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Reporting Summary

Nature Research 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 Research policies, see <u>Authors & Referees</u> and the <u>Editorial Policy Checklist</u>.

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	istics				
. 1		es, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.			
_	a Confirmed				
	★ The exact sam	ple size (n) for each experimental group/condition, given as a discrete number and unit of measurement			
$\boxtimes $	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.				
	A description of all covariates tested				
	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>				
	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 <i>d</i> , Pearson's <i>r</i>), indicating how they were calculated				
ı		Our web collection on <u>statistics for biologists</u> contains articles on many of the points above.			
Soft	ware and c	ode			
		ut <u>availability of computer code</u>			
	a collection	No software was used.			
Dat	a analysis	The algorithm C Cut is implemented in Matleh and available for non-commercial uses https://muwazhu@hithusket.org/muwazhu/gaut.git			
	Data analysis The algorithm G-Cut is implemented in Matlab and available for non commercial use: https://muyezhu@bitbucket.org/muyezhu/gcut.git				
		deposition in a community repository (e.g. GitHub). See the Nature Research guidelines for submitting code & software for further information.			
Dat	a				
All m - 4 - 4	nanuscripts must i Accession codes, uni A list of figures that l	ut <u>availability of data</u> nclude a <u>data availability statement</u> . This statement should provide the following information, where applicable: ique identifiers, or web links for publicly available datasets have associated raw data			
- A description of any restrictions on data availability The raw image data generated for the study (shown in Figure 1, 6, 7 and Supplementary Figure 6) are available from corresponding authors.					
ine ra	aw iiiiage data gene	rated for the study (shown in rigure 1, 0, 7 and supplementary rigure 6) are available from corresponding authors.			
Fie	ld-speci	fic reporting			
Please	e select the one b	elow that is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.			
∑ Lif	e sciences	Behavioural & social sciences Ecological, evolutionary & environmental sciences			

For a reference copy of the document with all sections, see $\underline{\mathsf{nature}.\mathsf{com}/\mathsf{documents}/\mathsf{nr}-\mathsf{reporting}-\mathsf{summary-flat}.\mathsf{pdf}}$

Life sciences study design

Materials & experimental systems

All studies must disclose on these points even when the disclosure is negative.			
Sample size	No sample size calculation was conducted, however the sample sizes in statistical analyses were greater than or comparable to those of typical life science publications.		
Data exclusions	No data were excluded.		
Replication	Accuracy of the g-cut algorithm was compared to alternative algorithms in various image stacks belonging to multiple brain regions, using multiple labeling strategy and collected with multiple image modalities. We found g-cut to be superior in all those image stacks.		
Randomization	Randomization was not applicable in this study. The g-cut algorithm needed to be evaluated against alternative algorithms on exact same input data.		
Blinding	Blinding was not applicable in this study. No human subjects were used.		

Reporting for specific materials, systems and methods

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.

n/a Involved in the study	n	n/a Involved in the study
Antibodies	!	ChIP-seq
Eukaryotic cell lines	!	Flow cytometry
Palaeontology	!	MRI-based neuroimaging
Animals and other org	ganisms	'
Human research parti	cipants	
Clinical data		
•		
Animals and other	organisms	
Policy information about stu	dies involving animals; ARF	RIVE guidelines recommended for reporting animal research
Laboratory animals (1) Mus musculus, C57		/6, male, 8 weeks
,	(2) Mus musculus, MORF	(Lu & Yang, 2017), male, 8 weeks
Wild animals	The study did not involve wild animals.	
Field-collected samples	The study did not involve samples collected from field.	
Ethics oversight	Identify the organization(:	s) that approved or provided guidance on the study protocol, OR state that no ethical approval or

Note that full information on the approval of the study protocol must also be provided in the manuscript.

guidance was required and explain why not.