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Reporting Summary

Statistics

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>.

For	all sta	atistical an	alyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.			
n/a	Con	firmed				
\boxtimes		The exact	sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement			
\times		A stateme	ent on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly			
\boxtimes	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 descript	ion of all covariates tested			
\boxtimes		A descript	ion of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons			
\boxtimes	11 1	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)				
\boxtimes	11 1	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.				
\boxtimes		For Bayes	ian 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.					
So	ftw	are an	d code			
Poli	cy inf	ormation a	about <u>availability of computer code</u>			
Data collection N/A		ollection	N/A			
Data analysis N/A		nalysis	N/A			

Data

Policy information about <u>availability of data</u>

All manuscripts must include a <u>data availability statement</u>. This statement should provide the following information, where applicable:

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.

- 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

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	Research involving hu	man participants,	their data,	or biological	material
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and sexual orientation and <u>rac</u>	e, edifficitly and racism.		
Reporting on sex and gender	N/A		
Reporting on race, ethnicity, other socially relevant groupings	or N/A		
Population characteristics	N/A		
Recruitment	N/A		
Ethics oversight	N/A		
Note that full information on the a	pproval of the study protocol must also be provided in the manuscript.		
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Please select the one below th	at is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.		
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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		
\boxtimes	Antibodies	\boxtimes	ChIP-seq		
\boxtimes	Eukaryotic cell lines	\boxtimes	Flow cytometry		
\boxtimes	Palaeontology and archaeology	\boxtimes	MRI-based neuroimaging		
\boxtimes	Animals and other organisms				
\boxtimes	Clinical data				
\boxtimes	Dual use research of concern				
\boxtimes	Plants				