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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>.

Statistics					
For all statistical analyse	es, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.				
n/a Confirmed					
The exact sam	ple size (n) for each experimental group/condition, given as a discrete number and unit of measurement				
A statement o	n whether measurements were taken from distinct samples or whether the same sample was measured repeatedly				
The statistical Only common te	test(s) used AND whether they are one- or two-sided states 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 descripti AND variation	on of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient) (e.g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals)				
For null hypot Give P values as	hesis testing, the test statistic (e.g. F , t , r) with confidence intervals, effect sizes, degrees of freedom and P value noted exact values whenever suitable.				
For Bayesian a	nalysis, information on the choice of priors and Markov chain Monte Carlo settings				
For hierarchical	al and complex designs, identification of the appropriate level for tests and full reporting of outcomes				
Estimates of e	ffect sizes (e.g. Cohen's d , Pearson's r), indicating how they were calculated				
'	Our web collection on <u>statistics for biologists</u> contains articles on many of the points above.				
Software and c	ode				
Policy information abou	ut <u>availability of computer code</u>				
Data collection	Blackrock Microsystems Central Software Suite v7				
Data analysis	Mathworks MATLAB 2018a, Connexon Systems dockex v0.1, Nurmikko Lab Neural Processing Toolkit v0.1 (custom)				
	or 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/reviewers verstrongly encourage code deposition in a community repository (e.g. GitHub). See the Nature Research guidelines for submitting code & software for further information.				
Data					
- Accession codes, uni - A list of figures that I	It <u>availability of data</u> Include a <u>data availability statement</u> . This statement should provide the following information, where applicable: que identifiers, or web links for publicly available datasets have associated raw data restrictions on data availability				
The data that supports th	e findings of this study are available upon request from the corresponding authors [C.H., J.L., A.N.].				
Field-speci	fic reporting				
Please select the one be	elow 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

Life sciences study design

All studies must disclose on these points even when the disclosure is negative.						
Sample size	Two non-human primates (NHPs) were implanted with microelectrode arrays in the superior temporal gyrus. We collected and analyzed data from a total of 3 arrays.					
Data exclusions	Any sections of data that contained audio noise recorded by the microphone (e.g. experimenters talking, animals moving) were discarded to ensure the data set consisted of only the desired audio (English words and NHP calls).					
Replication	Machine learning experiments were performed on data sets collected from 3 different microelectrode arrays implanted in the superior temporal gyrus. Intelligible audio was successfully reconstructed from all neural data across all arrays.					
Randomization	Audio samples (English words and NHP calls) played through the speaker were randomly ordered.					
Blinding	Blinding was not necessary for this work as there is no separate control and experimental group.					

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	n/a	Involved in the study
\boxtimes	Antibodies	\boxtimes	ChIP-seq
\boxtimes	Eukaryotic cell lines	\boxtimes	Flow cytometry
\boxtimes	Palaeontology	\boxtimes	MRI-based neuroimaging
	Animals and other organisms		
\times	Human research participants		
\times	Clinical data		

Animals and other organisms

Policy information about studies involving animals; ARRIVE guidelines recommended for reporting animal research

Laboratory animals This study included two male adult rhesus macaques. Wild animals This study did not involve wild animals. Field-collected samples This study did not involve samples collected from the field. All research protocols were approved and monitored by Brown University Institutional Animal Care and Use Committee, and all Ethics oversight research was performed in accordance with relevant NIH guidelines and regulations.

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