# 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>.

For all statistical analyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.

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n/a	Confirmed
	$\square$ 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
$\boxtimes$	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>
$\boxtimes$	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 <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 The code used to generate the main

The code used to generate the main results in this manuscript can be found in the following github repository:  $https://github.com/khaledsaab/eeg\_robustness$ 

Data analysis

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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

The Stanford clinical datasets used in this study are subject to restrictions regarding the availability of Protected Health Information. They were accessed with approval from the Institutional Review Board solely for the purpose of this specific study and are not accessible to the public.

		man participants, their data, or biological material		
		vith <u>human participants or human data</u> . See also policy information about <u>sex, gender (identity/presentation),</u> thnicity and racism.		
Reporting on sex and gender		We expand on this in the Methods section of our paper.		
Reporting on race, ethnicity, or other socially relevant groupings		We expand on this in the Methods section of our paper.		
Population charac	cteristics	We expand on this in the Methods section of our paper.		
Recruitment		We expand on this in the Methods section of our paper.		
Ethics oversight		We expand on this in the Methods section of our paper.		
lote that full informa	tion on the appr	oval of the study protocol must also be provided in the manuscript.		
Field-spe	citic re	porting		
lease select the or	ne below that is	s the best fit for your research. If you are not sure, read the appropriate sections before making your selection.		
Life sciences	В	ehavioural & social sciences		
or a reference copy of t	he document with	all sections, see <u>nature.com/documents/nr-reporting-summary-flat.pdf</u>		
ife scien	ices stu	udy design		
Il studies must dis	close on these	points even when the disclosure is negative.		
Sample size	We expand on t	this in the Methods section of our paper.		
Data exclusions	We expand on this in the Methods section of our paper.			
Replication	We expand on this in the Methods section of our paper.			
Randomization	We expand on this in the Methods section of our paper.			
Blinding	We expand on this in the Methods section of our paper.			
Reportin	g for sr	pecific materials, systems and methods		
Ve require information	on from authors	about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each materi your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response		
Materials & exp	perimental s	ystems Methods		
n/a Involved in the study		n/a Involved in the study		

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		

## Plants

Seed stocks	N/A
Novel plant genotypes	N/A
Authentication	N/A