

# Towards human SuperEEG

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## Abstract

Human *SuperEEG*<sup>1</sup> entails measuring ongoing neural activity with perfect precision and at arbitrarily high spatiotemporal resolution. Although true SuperEEG is impossible using existing methods, here we present a model-based method for *inferring* neural activity at millimeter-scale spatial resolutions and millisecond-scale temporal resolutions using standard human intracranial recordings. Our approach assumes that different people’s brains exhibit similar spatial correlations, and that (all else being equal) neural activity at nearby locations will tend to be similar. One can then ask, for an arbitrary individual’s brain: given recordings from a limited set of locations in that individual’s brain, along with the observed spatial correlations in other people’s recordings, what would recordings most likely have looked like at *other* locations in that individual’s brain?

**Keywords:** Electrocorticography (ECoG), intracranial electroencephalography (iEEG), local field potential (LFP), epilepsy, maximum likelihood estimation, Gaussian process regression

## Introduction

Modern human brain recording techniques are fraught with compromise [33]. Commonly used approaches include functional magnetic resonance imaging (fMRI), scalp electroencephalography (EEG), and magnetoencephalography (MEG). For each of these techniques, neuroscientists and electrophysiologists must choose to optimize spatial resolution at the cost of temporal resolution (e.g., as in fMRI) or temporal resolution at the cost of spatial resolution (e.g., as in EEG and MEG). A less widely used approach (due to requiring work with neurosurgical patients) is to record from electrodes implanted directly onto the cortical surface (electrocorticography; ECoG) or into deep brain structures (intracranial EEG; iEEG). However, these intracranial approaches also require compromise: the high temporal and

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<sup>1</sup>The term “SuperEEG” was coined by Robert J. Sawyer in his popular science fiction novel *The Terminal Experiment* [29]

25 spatial resolutions of intracranial recordings comes at the cost of substantially reduced brain coverage,  
26 since safety considerations limit the number of electrodes one may implant in a given patient's brain.  
27 Further, the locations of implanted electrodes are determined by clinical, rather than research, needs.

28 An increasingly popular approach is to improve the effective spatial resolution of MEG or scalp EEG  
29 data by using a geometric approach called *beamforming* to solve the biomagnetic or bioelectrical inverse  
30 problem [28]. This approach entails using detailed brain conductance models (often informed by high  
31 spatial resolution anatomical MRI images) along with the known sensor placements (localized precisely  
32 in 3D space) to reconstruct brain signals originating from theoretical point sources deep in the brain  
33 (and far from the sensors). Traditional beamforming approaches must overcome two obstacles. First,  
34 the inverse problem beamforming seeks to solve has infinitely many solutions. Researchers have made  
35 traction towards constraining the solution space by assuming that signal-generating sources are localized  
36 on a regularly spaced grid spanning the brain and that individual sources are small relative to their  
37 distances to the sensors [1, 11, 34]. The second, and in some ways much more serious, obstacle is that  
38 the magnetic fields produced by external (noise) sources are substantially stronger than those produced  
39 by the neuronal changes being sought (i.e., at deep structures, as measured by sensors at the scalp). This  
40 means that obtaining adequate signal quality often requires averaging the measured responses over tens  
41 to hundreds of responses or trials (e.g., see review by [11]).

42 Another approach to obtaining high spatial and temporal resolution neural data has been to collect  
43 fMRI and EEG data simultaneously. Simultaneous fMRI-EEG has the potential to balance the high spa-  
44 tial resolution of fMRI with the high temporal resolution of scalp EEG, thereby, in theory, providing the  
45 best of both worlds. In practice, however, the signal quality of both recordings suffers substantially when  
46 the two techniques are applied simultaneously (e.g., see review by [13]). In addition, the experimental  
47 designs that are ideally suited to each technique individually are somewhat at odds. For example, fMRI  
48 experiments typically lock stimulus presentation events to the regularly spaced image acquisition time  
49 (TR), which maximizes the number of post-stimulus samples. By contrast, EEG experiments typically

50 employ jittered stimulus presentation times to maximize the experimentalist’s ability to distinguish elec-  
51 trical brain activity from external noise sources such as from 60 Hz alternating current power sources.

52 The current “gold standard” for precisely localizing signals and sampling at high temporal resolution  
53 is to take (ECOG or iEEG) recordings from implanted electrodes (but from a limited set of locations in  
54 any given brain). This begs the following question: what can we infer about the activity exhibited by  
55 the rest of a person’s brain, given what we learn from the limited intracranial recordings we have from  
56 their brain and additional recordings taken from *other* people’s brains? Here we develop an approach,  
57 which we call *SuperEEG*, based on Gaussian process regression [27]. SuperEEG entails using data from  
58 multiple people to estimate activity patterns at arbitrary locations in each person’s brain (i.e., independent  
59 of their electrode placements). We test SuperEEG approach using two large datasets of intracranial  
60 recordings [7, 8, 12, 16–19, 21, 23, 30–32, 35, 40]. We show that the SuperEEG algorithm recovers  
61 signals well from electrodes that were held out of the training dataset. We also examine the factors that  
62 influence how accurately activity may be estimated (recovered), which may have important implications  
63 for electrode design and placement in neurosurgical applications.

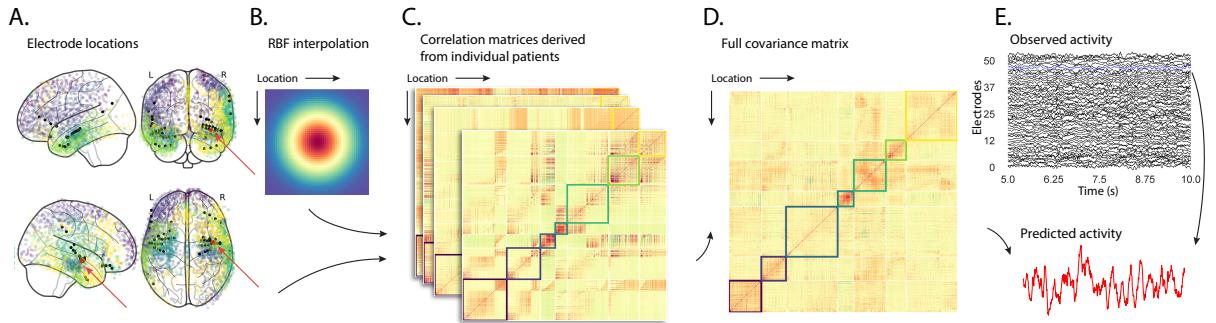
## 64 Approach

65 The SuperEEG approach to inferring high temporal resolution full-brain activity patterns is outlined and  
66 summarized in Figure 1. We describe (in this section) and evaluate (in *Results*) our approach using a two  
67 large previously collected dataset comprising multi-session intracranial recordings. Dataset 1 comprises  
68 multi-session recordings taken from 6876 electrodes implanted in the brains of 88 epilepsy patients [21,  
69 23, 30–32]. Each recording session lasted from XXX–XXX hours, and includes data recorded roughly  
70 from when the patients woke up each morning, to before they went to sleep at the end of each day. In  
71 addition to typical bed-ridden hospital patient activities (e.g., lying in bed, reading, watching television,  
72 using personal electronic devices, listening to music, visiting with family and friends, etc.), the patients  
73 also performed a variety of experimental cognitive tasks throughout their day (primarily list-learning

memory tasks). For the purposes of the Dataset 1 analyses presented here, we aggregated all data across each recording session, ignoring the particular activities or tasks the patients were performing at any given moment. We used Dataset 1 to develop and debug our main SuperEEG approach, and to examine the extent to which SuperEEG might be able to generate task-general predictions. Dataset 2 comprised multi-session recordings from XXX electrodes implanted in the brains of XXX epilepsy patients [7, 8, 12, 16–19, 35, 40]. Whereas Dataset 1 included recordings taken during a wide variety of behaviors, Dataset 2 included recordings taken as each patient performed each of two memory tasks: a random word list free recall task and a categorized word list free recall task. We used Dataset 2 to further examine the ability of SuperEEG to generalize its predictions within versus across tasks.

We first applied fourth order Butterworth notch filter to remove 60 Hz ( $\pm .5$  Hz) line noise from every recording (from every electrode). Next, we downsampled the recordings (regardless of the original samplerate) to 250 Hz. (This downsampling step served to both normalize for differences in sampling rates across patients and to ease the computational burden of our subsequent analyses.) We then excluded any electrodes that showed putative epileptiform activity. Specifically, we excluded from further analysis any electrode that exhibited an average kurtosis of 10 or greater across all of that patient’s recording sessions. We also excluded any patients with fewer than 2 electrodes that passed this criteria, as the SuperEEG algorithm requires measuring correlations between 2 or more electrodes from each patient. For Dataset 1, this yielded clean recordings from 4168 electrodes implanted throughout the brains of 67 patients (Fig. 1A); for Dataset 2, this yielded clean recordings from 2975 electrodes from 24 patients. Each individual patient contributes electrodes from a limited set of brain locations, which we localized in a common space [MNI152; 10]; an example Dataset 1 patient’s 54 electrodes that passed the predefined kurtosis test are highlighted in black and red.

The recording from a given electrode is maximally informative about the activity of the neural tissue immediately surrounding its recording surface. However, brain regions that are distant from the recording surface of the electrode also contribute to the recording, albeit (*ceteris paribus*) to a much lesser extent.



**Figure 1: Methods overview.** **A. Electrode locations.** Each dot reflects the location of a single electrode implanted in the brain of a Dataset 1 patient. A held-out recording location from one patient is indicated in red, and the patient’s remaining electrodes are indicated in black. The electrodes from the remaining patients are colored by  $k$ -means cluster (computed using the full-brain correlation model shown in Panel D). **B. Radial basis function kernel.** Each electrode contributed by the patient (black) weights on the full set of locations under consideration (all dots in Panel A, defined as  $\bar{R}$  in the text). The weights fall off with positional distance (in MNI space) according to an RBF. **C. Per-patient correlation matrices.** After computing the pairwise correlations between the recordings from each patient’s electrodes, we use RBF-weighted averages to estimate correlations between all locations in  $\bar{R}$ . We obtain an estimated full-brain correlation matrix using each patient’s data. **D. Merged correlation model.** We combine the per-patient correlation matrices (Panel C) to obtain a single full-brain correlation model that captures information contributed by every patient. Here we have sorted the rows and columns to reflect  $k$ -means clustering labels [using  $k=7$ ; 41], whereby we grouped locations based on their correlations with the rest of the brain (i.e., rows of the matrix displayed in the panel). The boundaries denote the cluster groups. The rows and columns of Panel C have been sorted using the Panel D-derived cluster labels. **E. Reconstructing activity throughout the brain.** Given the observed recordings from the given patient (shown in black; held-out recording is shown in blue), along with a full-brain correlation model (Panel D), we use Equation 12 to reconstruct the most probable activity at the held-out location (red).

One mechanism underlying these contributions is volume conduction. The precise rate of falloff due to volume conduction (i.e., how much a small volume of brain tissue at location  $x$  contributes to the recording from an electrode at location  $\eta$ ) depends on the size of the recording surface, the electrode's impedance, and the conductance profile of the volume of brain between  $x$  and  $\eta$ . As an approximation of this intuition, we place a Gaussian radial basis function (RBF) at the location  $\eta$  of each electrode's recording surface (Fig. 1B). We use the values of the RBF at any brain location  $x$  as a rough estimate of how much structures around  $x$  contributed to the recording from location  $\eta$ :

$$\text{rbf}(x|\eta, \lambda) = \exp \left\{ -\frac{\|x - \eta\|^2}{\lambda} \right\}, \quad (1)$$

where the width variable  $\lambda$  is a parameter of the algorithm (which may in principle be set according to location-specific tissue conductance profiles) that governs the level of spatial smoothing. In choosing  $\lambda$  for the analyses presented here, we sought to maximize spatial resolution (which implies a small value of  $\lambda$ ) while also maximizing the algorithm's ability to generalize to any location throughout the brain, including those without dense electrode coverage (which implies a large value of  $\lambda$ ). Using our prior work as a guide [22, 24], we set  $\lambda = 20$ , although this could in theory be optimized, e.g., using cross validation or a formal model [e.g., 24].

A second mechanism whereby a given region  $x$  can contribute to the recording at  $\eta$  is through anatomical connections between structures near  $x$  and  $\eta$ . We use spatial correlations in the data to estimate these anatomical connections [2]. Let  $\bar{R}$  be the set of locations at which we wish to estimate local field potentials, and let  $R_s$  be set of locations at which we observe local field potentials from patient  $s$  (excluding the electrodes that did not pass the kurtosis test described above). In the analyses below we define  $\bar{R} = \cup_{s=1}^S R_s$ . We can calculate the expected inter-electrode correlation matrix for patient  $s$ , where  $C_{s,k}(i, j)$  is the correlation between the time series of voltages for electrodes  $i$  and  $j$  from subject  $s$  during session  $k$ , using:

$$\bar{C}_s = r\left(\frac{1}{n}\left(\sum_{k=1}^n z(C_{s,k})\right)\right), \text{ where} \quad (2)$$

$$z(r) = \frac{\log(1+r) - \log(1-r)}{2} \text{ is the Fisher } z\text{-transformation and} \quad (3)$$

$$z^{-1}(z) = r(z) = \frac{\exp(2z) - 1}{\exp(2z) + 1} \text{ is its inverse.} \quad (4)$$

111 Next, we use Equation 1 to construct a number of to-be-estimated locations by number of patient elec-  
 112 trode locations weight matrix,  $W_s$ . Specifically,  $W_s$  approximates how informative the recordings at  
 113 each location in  $R_s$  are in reconstructing activity at each location in  $\bar{R}$ , where the contributions fall off  
 114 with an RBF according to the distances between the corresponding locations:

$$W_s(i,j) = \text{rbf}(i|j, \lambda). \quad (5)$$

115 Given this weight matrix,  $W_s$ , and the observed inter-electrode correlation matrix for patient  $s$ ,  $\bar{C}_s$ ,  
 116 we can estimate the correlation matrix for all locations in  $\bar{R}$  ( $\hat{C}_s$ ; Fig. 1C) using:

$$\hat{N}_s(x, y) = \sum_{i=1}^{|R_s|} \sum_{j=1}^{i-1} W(x, i) \cdot W(y, j) \cdot z(\bar{C}_s(i, j)) \quad (6)$$

$$\hat{D}_s(x, y) = \sum_{i=1}^{|R_s|} \sum_{j=1}^{i-1} W(x, i) \cdot W(y, j). \quad (7)$$

$$\hat{C}_s = r\left(\frac{\hat{N}_s}{\hat{D}_s}\right). \quad (8)$$

After estimating the numerator ( $\hat{N}_s$ ) and denominator ( $\hat{D}_s$ ) placeholders for each  $\hat{C}_s$ , we aggregate these estimates across patients to obtain a single expected full-brain correlation matrix ( $\hat{K}$ ; Fig. 1D):

$$\hat{K} = r\left(\frac{\sum \hat{N}_s}{\sum \hat{D}_s}\right). \quad (9)$$

117 Intuitively, the numerators capture the general structures of the patient-specific estimates of full-brain  
 118 correlations, and the denominators account for which locations were near the implanted electrodes in

119 each patient. To obtain  $\hat{K}$ , we compute a weighted average across the estimated patient-specific full-  
 120 brain correlation matrices, where patients with observed electrodes near a particular set of locations in  
 121  $\hat{K}$  contribute more to the estimate.

122 Having used the multi-patient data to estimate a full-brain correlation matrix at the set of locations  
 123 in  $\bar{R}$  that we wish to know about, we next use  $\hat{K}$  to estimate activity patterns everywhere in  $\bar{R}$ , given  
 124 observations at only a subset of locations in  $\bar{R}$  (Fig. 1E).

125 Let  $\alpha_s$  be the set of indices of patient  $s$ 's electrode locations in  $\bar{R}$ , and let  $\beta_s$  be the set of indices  
 126 of all other locations in  $\bar{R}$ . In other words,  $\beta_s$  reflects the locations in  $\bar{R}$  where we did not observe a  
 127 recording for patient  $s$  (these are the recording locations we will want to fill in using SuperEEG). We can  
 128 sub-divide  $\hat{K}$  as follows:

$$\hat{K}_{\beta_s, \alpha_s} = \hat{K}(\beta_s, \alpha_s), \text{ and} \quad (10)$$

$$\hat{K}_{\alpha_s, \alpha_s} = \hat{K}(\alpha_s, \alpha_s). \quad (11)$$

129 Here  $\hat{K}_{\beta_s, \alpha_s}$  represents the correlations between the “unknown” activity at the locations in  $\beta_s$  and the  
 130 observed activity at the locations in  $\alpha_s$ , and  $\hat{K}_{\alpha_s, \alpha_s}$  represents the correlations between the observed  
 131 recordings (at the locations in  $\alpha_s$ ).

132 Let  $Y_{s,k,\alpha_s}$  be the number-of-timepoints ( $T$ ) by  $\text{length}(\alpha_s)$  matrix of (observed) voltages from the  
 133 electrodes in  $\alpha_s$  during session  $k$  from patient  $s$ . Then we can estimate the voltage from patient  $s$ 's  $k^{th}$   
 134 session at the locations in  $\beta_s$  using [27]:

$$\hat{Y}_{s,k,\beta_s} = ((\hat{K}_{\beta_s, \alpha_s} \cdot \hat{K}_{\alpha_s, \alpha_s}^{-1}) \cdot Y_{s,k,\alpha_s}^T)^T. \quad (12)$$

135 This equation is the foundation of the SuperEEG algorithm. Whereas we observe recordings only at the  
 136 locations in  $\alpha_s$ , Equation 12 allows us to estimate the recordings at all locations in  $\beta_s$ , which we can

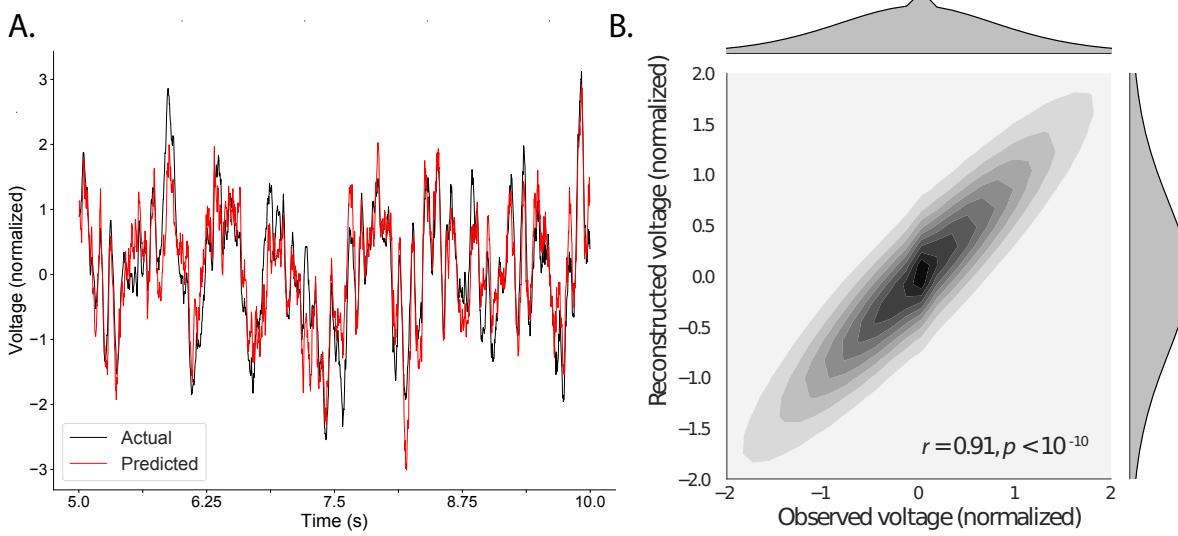
137 define *a priori* to include any locations we wish, throughout the brain. This yields estimates of the time-  
138 varying voltages at *every* location in  $\bar{R}$ , provided that we define  $\bar{R}$  in advance to include the union of all  
139 of the locations in  $\alpha_s$  and all of the locations at which we wish to estimate recordings (i.e., a timeseries  
140 of voltages).

141 We designed our approach to be agnostic to electrode impedances, as electrodes that do not exist  
142 do not have impedances. Therefore our algorithm recovers voltages in standard deviation ( $z$ -scored)  
143 units rather than attempting to recover absolute voltages. (This property reflects the fact that  $\hat{K}_{\beta_s, \alpha_s}$  and  
144  $\hat{K}_{\alpha_s, \alpha_s}$  are correlation matrices rather than covariance matrices.) Also, note that Equation 12 directly  
145 requires computing a  $T$  by  $T$  matrix, which can become computationally intractable when  $T$  is very  
146 large (e.g., for the patient highlighted in Fig. 2,  $T = 20458799$ ). However, because Equation 12 is time  
147 invariant, we may compute  $Y_{s,k,\beta_s}$  in a piecewise manner by filling in  $Y_{s,k,\beta_s}$  one row at a time (using  
148 the corresponding samples from  $Y_{s,k,\alpha_s}$ ).

149 The SuperEEG algorithm described above and in Figure 1 allows us to estimate, up to a constant  
150 scaling factor, local field potentials (LFPs) for each patient at all arbitrarily chosen locations in the set  
151  $\bar{R}$ , even if we did not record that patient's brain at all of those locations. We next turn to an evaluation  
152 of the accuracy of those estimates.

## 153 Results

154 We used a cross-validation approach to test the accuracy with which the SuperEEG algorithm recon-  
155 structs activity throughout the brain. For each patient in turn, we estimated full-brain correlation matrices  
156 (Eqn. 9) using data from all of the *other* patients. This step ensured that the data we were reconstruct-  
157 ing could not also be used to estimate the between-location correlations that drove the reconstructions  
158 via Equation 12 (otherwise the analysis would be circular). For that held-out patient, for each of their  
159 electrodes in turn, we used Equation 12 to reconstruct activity at the held-out electrode location, using  
160 the correlation matrix trained on all other patients' data at  $\hat{K}$ , and using activity recorded from the other



**Figure 2: Observed and reconstructed LFP from a single electrode. A. Example LFP.** A 5 s recording from the red electrode in Figure 1A is displayed in red, and the reconstructed LFP during the same time window is shown in blue. **B. Observed versus reconstructed LFP over 14.2 hours.** The two-dimensional histogram reflects the relation between distributions of observed versus reconstructed voltages from one patient, across the 14.2 hours of recorded data collected over 6 recording sessions. The correlation reported in the panel is between the observed and reconstructed voltages. Both panels: all voltages are represented in standard deviation units (computed within session).

161 electrodes from the held-out patient as  $Y_{s,k,\alpha_s}$ . We then asked: how closely did each of the SuperEEG-  
 162 estimated recordings at those electrodes match the observed recordings from those electrodes (i.e., how  
 163 closely did the estimated  $\hat{Y}_{s,k,\beta_s}$  match the observed  $Y_{s,k,\beta_s}$ ?).

164 To illustrate our approach, we first examine and individual held-out raw LFP trace and its associ-  
 165 ated SuperEEG-derived reconstruction. Figure 2A displays the observed LFP from the red electrode in  
 166 Figure 1A (red), and its associated reconstruction (blue), during a 5 s time window during one of the  
 167 example patient’s six recording sessions. The two traces match closely ( $r = XXX, p = XXX$ ). Fig-  
 168 ure 2B displays a two-dimensional histogram of the actual versus reconstructed voltages for the entire  
 169 14.2 total hours of recordings from the example electrode (correlation:  $r = XXX, p = XXX$ ). This  
 170 example confirms that the SuperEEG algorithm recovers the recordings from this single electrode well.  
 171 Next, we quantify the algorithm’s performance across the full dataset.

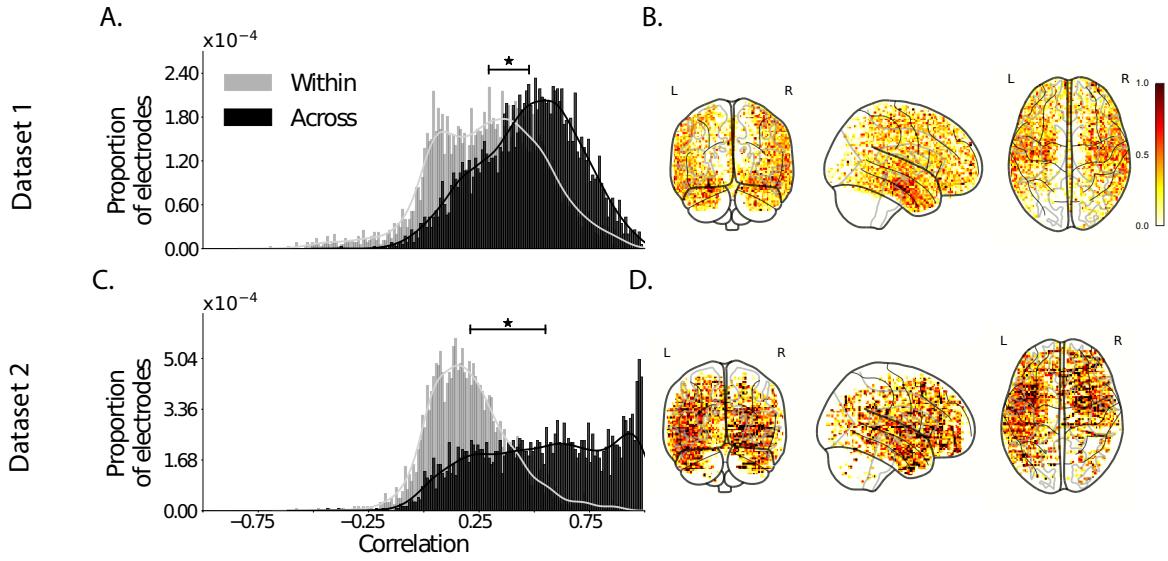
172 For each held-out electrode, from each held-out patient in turn, we computed the average corre-  
173 lation (across recording sessions) between the SuperEEG-reconstructed voltage traces and the observed  
174 voltage traces from that electrode. For this analysis we set  $\bar{R}$  to be the union of all electrode locations  
175 across all patients. This yielded a single correlation coefficient for each electrode location in  $\bar{R}$ , reflecting  
176 how well the SuperEEG algorithm was able to recover the recording at that location by incorporating data  
177 across patients (black histogram in Fig. 3A, map in Fig. 3C). The observed distribution of correlations  
178 was centered well above zero (mean: XXX;  $t$ -test comparing mean of distribution of  $z$ -transformed per-  
179 electrode correlation coefficients to 0:  $t(XXX) = XXX, p = XXX$ ), indicating that the SuperEEG  
180 algorithm recovers held-out activity patterns substantially better than random guessing.

181 As a stricter benchmark, we compared the quality of these across-participant reconstructions (i.e.,  
182 computed using a correlation model derived from other patients' data) to reconstructions generated using  
183 a correlation model trained using the in-patient's data. In other words, for this within-patient benchmark  
184 analysis we estimated  $\hat{C}_s$  (Eqn. 8) for each patient in turn, using recordings from all of that patient's  
185 electrodes except at the location we were reconstructing. These within-patient reconstructions serve as  
186 an estimate of how well data from all of the other electrodes from a single patient may be used to esti-  
187 mate held-out data. This allows us to ask how much information about the activity at a given electrode  
188 might be inferred through (a) volume conductance or other sources of "leakage" from activity patterns  
189 measured from the patient's other electrodes and (b) across-electrode correlations learned from that sin-  
190 gle patient. As shown in Figure 3A (gray histogram), the distribution of within-patient correlations was  
191 centered well above zero (mean: XXX;  $t$ -test comparing mean of distribution of  $z$ -transformed per-  
192 electrode correlation coefficients to 0:  $t(XXX) = XXX, p = XXX$ ). However, the across-patient  
193 correlations were substantially higher ( $t$ -test comparing average  $z$ -transformed within versus across pa-  
194 tient electrode correlations:  $t(XXX) = XXX, p = XXX$ ). This is an especially conservative test,  
195 given that the across-patient SuperEEG reconstructions exclude (from the correlation matrix estimates)  
196 all data from the patient whose data is being reconstructed. We repeated each of these analyses on a

197 second independent dataset and found similar results (Fig. 3B, D; within versus across reconstruction  
198 accuracy:  $t(23) = 6.93, p < 10^{-5}$ ). We also replicated this result separately for each of the two ex-  
199 periments from Dataset 2 (Fig. S2). This overall finding, that when reconstructing held-out data from a  
200 given patient correlation models derived from *other* patient's data yield higher reconstruction accuracy  
201 than correlation models derived from that patient, has two important implications. First, it implies that  
202 distant electrodes provide additional predictive power to the data reconstructions beyond the information  
203 contained solely in nearby electrodes. (This follows from the fact that each patient's electrodes are im-  
204 planted in a unique set of locations, so for any given electrode the closest electrodes in the full dataset  
205 are likely to come from the same patient.) Second, it implies that the spatial correlations derived from  
206 the SuperEEG algorithm are, to some extent, similar across people.

207 The recordings we analyzed from Dataset 1 comprised data collected as the patients performed a  
208 variety of (essentially uncontrolled) tasks throughout each day's recording session. That we observed  
209 reliable reconstruction accuracy across patients suggests that the spatial correlations derived from the  
210 SuperEEG algorithm are, to some extent, similar across tasks. We tested this finding more explicitly  
211 using Dataset 2. In Dataset 2, the recordings were limited to times when each patient was participating  
212 in each of two experiments (Experiment 1, a random-word list free recall task, and Experiment 2, a  
213 categorized list free recall task). We wondered whether a correlation model trained using data only from  
214 one experiment might yield good predictions for data from the other experiment. Further, we wondered  
215 about the extent to which it might be beneficial or harmful to combine data across tasks.

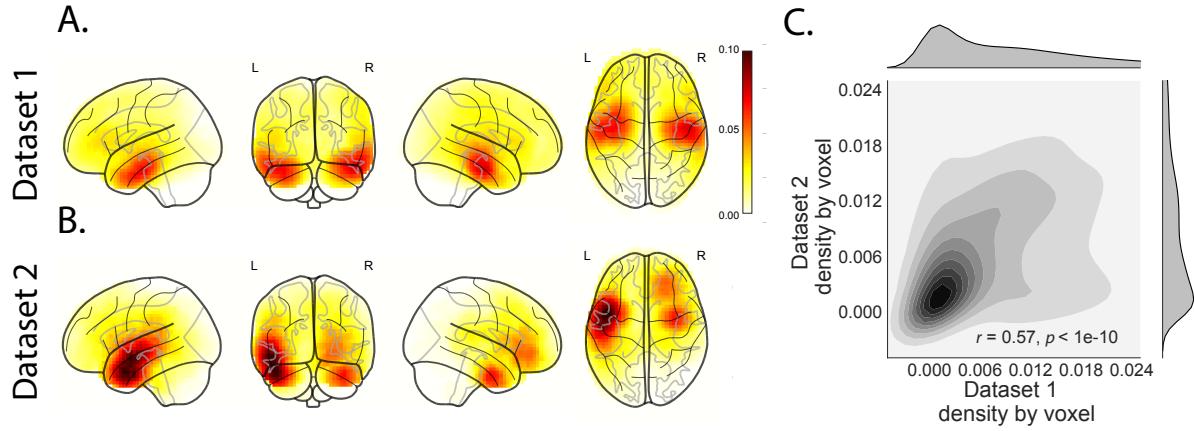
216 To test the task-specificity of the SuperEEG-derived correlation models, we repeated our within-  
217 and across-patient cross validation procedures separately for Experiment 1 and Experiment 2 data from  
218 Dataset 2. We then compared the reconstruction accuracies of held-out electrodes, for models trained  
219 within versus across the two experiments, or combining across both experiments (Fig. S3). In every case  
220 we found that across-patient models trained using data from all other patients out-performed within-  
221 patient models trained on data only from the subject contributing the given electrode ( $ts(XXX) >$



**Figure 3: Reconstruction quality across all electrodes in two ECoG datasets. A. Distributions of correlations between observed versus reconstructed activity by electrode, for Dataset 1.** The across-patient distribution (black) reflects reconstruction performance using a correlation model trained on all but one patient, and then applied to the held-out patient's data. The within-patient distribution (gray) reflects performance using a correlation model trained on the same patient who contributed the to-be-reconstructed electrode. **B. Distributions of correlations for Dataset 2.** This panel is in the same format as Panel A, but reflects results obtained from Dataset 2. **C.-D. Reconstruction performance by location.** Each dot reflects the location of a single implanted electrode from Dataset 1 (Panel C) or Dataset 2 (Panel D). The dot colors denote the average across-session correlation, using the across-patient correlation model, between the observed and reconstructed activity at the given electrode location.

222  $XXX, ps < XXX$ ). All reconstruction accuracies also reliably exceeded chance performance ( $ts(XXX) >$   
223  $XXX, ps < XXX$ ). Average reconstruction accuracy was highest for the across-patient models limited  
224 to data from the same experiment (mean accuracy: 0.55), next-highest for the across-patient models that  
225 combined data across both experiments (mean accuracy: 0.50), and lowest for models trained across task  
226 (mean accuracy: 0.37). This result also held for each of the Dataset 2 experiments individually (Fig. S4).  
227 Taken together, these results indicate that there are reliable commonalities in the spatial correlations of  
228 full-brain activity across tasks, but that there are also reliable differences in these spatial correlations  
229 across tasks. Whereas reconstruction accuracy benefits from incorporating data from other patients, re-  
230 construction accuracy is highest when constrained to within-task data, or data that includes a variety of  
231 tasks (e.g., Dataset 1, or combining across the two Dataset 2 experiments).

232 Although both datasets we examined provide good full-brain coverage (when considering data from  
233 every patient; e.g. Fig. 3B, D), electrodes are not placed uniformly throughout the brain. For example,  
234 electrodes are more likely to be implanted in regions like the medial temporal lobe (MTL), and are rarely  
235 implanted in occipital cortex (Fig. 4A, B). Separately for each dataset, for each voxel in the 4 mm<sup>3</sup> voxel  
236 MNI152 brain, we computed the proportion of electrodes contained within a 20 MNI unit radius sphere  
237 centered on that voxel. We defined the *density* at that location as this proportion. Across Datasets 1  
238 and 2, the electrode placement densities were similar (correlation by voxel:  $r = 0.57, p < 10^{-10}$ ). We  
239 wondered whether regions with good coverage might be associated with better reconstruction accuracy  
240 (e.g. Fig. 3B, D indicate that many electrodes in the MTL have relatively high reconstruction accuracy,  
241 and occipital electrodes tend to have relatively low reconstruction accuracy). To test whether this held  
242 more generally across the entire brain, for each dataset we computed the electrode placement density for  
243 each electrode from each patient (using the proportion of *other* patients' electrodes within 20 MNI units  
244 of the given electrode). We then correlated these density values with the across-patient reconstruction  
245 accuracies for each electrode. Contrary to our expectation, rather than positive correlations, we found  
246 weak (but reliable) negative correlations between reconstruction accuracy and density for both datasets



**Figure 4: Electrode sampling density by location.** **A. Electrode sampling density by voxel in Dataset 1.** Each voxel is colored by the proportion of total electrodes in the dataset that are located within a 20 MNI unit radius sphere centered on the given voxel. **B. Electrode sampling density by voxel in Dataset 2.** This panel displays the sampling density map for Dataset 2, in the same format as Panel A. **C. Correspondence in sampling density by voxel across Datasets 1 and 2.** The two-dimensional histogram displays the by-voxel densities in the two Datasets, and the one-dimensional histograms display the proportions of voxels in each dataset with the given density value. The correlation reported in the panel is across voxels in the  $4 \text{ mm}^3$  MNI brain.

(Dataset 1:  $r = XXX, p = XXX$ ; Dataset 2:  $r = -0.16, p < 10^{-10}$ ). This indicates that the reconstruction accuracies we observed are not driven solely by sampling density, but rather may also reflect higher order properties of neural dynamics such as functional correlations between distant voxels [3].

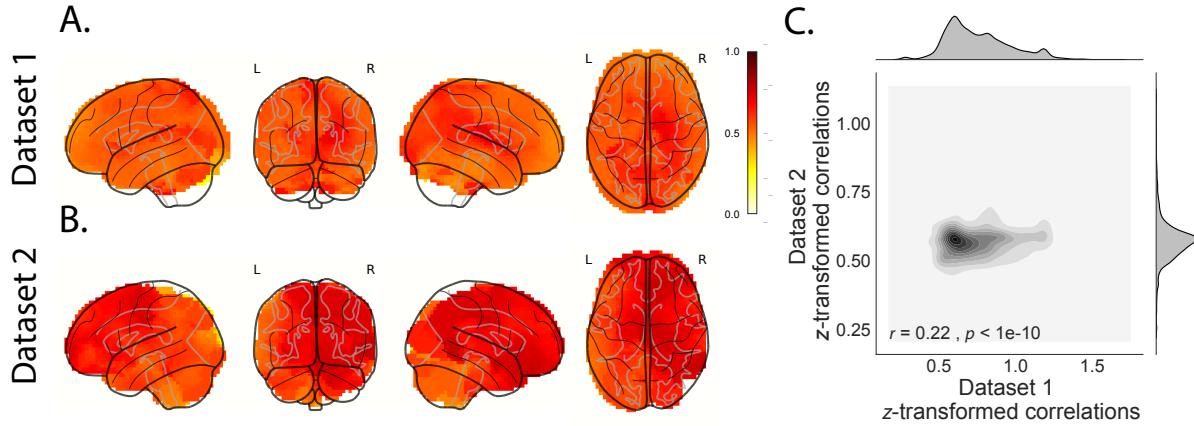
In neurosurgical applications where one wishes to infer full-brain activity patterns, can our framework yield insights into where the electrodes should be placed? A basic assumption of our approach (and of most prior ECoG work) is that electrode recordings are most informative about the neural activity near the recording surface of the electrode. But if we consider that activity patterns throughout the brain are meaningfully correlated, are there particular implantation locations that, if present in a patient's brain, yield especially high reconstruction accuracies throughout the rest of the brain? For example, one might hypothesize that brain structures that are heavily interconnected with many other structures could be more informative about full-brain activity patterns than comparatively isolated structures.

To gain insights into whether particular electrode locations might be especially informative, we first

259 computed the average reconstruction accuracy across all of each patient’s electrodes (using the across-  
260 patients cross validation test; black histograms in Fig. 3A and C). We labeled each patient’s electrodes in  
261 each dataset with the average reconstruction accuracy for that patient. In other words, we assigned every  
262 electrode from each patient the same value, reflecting how well the activity patterns at those electrodes  
263 were reconstructed on average. Next, for each voxel in the 4 mm<sup>3</sup> MNI brain, we computed the average  
264 value across any electrode (from any patient) that came within 20 MNI units of that voxel’s center.  
265 Effectively, we computed an *information score* for each voxel, reflecting the average reconstruction  
266 accuracy across any patients with electrodes near each voxel— where the averages were weighted to  
267 reflect patients who had more electrodes implanted near that location. This yielded a single map for each  
268 dataset, highlighting regions that are especially promising implantation targets in terms of providing full-  
269 brain activity information via SuperEEG (Fig. 5A, B). Despite task and patient differences across the two  
270 datasets, we nonetheless found that the maps of the most promising implantation targets derived from  
271 both datasets were similar (voxelwise correlation between information scores across the two datasets:  
272  $r = XXX, p = XXX$ ). While the correspondence between the two maps was not perfect, our finding  
273 that there were some commonalities between the two maps lends support to the notion that different brain  
274 areas are differently informative about full-brain activity patterns. Further, regions like XXX, XXX, and  
275 XXX that exhibited strong information scores in both datasets may be especially promising implantation  
276 targets.

## 277 Discussion

278 Are our brain’s networks static or dynamic? And to what extent are the network properties of our brains  
279 stable across people and tasks? One body of work suggests that our brain’s *functional* networks are  
280 dynamic [e.g., 24], person-specific [e.g., 9], and task-specific [e.g., 39]. In contrast, although the gross  
281 anatomical structure of our brains changes meaningfully over the course of years as our brains develop,  
282 on the timescales of typical neuroimaging experiments (i.e., hours to days) our anatomical networks



**Figure 5: Most informative electrode locations.** **A. Dataset 1 information score by voxel.** The voxel colors reflect the weighted average reconstruction accuracy across all electrodes from any patients with at least one electrode within 20 MNI units of the given voxel. **B. Dataset 2 information score by voxel.** This panel is in the same format as Panel A. **C. Correspondence in information scores by voxel across Datasets 1 and 2.** Same format as Figure 4C.

are largely stable [e.g., 4]. Further, many aspects of brain anatomy, including white matter structure, is largely preserved across people [e.g., 15, 26, 37] There are several possible means of reconciling this apparent inconsistency between dynamic person- and task-specific functional networks versus stable anatomical networks. For example, relatively small magnitude anatomical differences across people may be reflected in reliable functional connectivity differences. Along these lines, one recent study found that diffusion tensor imaging (DTI) structural data is similar across people, but may be used to predict person-specific resting state functional connectivity data [2]. Similarly, other work indicates that task-specific functional connectivity may be predicted by resting state functional connectivity data [5, 38]. Another (potentially complementary) possibility is that our functional networks are constrained by anatomy, but nevertheless exhibit (potentially rapid) task-dependent changes [e.g., 36].

Here we have taken a model-based approach to studying whether high spatiotemporal resolution activity patterns throughout the human brain may be explained by a static connectome model that is shared across people and tasks. Specifically, we trained a model to take in recordings from a subset of brain locations, and then predicted activity patterns during the same interval, but at *other* locations

297 that were held out from the model. Our model, based on Gaussian process regression, was built on  
298 three general hypotheses about the nature of the correlational structure of neural activity (each of which  
299 we tested). First, we hypothesized that functional correlations are stable over time and across tasks.  
300 We found that, although aspects of the patients' functional correlations that were stable across tasks, we  
301 achieved better reconstruction accuracy when we trained the model on within-task data [we acknowledge  
302 that our general approach could potentially be extended to better model across-task changes, following  
303 5, 38, and others]. Second, we hypothesized that some of the correlational structure of people's brain  
304 activity is similar across individuals. Consistent with this hypothesis, our model explained the data  
305 best when we trained the correlation model using data from *other* patients— even when compared to a  
306 correlation model trained using the same patient's data. Third, we resolved ambiguities in the data by  
307 hypothesizing that neural activity from nearby sources will tend to be similar, all else being equal. This  
308 hypothesis was supported through our finding that all of the models we trained that incorporated this  
309 spatial smoothness assumption predicted held-out data well above chance.

310 One potential limitation of our approach is that it does not provide a natural means of estimating the  
311 precise timing of single-neuron action potentials. Prior work has shown that gamma band and broad-  
312 band activity in the LFP may be used to estimate the firing rates of neurons that underly the population  
313 contributing to the LFP [6, 14, 20, 25]. Because SuperEEG reconstructs LFPs throughout the brain, one  
314 could in principle use gamma or broadband power in the reconstructed signals to estimate the corre-  
315 sponding firing rates (though not the timings of individual action potentials).

316 Beyond providing a means of estimating ongoing activity throughout the brain using already im-  
317 planted electrodes, our work also has implications for where to place the electrodes in the first place.  
318 Electrodes are typically implanted to maximize coverage of suspected epileptogenic tissue. However,  
319 our findings suggest that this approach could be further optimized. Specifically, one could leverage not  
320 only the non-invasive recordings taken during an initial monitoring period (as is currently done rou-  
321 tinely), but also recordings collected from other patients. We could then ask: given everything we know

322 about the other patients and from the scalp recordings of this new patient, where should we place a fixed  
323 number of electrodes to maximize our ability to map seizure foci? As shown in Figure 5, recordings  
324 from different locations are differently informative in terms of reconstructing the spatiotemporal patterns  
325 throughout the brain. This property might be leveraged in decisions about where to surgically implant  
326 electrodes in future patients.

## 327 **Concluding remarks**

328 Over the past several decades, neuroscientists have begun to leverage the strikingly profound mathemati-  
329 cal structure underlying the brain’s complexity to infer how our brains carry out computations to support  
330 our thoughts, actions, and physiological processes. Whereas traditional beamforming techniques rely on  
331 geometric source-localization of signals measured at the scalp, here we propose an alternative approach  
332 that leverages the rich correlational structure of a large dataset of human intracranial recordings. In do-  
333 ing so, we are one step closer to observing, and perhaps someday understanding, the full spatiotemporal  
334 structure of human neural activity.

## 335 **Code availability**

336 We have published an open-source toolbox implementing the SuperEEG algorithm. It may be down-  
337 loaded [here](#).

## 338 **Data availability**

339 The dataset analyzed in this study was generously shared by Michael Kahana. A portion of Dataset 1  
340 may be downloaded [here](#). Dataset 2 may be downloaded [here](#).

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<sup>349</sup> **Author Contributions**

<sup>350</sup> J.R.M conceived and initiated the project. L.L.W.O. and A.C.H. performed the analyses. J.R.M. and  
<sup>351</sup> L.L.W.O. wrote the manuscript.

<sup>352</sup> **Author Information**

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<sup>358</sup> **References and Notes**

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