

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](#).

Statistics

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

- 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.
- 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. F , t , r) with confidence intervals, effect sizes, degrees of freedom and P value noted
Give P 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

Our web collection on [statistics for biologists](#) contains articles on many of the points above.

Software and code

Policy information about [availability of computer code](#)

Data collection

We used intraoperative photography, intraoperative video recordings, clinical records (Dxcare, Epione, Simusante), speech therapist records and neuroimaging data obtained from standard clinical practice. Direct electrostimulation mapping was performed with a Nimbus stimulator (Newmedic, France).

Data analysis

Matlab (Release 2018a, The MathWorksInc, Natick, USA), SPM12, MRICron software (NITRC, University of Massachusetts, MA, USA) Brain VISA/Abatomist Software (5.0, CEA I2BM, CATI nNeuroimaging, INSERM IFR49 and CNRS, France), Marsbar Toolbox (implemented in Matlab), DSI Studio software (<http://dsi-studio.labsolver.org>), RStudio 1.3 (R Foundation, for Statistical Computing, Vienna, Austria), Chorddiag 0.1.3 package implemented in RStudio.

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](#)

Data necessary to the reproducibility of the results are provided in the Supplementary files. Other data can be obtained upon request by contacting Dr Sam Ng (s-ng@chu-montpellier.fr)

Field-specific reporting

Please select the one below that is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.

Life sciences Behavioural & social sciences Ecological, evolutionary & environmental sciences

For a reference copy of the document with all sections, see [nature.com/documents/nr-reporting-summary-flat.pdf](https://www.nature.com/documents/nr-reporting-summary-flat.pdf)

Life sciences study design

All studies must disclose on these points even when the disclosure is negative.

Sample size	<input type="text" value="No sample size calculation was performed"/>
Data exclusions	<input type="text" value="N/A"/>
Replication	<input type="text" value="All electrostimulation applications inducing an intraoperative disturbance were replicated at least 3 times. Stimulations inducing reading/language disorders at least 3 times on the same subcortical/cortical site were selected for analysis."/>
Randomization	<input type="text" value="N/A"/>
Blinding	<input type="text" value="All intraoperative neurocognitive/ reading disorders were collected by a senior speech therapist who remained blinded to electrostimulation application."/>

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	<input type="checkbox"/> Involved in the study	n/a	<input type="checkbox"/> Involved in the study
<input checked="" type="checkbox"/>	<input type="checkbox"/> Antibodies	<input checked="" type="checkbox"/>	<input type="checkbox"/> ChIP-seq
<input checked="" type="checkbox"/>	<input type="checkbox"/> Eukaryotic cell lines	<input checked="" type="checkbox"/>	<input type="checkbox"/> Flow cytometry
<input checked="" type="checkbox"/>	<input type="checkbox"/> Palaeontology and archaeology	<input type="checkbox"/>	<input checked="" type="checkbox"/> MRI-based neuroimaging
<input checked="" type="checkbox"/>	<input type="checkbox"/> Animals and other organisms		
<input checked="" type="checkbox"/>	<input type="checkbox"/> Human research participants		
<input type="checkbox"/>	<input checked="" type="checkbox"/> Clinical data		
<input checked="" type="checkbox"/>	<input type="checkbox"/> Dual use research of concern		

Clinical data

Policy information about [clinical studies](#)

All manuscripts should comply with the ICMJE [guidelines for publication of clinical research](#) and a completed [CONSORT checklist](#) must be included with all submissions.

Clinical trial registration	<input type="text" value="Approval for this study was granted by the Institutional Review Board of Montpellier University Center (N°202000557)"/>
Study protocol	<input type="text" value="The study protocol is provided in the manuscript and was previously published in cited papers."/>
Data collection	<input type="text" value="All data were obtained from a clinical context without additional intervention."/>
Outcomes	<input type="text" value="Electrostimulation mapping findings, based on an asleep-awake-asleep protocol during brain tumor resection, i.e. intraoperative language disorders (either verbal and non-verbal disorders), reading disorders, visuospatial disorders, mentalizing disorders or self-evaluation disorders."/>

Magnetic resonance imaging

Experimental design

Design type	<input type="text" value="Standard structural neuroimaging (FLAIR, T1, T sequences)"/>
Design specifications	<input type="text" value="NA"/>

Behavioral performance measures

Neuroimaging data were confronted to intraoperative behavioral findings.

Acquisition

Imaging type(s)

Structural

Field strength

3T

Sequence & imaging parameters

T1, T1 with Gadolinium, T2, FLAIR either preoperatively and 3 months after surgery

Area of acquisition

Whole brain

Diffusion MRI

 Used Not used

Parameters Diffusion data (population-averaged streamlines) from the Human Connectome Project were used.

Preprocessing

Preprocessing software

Diffusion data from the HCP were used (pre-processing steps are fully detailed in cited publications)

Normalization

Structural data were normalized using SPM12, implemented in the MATLAB environment.

Normalization template

N/A

Noise and artifact removal

N/A

Volume censoring

N/A

Statistical modeling & inference

Model type and settings

N/A

Effect(s) tested

N/A

Specify type of analysis:

Whole brain

ROI-based

Both

Anatomical location(s)

Stimulation points were used as ROIs to generate streamline disconnection and disconnection matrices in DSI studio (HCP datasets).

Statistic type for inference
(See [Eklund et al. 2016](#))

N/A

Correction

FDR

Models & analysis

n/a

Involved in the study

 Functional and/or effective connectivity Graph analysis Multivariate modeling or predictive analysis