nature portfolio

Corresponding author(s):	Matthew Patterson
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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 Editorial Policies and the Editorial Policy Checklist.

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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
	\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
\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.
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
\boxtimes	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	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.
So	ftware and code
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Policy information about availability of computer code

Open source data was used in this study that was in csv format Data collection

Data analysis

Custom scripts in Python 3.9.0 were used for the analysis of this code. This code is open-sourced and is available in the Code Availability

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 Newcastle Polysomnography dataset can be found at https://zenodo.org/record/1160410#.YynNvOzMJz8

Human rese	arch parti	cipants		
Policy information	about <u>studies i</u>	nvolving human research participants and Sex and Gender in Research.		
Reporting on sex	and gender	Sex based analysis was not performed because of the small sample size.		
Population chara	cteristics	20 of the 28 participants had a sleep disorder. Sleep disorders included idiopathic hypersomnia, restless leg syndrome, sleep apnea, narcolepsy, sleep paralysis, nocturia, obstructive sleep apnea, REM sleep disorder, parasomnia and insomnia.		
Recruitment		Patients who were scheduled for a one-night polysomnography assessment at the Freeman Hospital, Newcastle upon Tyne, UK, as part of their routine clinical assessment were invited to participate in the study.		
Ethics oversight		NRES North East Sunderland Ethics Committee		
Note that full informa	ation on the appr	roval of the study protocol must also be provided in the manuscript.		
Life sciences	ne below that i	s the best fit for your research. If you are not sure, read the appropriate sections before making your selection. Behavioural & social sciences		
Life scier	nces sti	udy design		
All studies must dis	sclose on these	points even when the disclosure is negative.		
Sample size	28			
Data exclusions	No data exclusi	ons		
Replication	Analysis code v	Analysis code was peer reviewed on GitHub		
Randomization	Group allocation	Group allocation is not relevant to this study because it is a validation study		
Blinding	Blinding is not relevant to this study because this is a validation study			
		pecific materials, systems and methods		
system or method list	ted is relevant to	about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material, 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	·	·		
<u>'_</u> '		n/a Involved in the study ☐ ChIP-seq		
Eukaryotic cell lines		Flow cytometry		
Palaeontology and archaeology		logy MRI-based neuroimaging		

Dual use research of concern

Clinical data

Animals and other organisms

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