## nature research

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## **Reporting Summary**

No datasets were generated or analyzed for this study.

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Statistics					
For all statistical an	alyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.				
n/a Confirmed					
The exact	ct sample size $(n)$ for each experimental group/condition, given as a discrete number and unit of measurement				
A stateme	tement 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.				
A descript	A description of all covariates tested				
A descript	ription 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 Give <i>P</i> values as exact values whenever suitable.					
For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings					
For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes					
Estimates of effect sizes (e.g. Cohen's $d$ , Pearson's $r$ ), indicating how they were calculated					
1	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	No software was used in data collection for this study.				
Data analysis	The protocols described herein use the Lathe software, found at https://github.com/bhattlab/lathe.				
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 Research guidelines for submitting code & software for further information.					
Data					
All manuscripts m - Accession codes - A list of figures	about <u>availability of data</u> ust include a <u>data availability statement</u> . This statement should provide the following information, where applicable: s, unique identifiers, or web links for publicly available datasets that have associated raw data				

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All studies must dis	sclose on these points even when t	he disclosure is negative.		
Sample size	n/a			
Data exclusions	n/a			
Replication	n/a			
Randomization	n/a			
Blinding	n/a			
<b>D</b>				
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Palaeontology and archaeology MRI-based neuroimaging		MRI-based neuroimaging		
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Human research participants

Clinical data

Dual use research of concern