# 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
$\boxtimes$	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
$\boxtimes$	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 statistics for biologists contains articles on many of the points above.

#### Software and code

Policy information about availability of computer code

Data collection

Agilent 1290II-6460, Shimadzu TQ8050 NX, NanoDrop ND-1000 UV-Vis Spectrophotometer, 5-L fermenter (T&J Bio-engineering, Shanghai Co., LTD), Glucose-lactate analyzer (M100, Shenzhen Sieman Technology Co., Ltd), ReadMax 1900Plus Light absorption full wavelength microplate reader, 600 MBruker Avance NEO 600M NMR Spectroscopy.

Data analysis

Origin 2021 64bit, Microsoft Excel 2016, Microsoft Word 2016, Microsoft PowerPoint 2016, Adobe Illustrate, SPSS v13.0, Pymol 2.4.0, Gaussian 16, Gaussian view, AutoDock Vina, ChemSketch 2021, Mega-X, Gromacs 2022, Amber 18, MestReNova v14.0.

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

Data supporting the findings of this work are available within the manuscript and supplementary files. The datasets generated and analyzed during the current study are also available from the corresponding author.

#### 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, ethnicity and racism</u>.

This is not applicable in this study.	
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Note that full information on the approval of the study protocol must also be provided in the manuscript.

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

### Field-specific reporting

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

No sample-size calculations were performed. As indicated in the text, all experiments were performed from a single colony, which provides some limited information about the distribution of the measurements and is typical of similar experiments and studies in the field.

Data exclusions

No data were excluded form the analyses.

All the biochemical and biological experiments were performed in three replicated or more. Data were repeatable on different date.

Randomization The cells are uniformly and randomly distributed in the medium and buffer. Pipet tips were used to scoop few cells for culturing, the scooping locations are all random.

locations are all random

Blinding

Blinding is not relevant to our study because none of our data is based on qualitative scoring metrics nor does it involve animals or human research participants. As described in the above section for randomization, blinding during group allocation is irrelevant because the samples of cell cultures that were split into different conditions were random samplings and there is no control over which cells will be selected and thus, no bias during group allocation.

### 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.

Ma	terials & experime	ntal systems Me	ethods
n/a	Involved in the study	n/a	Involved in the study
$\boxtimes$	Antibodies	$\boxtimes$	ChIP-seq
$\times$	Eukaryotic cell lines	$\boxtimes$	Flow cytometry
$\times$	Palaeontology and a	rchaeology	MRI-based neuroimaging
$\times$	Animals and other or	rganisms	
$\times$	Clinical data		
$\times$	Dual use research of concern		
$\boxtimes$	Plants		
Plants			
Se	ed stocks	This is not applicable in this study	y.
No	vel plant genotypes	This is not applicable in this study	y.

Authentication

This is not applicable in this study.