

Field conditions

Location

Access & import/export

Disturbance

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

n/a ☐ Involved in the study

☒ Antibodies

☒ Eukaryotic cell lines

☐ Palaeontology and archaeology

☐ Animals and other organisms

☐ Clinical data

☐ Dual use research of concern

Methods

n/a ☐ Involved in the study

☒ ChIP-seq

☐ Flow cytometry

☒ MRI-based neuroimaging

Antibodies

Antibodies used Details of all antibodies used in this article were provided with catalog number, lot number and commercial sources supplied as below:

Validation Antibodies obtained from the commercial source were validated by the suppliers, detailed validation analysis relevant literatures are

Eukaryotic cell lines

Policy information about [cell lines](#) and [Sex and Gender in Research](#)

Cell line source(s) Human fibroblasts sourced from ThermoFisher (Catalogue number, C-013-5C; lot#1029000 for 38F; lot#1528526 for 55F and

Authentication Human dermal fibroblasts were derived from tissues of donor (age and sex identified) and authenticated by ThermoFisher (via

Mycoplasma contamination Fibroblasts lines were tested by ThermoFisher. Furthermore, cell lines were regularly tested and were mycoplasma negative.

Commonly misidentified lines (See [ICLAC](#) register) No commonly misidentified cell lines were used in this study.

Palaeontology and Archaeology

Specimen provenance

Specimen deposition

Dating methods

☐ Tick this box to confirm that the raw and calibrated dates are available in the paper or in Supplementary Information.

Ethics oversight

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

Animals and other research organisms

Policy information about [studies involving animals](#); [ARRIVE guidelines](#) recommended for reporting animal research, and [Sex and Gender in Research](#)

Laboratory animals	
Wild animals	
Reporting on sex	
Field-collected samples	
Ethics oversight	

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

Clinical data

Policy information about [clinical studies](#)

All manuscripts should comply with the ICMJE [guidelines for publication of clinical research](#) and a completed [CONSORT checklist](#) must be included with all submissions.

Clinical trial registration	
Study protocol	
Data collection	
Outcomes	

Dual use research of concern

Policy information about [dual use research of concern](#)

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:

No	Yes
<input type="radio"/>	<input checked="" type="radio"/> Public health
<input type="radio"/>	<input checked="" type="radio"/> National security
<input type="radio"/>	<input checked="" type="radio"/> Crops and/or livestock
<input type="radio"/>	<input checked="" type="radio"/> Ecosystems
<input type="radio"/>	<input checked="" type="radio"/> Any other significant area

Experiments of concern

Does the work involve any of these experiments of concern:

No	Yes
<input type="radio"/>	<input checked="" type="radio"/> Demonstrate how to render a vaccine ineffective
<input type="radio"/>	<input checked="" type="radio"/> Confer resistance to therapeutically useful antibiotics or antiviral agents
<input type="radio"/>	<input checked="" type="radio"/> Enhance the virulence of a pathogen or render a nonpathogen virulent
<input type="radio"/>	<input checked="" type="radio"/> Increase transmissibility of a pathogen
<input type="radio"/>	<input checked="" type="radio"/> Alter the host range of a pathogen
<input type="radio"/>	<input checked="" type="radio"/> Enable evasion of diagnostic/detection modalities
<input type="radio"/>	<input checked="" type="radio"/> Enable the weaponization of a biological agent or toxin
<input type="radio"/>	<input checked="" type="radio"/> Any other potentially harmful combination of experiments and agents

ChIP-seq

Data deposition

☐ Confirm that both raw and final processed data have been deposited in a public database such as [GEO](#).

☐ Confirm that you have deposited or provided access to graph files (e.g. BED files) for the called peaks.

Data access links <i>May remain private before publication</i>	
Files in database submission	
Genome browser session (e.g. UCSC)	

Methodology

Replicates	<input type="text"/>
Sequencing depth	<input type="text"/>
Antibodies	<input type="text"/>
Peak calling parameters	<input type="text"/>
Data quality	<input type="text"/>
Software	<input type="text"/>

Flow Cytometry

Plots

Confirm that:

- ☐ The axis labels state the marker and fluorochrome used (e.g. CD4-FITC).
- ☐ The axis scales are clearly visible. Include numbers along axes only for bottom left plot of group (a 'group' is an analysis of identical markers).
- ☐ All plots are contour plots with outliers or pseudocolor plots.
- ☐ A numerical value for number of cells or percentage (with statistics) is provided.

Methodology

Sample preparation	<input type="text"/>
Instrument	<input type="text"/>
Software	<input type="text"/>
Cell population abundance	<input type="text"/>
Gating strategy	<input type="text"/>

☐ Tick this box to confirm that a figure exemplifying the gating strategy is provided in the Supplementary Information.

Magnetic resonance imaging

Experimental design

Design type	<input type="text"/>
Design specifications	<input type="text"/>
Behavioral performance measures	<input type="text"/>

Acquisition

Imaging type(s)	<input type="text"/>
Field strength	<input type="text"/>
Sequence & imaging parameters	<input type="text"/>
Area of acquisition	<input type="text"/>
Diffusion MRI	<input checked="" type="radio"/> Used <input type="radio"/> Not used

Preprocessing

Preprocessing software	<input type="text"/>
Normalization	<input type="text"/>
Normalization template	<input type="text"/>
Noise and artifact removal	<input type="text"/>
Volume censoring	<input type="text"/>

Statistical modeling & inference

Model type and settings	<input type="text"/>
Effect(s) tested	<input type="text"/>
Specify type of analysis:	<input checked="" type="radio"/> Whole brain <input type="radio"/> ROI-based <input type="radio"/> Both
Statistic type for inference (See Eklund et al. 2016)	<input type="text"/>
Correction	<input type="text"/>

Models & analysis

n/a	Involvement in the study
<input type="checkbox"/>	Functional and/or effective connectivity
<input type="checkbox"/>	Graph analysis
<input type="checkbox"/>	Multivariate modeling or predictive analysis
	Functional and/or effective connectivity
	Graph analysis
	Multivariate modeling and predictive analysis

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