

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 <i>Only common tests should be described solely by name; describe more complex techniques in the Methods section.</i>
☒	☐ 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 <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 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	Custom software was developed in C and Python 3.9.11 for running experimental tasks, recording neural data, and decoding speech, cursor, and gestures from neural data. Software packages included PyTorch 2.9, Redis 7.4, StyleTTS2, Numpy 1.22.3, Numba 0.56.4, Scipy 1.9.1, Sounddevice 0.4.7.
Data analysis	Data was analyzed using custom Python 3.9.11 code. Code required for reproducing the figures will be shared via GitHub upon publication.

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](#)

Neural data and speech decoding transcripts recorded during the participant's independent use of the BCI system cannot be shared to preserve the participant's

Research involving human participants, their data, or biological material

Policy information about studies with [human participants or human data](#). See also policy information about [sex, gender \(identity/presentation\), and sexual orientation](#) and [race, ethnicity and racism](#).

Reporting on sex and gender	This study included data from one participant, T15, who is a biological male and identifies as a man. This information was self-reported. No sex or gender based analyses were performed given there was only a single participant and the study was developing a brain-computer interface application.
Reporting on race, ethnicity, or other socially relevant groupings	This study assessed brain-computer interface performance for a single participant. No variables related to race, ethnicity, or other socially relevant groupings were reported or analyzed.
Population characteristics	This study includes data from one participant (T15) who gave informed consent and was enrolled in the BrainGate2 clinical trial (ClinicalTrials.gov Identifier: NCT00912041, registered on June 3, 2009) but this study did not report clinical trial results. T15 is a left-handed man with ALS, 45-47 years old at the times of data collection. His ALS symptoms began five years prior to enrollment into this study.
Recruitment	Participant T15 was enrolled in the BrainGate2 clinical trial after meeting inclusion criteria based in part on disease characteristics. Inclusion and exclusion criteria are available online (ClinicalTrials.gov).
Ethics oversight	The BrainGate2 Neural Interface System clinical trial was approved under an Investigational Device Exemption (IDE) by the US Food and Drug Administration (IDE #G09003). Permission was also granted by the Institutional Review Boards at the University of California, Davis (protocol #1843264 and #1950310) and Mass General Brigham (#2009P000505). All research was performed in accordance with relevant guidelines and regulations.

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

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	No sample-size calculation was performed. Data were collected in a single participant to characterize the performance of a brain-computer interface.
Data exclusions	This study is based on brain-computer interface performance evaluation data collected over months. All available days of data are reported in the study.
Replication	This study assessed brain-computer interface performance with a single performance. Results were replicated across hundreds of independent days of device usage.
Randomization	Randomization into groups is not relevant for this study as only one participant is included in this study.
Blinding	Blinding is not relevant to this study as only one participant was included to assess the performance of a brain-computer interface.

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	Involved in the study
☒	☐ Antibodies
☒	☐ Eukaryotic cell lines
☒	☐ Palaeontology and archaeology
☒	☐ Animals and other organisms
☒	☐ Clinical data
☒	☐ Dual use research of concern
☒	☐ Plants

n/a	Involved in the study
☒	☐ ChIP-seq
☒	☐ Flow cytometry
☒	☐ MRI-based neuroimaging

Plants

Seed stocks

N/A

Novel plant genotypes

N/A

Authentication

N/A