# nature portfolio

Corresponding author(s):	Susumu Goyama
Last updated by author(s):	Oct 18, 2023

### **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>.

< ∙	トつ	1	101	- 1	$\sim$
. )	ıa	ш	וכו		CS

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.
$\boxtimes$	A description of all covariates tested
X	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>
$\times$	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
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

Provide a description of all commercial, open source and custom code used to collect the data in this study, specifying the version used OR state that no software was used.

Data analysis

Provide a description of all commercial, open source and custom code used to analyse the data in this study, specifying the version used OR state that no software was used.

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 <u>guidelines for submitting code & software</u> 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

We deposited all the RNA-seq data into GEO.

#### Human research participants

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

Reporting on sex and gender

Use the terms sex (biological attribute) and gender (shaped by social and cultural circumstances) carefully in order to avoid confusing both terms. Indicate if findings apply to only one sex or gender; describe whether sex and gender were considered in study design whether sex and/or gender was determined based on self-reporting or assigned and methods used. Provide in the source data disaggregated sex and gender data where this information has been collected, and consent has been obtained for sharing of individual-level data; provide overall numbers in this Reporting Summary. Please state if this information has not been collected. Report sex- and gender-based analyses where performed, justify reasons for lack of sex- and gender-based analysis.

Population characteristics

Describe the covariate-relevant population characteristics of the human research participants (e.g. age, genotypic information, past and current diagnosis and treatment categories). If you filled out the behavioural & social sciences study design questions and have nothing to add here, write "See above."

Recruitment

Describe how participants were recruited. Outline any potential self-selection bias or other biases that may be present and how these are likely to impact results.

Ethics oversight

Identify the organization(s) that approved the study protocol.

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	w 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

For a reference copy of the document with all sections, see <u>nature.com/documents/nr-reporting-summary-flat.pdf</u>

#### Life sciences study design

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

Sample size	Sample size was decided on the basis of our previous experience in the field, not predetermined by a statistical method.
Data exclusions	No data were excluded.
Replication	All data were confirmed by at least one biologically independent experiments
Randomization	In animal experiments, mice were randomly assigned to several groups.
Blinding	Experiments were not blinded.

# Behavioural & social sciences study design

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

Study description

Briefly describe the study type including whether data are quantitative, qualitative, or mixed-methods (e.g. qualitative cross-sectional, quantitative experimental, mixed-methods case study).

Research sample

State the research sample (e.g. Harvard university undergraduates, villagers in rural India) and provide relevant demographic information (e.g. age, sex) and indicate whether the sample is representative. Provide a rationale for the study sample chosen. For studies involving existing datasets, please describe the dataset and source.

Sampling strategy

Describe the sampling procedure (e.g. random, snowball, stratified, convenience). Describe the statistical methods that were used to predetermine sample size OR if no sample-size calculation was performed, describe how sample sizes were chosen and provide a rationale for why these sample sizes are sufficient. For qualitative data, please indicate whether data saturation was considered, and what criteria were used to decide that no further sampling was needed.

Data collection

Provide details about the data collection procedure, including the instruments or devices used to record the data (e.g. pen and paper, computer, eye tracker, video or audio equipment) whether anyone was present besides the participant(s) and the researcher, and whether the researcher was blind to experimental condition and/or the study hypothesis during data collection.

Timing	Indicate the start and stop dates of data collection. If there is a gap between collection periods, state the dates for each sample cohort.
Data exclusions	If no data were excluded from the analyses, state so OR if data were excluded, provide the exact number of exclusions and the rationale behind them, indicating whether exclusion criteria were pre-established.
Non-participation	State how many participants dropped out/declined participation and the reason(s) given OR provide response rate OR state that no participants dropped out/declined participation.

If participants were not allocated into experimental groups, state so OR describe how participants were allocated to groups, and if

# Ecological, evolutionary & environmental sciences study design

allocation was not random, describe how covariates were controlled.

Randomization

Study description	Briefly describe the study. For quantitative data include treatment factors and interactions, design structure (e.g. factorial, nested, hierarchical), nature and number of experimental units and replicates.
Research sample	Describe the research sample (e.g. a group of tagged Passer domesticus, all Stenocereus thurberi within Organ Pipe Cactus National Monument), and provide a rationale for the sample choice. When relevant, describe the organism taxa, source, sex, age range and any manipulations. State what population the sample is meant to represent when applicable. For studies involving existing datasets, describe the data and its source.
Sampling strategy	Note the sampling procedure. Describe the statistical methods that were used to predetermine sample size OR if no sample-size calculation was performed, describe how sample sizes were chosen and provide a rationale for why these sample sizes are sufficient.
Data collection	Describe the data collection procedure, including who recorded the data and how.
Timing and spatial scale	Indicate the start and stop dates of data collection, noting the frequency and periodicity of sampling and providing a rationale for these choices. If there is a gap between collection periods, state the dates for each sample cohort. Specify the spatial scale from which the data are taken
Data exclusions	If no data were excluded from the analyses, state so OR if data were excluded, describe the exclusions and the rationale behind them, indicating whether exclusion criteria were pre-established.
Reproducibility	Describe the measures taken to verify the reproducibility of experimental findings. For each experiment, note whether any attempts to repeat the experiment failed OR state that all attempts to repeat the experiment were successful.
Randomization	Describe how samples/organisms/participants were allocated into groups. If allocation was not random, describe how covariates were controlled. If this is not relevant to your study, explain why.
Blinding	Describe the extent of blinding used during data acquisition and analysis. If blinding was not possible, describe why OR explain why blinding was not relevant to your study.
Did the study involve fiel	d work? Yes No

Field conditions	Describe the study conditions for field work, providing relevant parameters (e.g. temperature, rainfall).
Location	State the location of the sampling or experiment, providing relevant parameters (e.g. latitude and longitude, elevation, water depth).
Access & import/export	Describe the efforts you have made to access habitats and to collect and import/export your samples in a responsible manner and in compliance with local, national and international laws, noting any permits that were obtained (give the name of the issuing authority, the date of issue, and any identifying information).
Disturbance	Describe any disturbance caused by the study and how it was minimized.

### 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 & experime	ntal sy	ystems Methods			
n/a Involved in the study n/a Involved in the study					
Antibodies		ChIP-seq			
Eukaryotic cell lines		Flow cytometry			
Palaeontology and a	Palaeontology and archaeology MRI-based neuroimaging				
Animals and other o	organisms				
Clinical data					
Dual use research of	f concerr				
Antibodies					
Antibodies used	The info	ormation is provided as Supplementary tables.			
Validation	Yes, we	validated it.			
Eukaryotic cell lin	es				
Policy information about <u>ce</u>	ell lines a	and Sex and Gender in Research			
Cell line source(s)		The information is provided in Methods section.			
Authentication		Cells are gotten from ATCC.			
Mycoplasma contaminati	on (	Cells were not tested for mycoplasma contamination.			
Commonly misidentified I (See <u>ICLAC</u> register)	lines	We did not use commonly misidentified cell lines in this study.			
Animals and othe	r rese	earch organisms			
Policy information about <u>str</u> <u>Research</u>	<u>udies in</u>	volving animals; ARRIVE guidelines recommended for reporting animal research, and Sex and Gender in			
Laboratory animals		pe (WT) C57BL/6J mice and C57BL/6J-Rag2em3Lutzy/J (Rag2-/-) mice were purchased from Sankyo Labo Service corporation . NSG mice were purchased from Charles River Laboratories Japan. All experiments were performed with 8—12-week-old mice.			
Wild animals	This stu	dy did not use wild animals.			
Reporting on sex	All the i	nformation about sex of the mice used are provided in the figure legends.			
Field-collected samples	This stu	dy did not involve the samples collected from Fields.			
Ethics oversight  All animal studies were approved by the Animal Care Committee of the Institute of Medical Science at the University of Tokyo (approval number: PA15-100, PA18-46), and were conducted in accordance with the Regulation on Animal Experimentation at University of Tokyo based on International Guiding Principles for Biomedical Research Involving Animals.					
Note that full information on the	he appro	val of the study protocol must also be provided in the manuscript.			
Flow Cytometry					
Plots					
Confirm that:					
<u></u>	he mark	er and fluorochrome used (e.g. CD4-FITC).			
		ble. Include numbers along axes only for bottom left plot of group (a 'group' is an analysis of identical markers).			
		th outliers or pseudocolor plots.			
		r of cells or percentage (with statistics) is provided.			
Methodology					
Sample preparation We described it in Methods section.					
Instrument	We described it in Methods section.				

	$\overline{}$	í
	Ξ	<b>+</b>
	C	t
	<u>=</u>	
	Œ	)
	~	5
	≍	ί
	⋍	ζ.
	_	
	$\overline{}$	₹
		<u>_</u>
	=	ζ,
	_	
Z		
	_	5
_	π	5
	_ α	5
	≍	5
	Č	
	Č	700
	<u> </u>	100
	<u> </u>	
Ĺ	<u> </u>	
	<u> </u>	200
Ĺ	<u> </u>	2007-200-2
Ĺ	<u> </u>	
Ĺ	<u> </u>	
Ú		
Ĺ		

≤	ŝ	
۵		
כ		
2		
₹		
_		

Software	We described it in Methods section.	
Cell population abundance	We provided the information as supplemental information.	
Gating strategy	We provided the information as supplemental information.	
Tick this box to confirm that a figure exemplifying the gating strategy is provided in the Supplementary Information.		