

EEG and MEG Inversion Using Convolutional and Recurrent Neural Networks

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Abstract

Real-time localization of neuronal activity has a number of uses, including brain-computer interfaces and medical diagnostics. Generally, this is done by taking measurements of the brain's magnetic and electric fields (magnetoencephalography [MEG] and electroencephalography [EEG]), and inverting the measurements. Most approaches are physics-based, and attempt to find the best estimate of the lead-field matrix, which relates the dipole activation to the field strength by minimizing the error of an estimate. To date, most approaches using the lead-field matrix are complicated and too slow in real time, aimed primarily at elucidating brain structural functional relationships from experimental data. We propose a new technique in which the location of peak neuronal current is estimated by treating EEG and MEG as a two-channel image, or time-series of images, which is processed by a neural network which returns the location of the dipole of peak magnitude. Four architectures are tested: 2-layer perceptron, convolutional neural network (CNN), recurrent neural network (RNN), and CNN feeding RNN. In the absence of true measures of neuronal activity, we used two publicly available MEG/EEG datasets, and treated the estimates of the traditional minimum-norm estimate (MNE) as true estimates. We test the four variations of the network architecture, and in the best case (CNN only), we achieve test dataset errors (RMSE of max dipole location) of 1.9 and 5.1 mm on two separate datasets of MEG/EEG time series from subjects with different types of stimulus.

Index Terms

Electrophysical imaging, Brain, Neural networks, Dipole.

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I. INTRODUCTION

There is a great need for interpretation of brain signals for both use in control of devices, for prosthetics, for example, or for disease diagnostics. Knowledge of which neurons are most active in the brain (known as localization) at a given time can help in a number of areas. It can aid in diagnosis and presurgical evaluation of epileptics, for instance [1]. Mapping brain activity with high resolution in time and space also helps with fundamental understanding of the human brain and cognition [2]. Brain-computer interfaces have been developed using this sort of neuron data as well [3]. Two sources of data used in neuron localization are MEG and EEG. Magnetoencephalography (MEG), the measurement of the brain's magnetic field, is typically done using cryogenic superconducting quantum interference devices (SQUID) on a dense grid surrounding the skull. Additionally, electroencephalography (EEG), is also used for BCI as well as medical diagnostics and study of brain structure and function [4]. Particularly for the latter, there is great utility in using the external measurements of MEG and EEG to determine which parts of the brain are most electrically active, a problem known as localization or inversion. Functional magnetic resonance imaging (fMRI) can be used but instead measures blood flow, and is thus not a direct measurement of electrical activity [5].

The fundamental model for describing the relationship between MEG/EEG measurements and neural activity is the dipole model [6]. This approximates each neuron as a current dipole on a fixed grid, usually obtained through MRI. Dipoles are either considered to be free to rotate or fixed, and only the radial vector component of the dipole is considered for its contribution to the measured fields. Localization, or inversion, is the process of inverting the measured magnetic and/or electric fields to obtain the spatial location and magnitudes of the neuronal current. Two approaches, determining the location of a single dipole which would best produce the measured fields (single dipole fit), and determining the current dipole at every point in the grid, have been used. In our approach, we instead opt to find the location of the maximum dipole, rather than all or a single best-fit. See Figs 1.

Classically, MEG/EEG inversion has followed one of a few approaches, with incremental enhancements as sophistication and computational power has increased. [8] provides an excellent review of techniques. [6] established the original framework of the dipole in a sphere model, in which a lead-field matrix based on magnetostatic and electrostatic equations, and the positions of dipoles, transforms a vector of dipole magnitudes into a vector of magnetic flux and electric

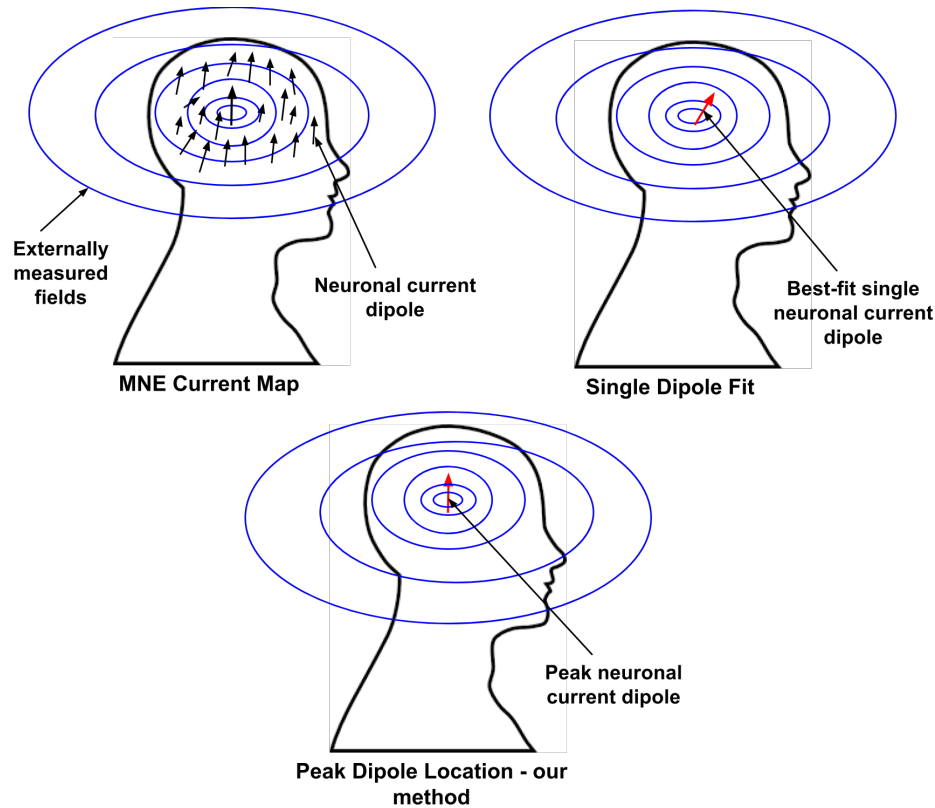


Fig. 1. Simplified drawing of the neuron/fields model used in this paper and the two others commonly used in MEG/EEG inversion.

potential at sensor locations. The problem of localization becomes a problem of finding the locations and magnitudes of dipoles that minimize the square error of predicted versus observed fields. Using this formalism and a finite-element model, [9] localized dipoles from MEG data. [10] used Kalman filtering to solve the dynamical inverse problem. The signal-space separation method transforms the data into basis functions in signal space to better filter background noise sources [11]. Beamformer techniques have also been applied for MEG/EEG inversion [12]. More recently, EEG has been inverted using artificial neural networks [13].

Neural networks have proven an important tool for accurate predictions from noisy and large datasets. A recent technique is convolutional neural networks, in which image inputs are convolved with weighted kernels [14]. To treat time-series or sequences, recurrent neural networks, incorporating temporal feedback, have been developed, such as the long-short-term memory RNN [15]. CNN and RNN have been combined in [16] to translate video sequences into text descriptions of the video.

Our proposed idea is to apply the combined CNN/RNN framework to sequences of MEG/EEG data, and predict the location of only the peak dipole. In this way we simplify the problem, and with appropriately trained networks, we could make quick predicts from data, possibly in real time. To demonstrate our technique, we use publicly available datasets, and use the dipole currents predicted by a well established technique (MNE) as the true value to which we compare our estimates and train our network [17]. Further, we investigate four variations of the network architecture (2-layer MLP, RNN, CNN, and CNN with RNN), to establish the advantages and disadvantages of each.

II. METHODS

In this section, we describe our methodology, including datasets, preprocessing, and neural network architectures.

1) Datasets: To test our method, we used two multimodal datasets. Both were processed using the python-mne package [17]. In each, we use the current map predicted by MNE as the truth; from this we find the location of the maximum dipole as the truth on which to train our networks. This is because there is no real, publicly available dataset which includes both MEG/EEG electrocorticographic (ECoG) measurements (placing electrodes directly in the brain to measure activity, the only way to exactly measure the location of peak neural current). The spatial location error associated with inversion is estimated at around 3mm [18], so errors we report are in addition to that error. This is a limitation of the datasets available, and not the fundamentals of our proposed technique - with a training dataset which has ECoG we could achieve higher spatial accuracy.

One dataset [19] consisted of 19 subjects presented with stimulus of famous, unfamiliar, or scrambled faces. This data was obtained from the OpenfMRI database. Its accession number is ds000117. There were 102 magnetometers, 204 planar gradiometers, and 70 electrodes, with 3 electrodes used for ocular and cardiac artifacts. We excluded gradiometers, as our study is to develop an inversion technique for use with our own MEG system, which only has the radial magnetometers. We considered one subject only (sub007) as that subject's MRI was the only one to segment properly into skin, skull, and cortex layers (an important feature for the inversion step). We used the data processing pipeline provided with the dataset here: <http://mne-tools.github.io/mne-biomag-group-demo/>. The sampling rate was 1100 Hz, with 551 time points per trial, and oct6 (source spacing 4.9 mm) grid for the dipole mesh used in MNE. There

were a total of 879 trials. We studied subjects 7 and 8, as their MRIs were best segmented by the MNE toolbox - accurate segmentation into skull, skin, and cortex is necessary for classical inversion of MEG/EEG.

The second dataset [20] is included with MNE, and consists of trials where multiple subjects received audio or visual stimulus: checkerboard patterns were presented into the left and right visual field, interspersed by tones to the left or right ear. The dataset includes, for each trial, one ocular channel (for artifact removal), 59 EEG channels 102, and MEG channels. The source grid used was oct5 (9.9 mm). The sampling rate was 150 Hz, for 106 time steps, with 239 trials.

A. Preprocessing

For both datasets, channels were preprocessed through lowpass filtering, followed by electrooculogram/electrocardiogram (EOG/ECG) artifact removal using direct measurements of EOG or ECG and/or independent component analysis. Data was epoched (cut into trials) but not averaged over each type of stimulus - each trial is considered as a separate datapoint. Further, prior to input to the neural network, the data was transformed by principle component analysis, to avoid numerical problems due to the small sensor units (fT and μV). To obtain the “true” location of the peak dipole, the MNE method is used on the pre-processed data, a neuronal current map is estimated; and the location of the peak neuronal current is found. This is the true value we use for training our model.

B. Description of Neural Networks

We considered four variations of the network structure, each of which was programmed using Google’s Tensorflow API in Python (tensorflow.com). The simplest configuration was a multi-layer perceptron, where the PCA-transformed data is input as a vector at each timestep, and transformed by an input layer with ReLU activation functions, and then an output layer, with linear activation functions, to give the coordinate of the maximum current neuron at each timestep (Fig. 2). The next network type was a convolutional neural network, where the data at each timestep was interpolated to an 11 by 11 spatial grid, using the sensors’ physical locations, such that the data forms a sequence of 2-channel (EEG and MEG) images. The grid size was chosen such that increasing the number of pixels had no marginal benefit on the model error. There were two convolutional layers, with kernel size 3 by 3. This was followed by a dense layer which is mapped to the output with a linear activation function, to give the coordinate of the

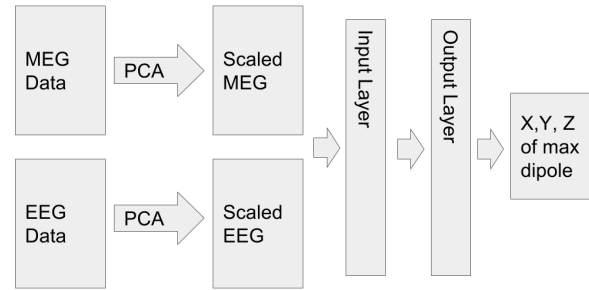


Fig. 2. Block diagram of MLP neural network.

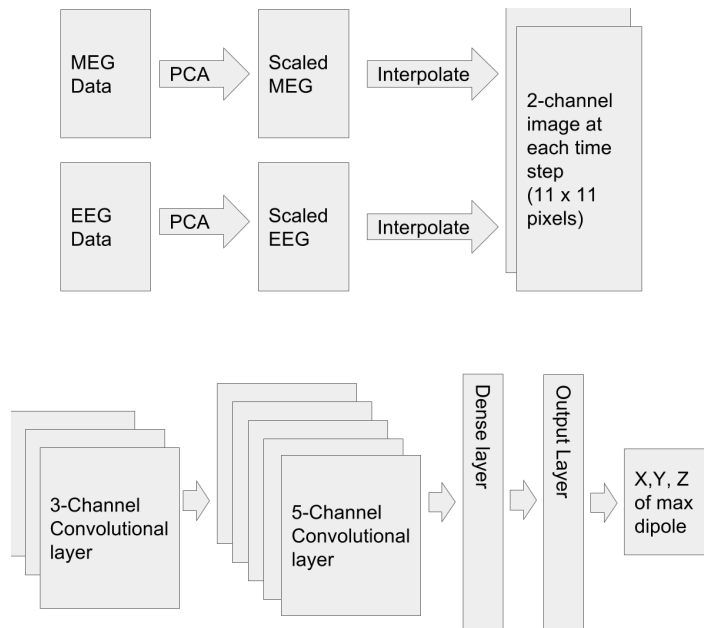


Fig. 3. Block diagram of CNN neural network.

maximum current neuron at each timestep. We also considered a recurrent neural network, with long short term memory (LSTM) cells, with an input layer, recurrent layer, and output layer, returning the dipole location of the maximum current neuron over the entire timeseries, not at

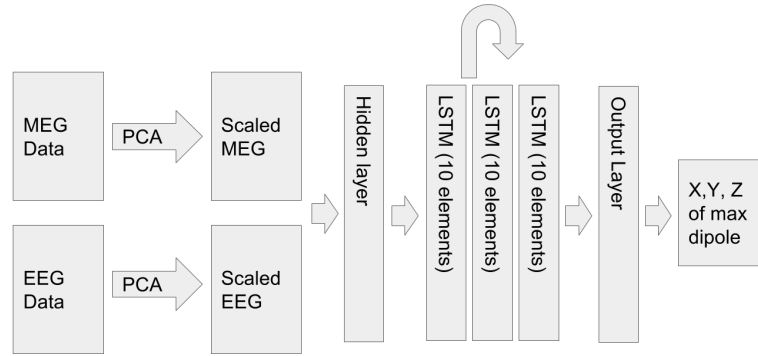


Fig. 4. Block diagram of RNN neural network.

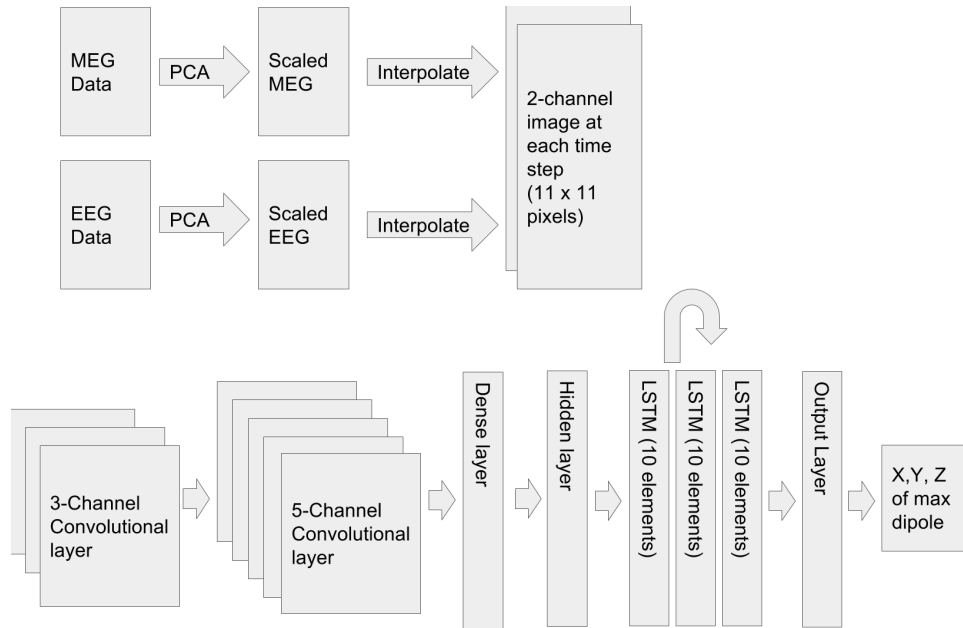


Fig. 5. Block diagram of CNN+RNN neural network.

each timestep. Finally, we considered a combination of the CNN and RNN networks (Fig. 5). Network parameters are given in Table I.

TABLE I
NETWORK PARAMETERS.

Network	CNN	Hidden layer	RNN
MLP	N/A	10 units	N/A
CNN	2 layers (3, then 5 channels)	N/A	N/A
RNN	N/A	N/A	10 units, 3 layers
CNN+RNN	2 layers (3, then 5 channels)	N/A	10 units, 3 layers

C. Training and testing

To train the networks, on each dataset the data was divided into randomly selected test, validation, and training sets, where the test set was 20% of the total dataset, validation was 20% of the remainder, and the remainder was divided into batches for training of 20% each. Cost was the root mean square error (RMSE) of the location of the maximum current dipole, where the true value was taken as the Minimum Norm Estimate found using the MNE Python package. Validation error was logged every 100 steps, and each batch was trained for 1250 steps (giving 5000 train steps total). We used Adam optimization to train the network parameters [21] with a learning rate of 0.005. Five train/validation/test sets were randomly shuffled to ensure the results were reliable. To further prevent overtraining, the training set was divided into batches, the networks were trained on a sequence of batches, and as a consequence the training curves show a series of spikes. Whenever a new batch of training examples is selected, the training error temporarily spikes.

III. RESULTS

TABLE II
TEST SET RMSE FROM 5 CROSS-VALIDATION RUNS (MEAN, STANDARD DEVIATION)

Subject	MLP	CNN	RNN	CNN+RNN
Auditory (1 dipole)	5.606, 0.330	4.981, 0.387	37.647, 3.408	33.879, 4.572
Faces, subject #7 (1 dipole)	1.920, 0.015	1.917, 0.018	47.391, 4.828	42.955, 1.410
Faces, subject #8 (1 dipole)	1.137, 0.062	1.235, 0.030	25.745, 1.103	19.754, 0.814
Auditory (Best of 100 dipoles)	2.206, 0.179	2.176, 0.133	11.994, 2.915	10.200, 1.570
Faces, subject #7 (Best of 100 dipoles)	1.149, 0.030	1.137, 0.023	18.589, 2.200	17.787, 0.747
Faces, subject #8 (Best of 100 dipoles)	0.683, 0.068	0.632, 0.023	8.560, 1.116	9.043, 1.714

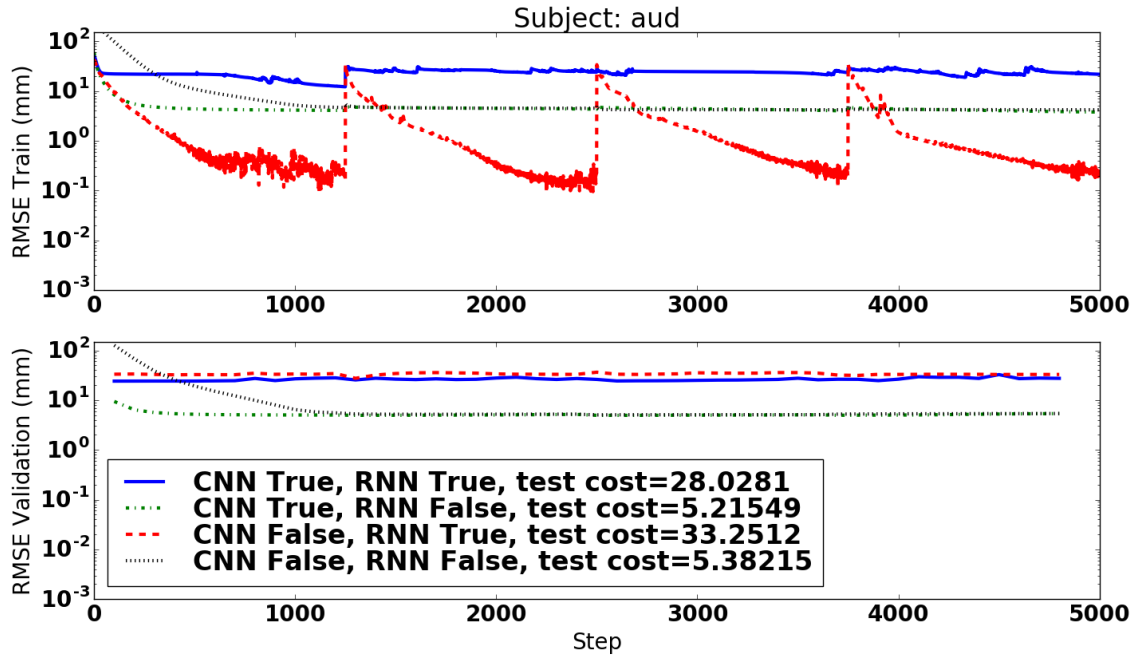


Fig. 6. Training/validation results for auditory stimulus dataset (1 dipole).

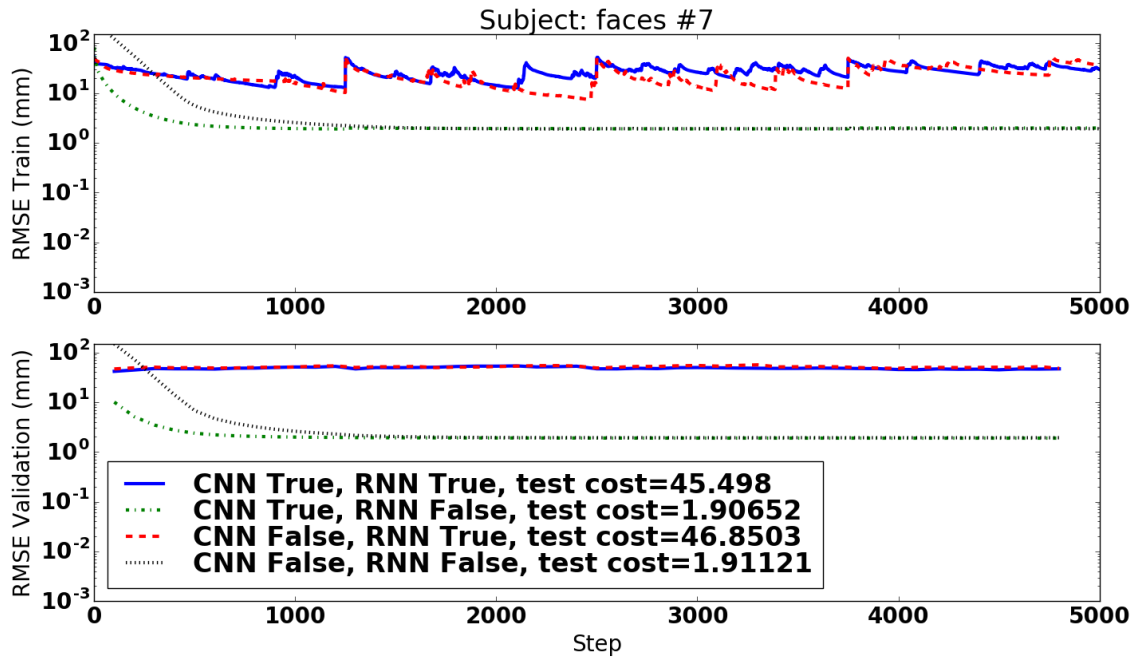


Fig. 7. Training/validation results for faces stimulus dataset on subject #7 (1 dipole).

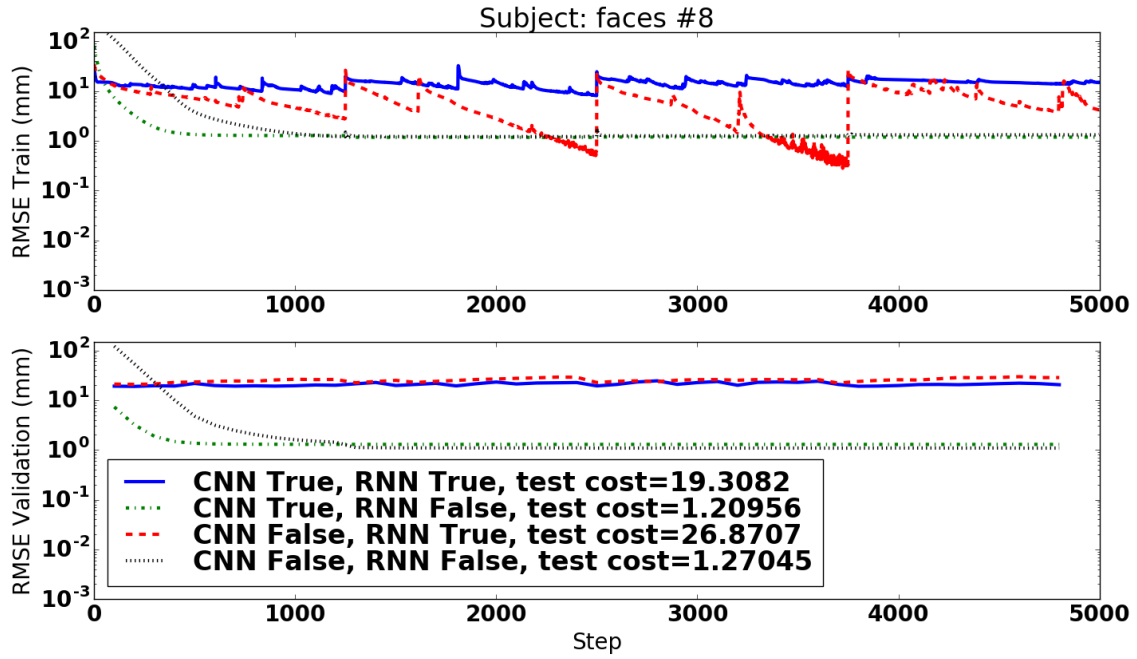


Fig. 8. Training/validation results for faces stimulus dataset on subject #8 (1 dipole).

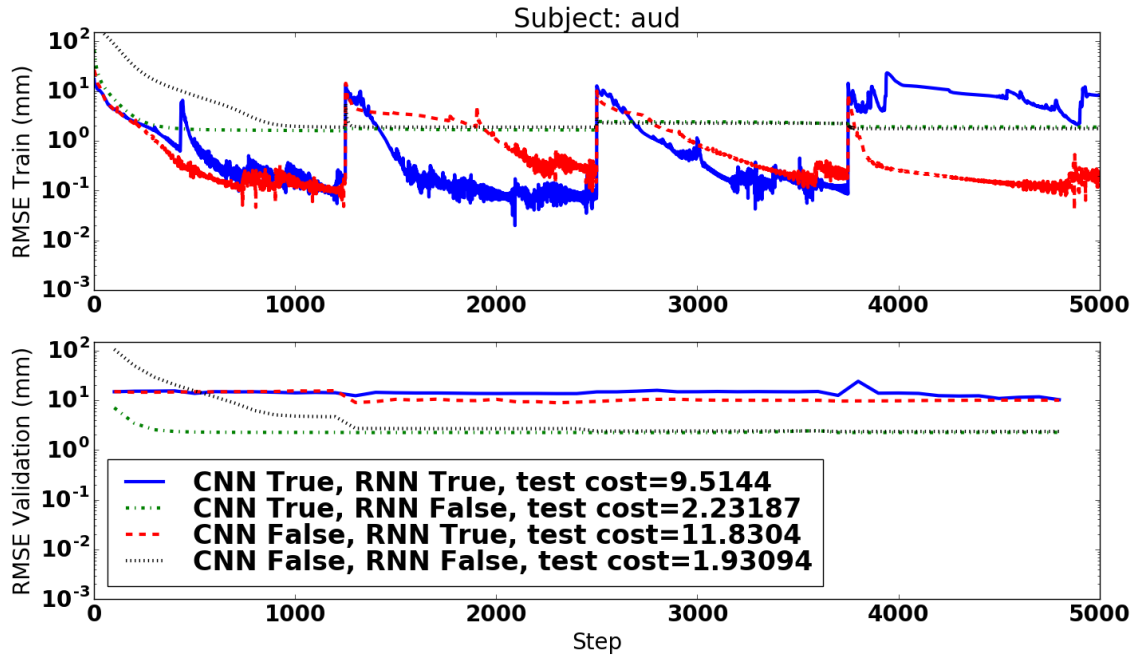


Fig. 9. Training/validation results for auditory stimulus dataset (Best of 100 dipoles).

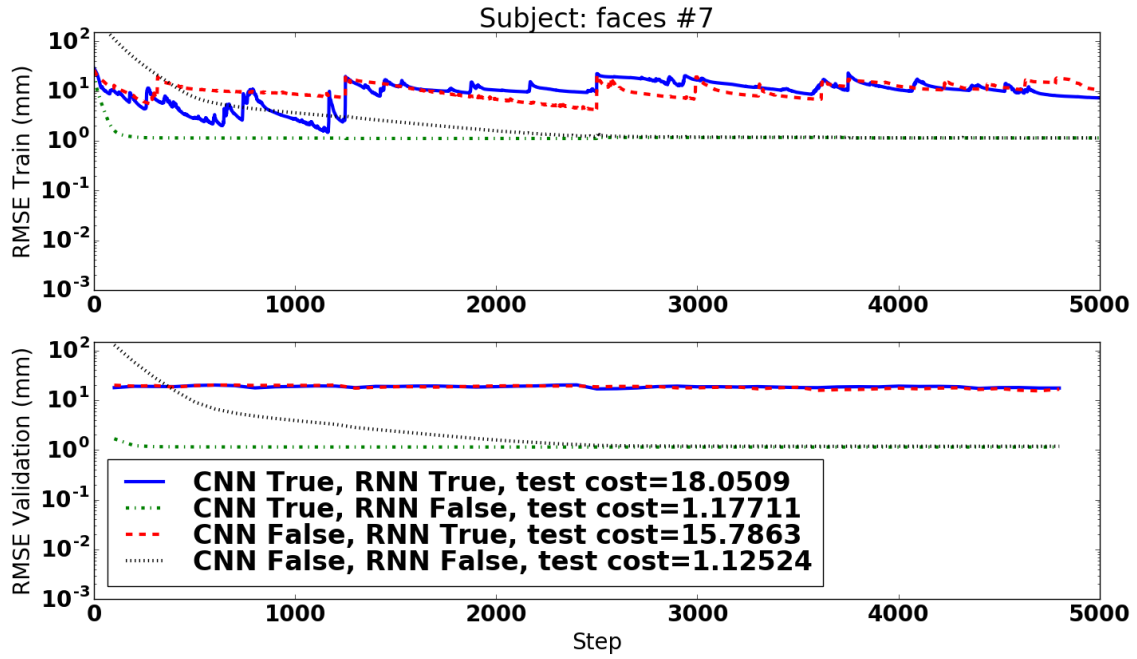


Fig. 10. Training/validation results for faces stimulus dataset, subject #7 (Best of 100 dipoles).

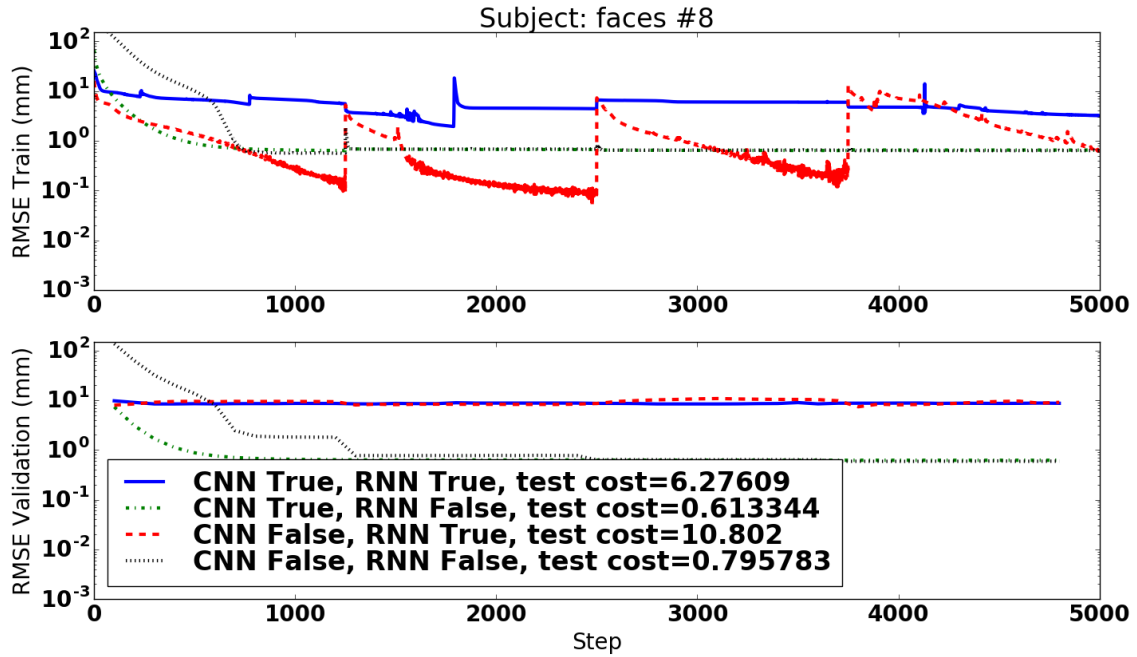


Fig. 11. Training/validation results for faces stimulus dataset, subject #8 (Best of 100 dipoles).

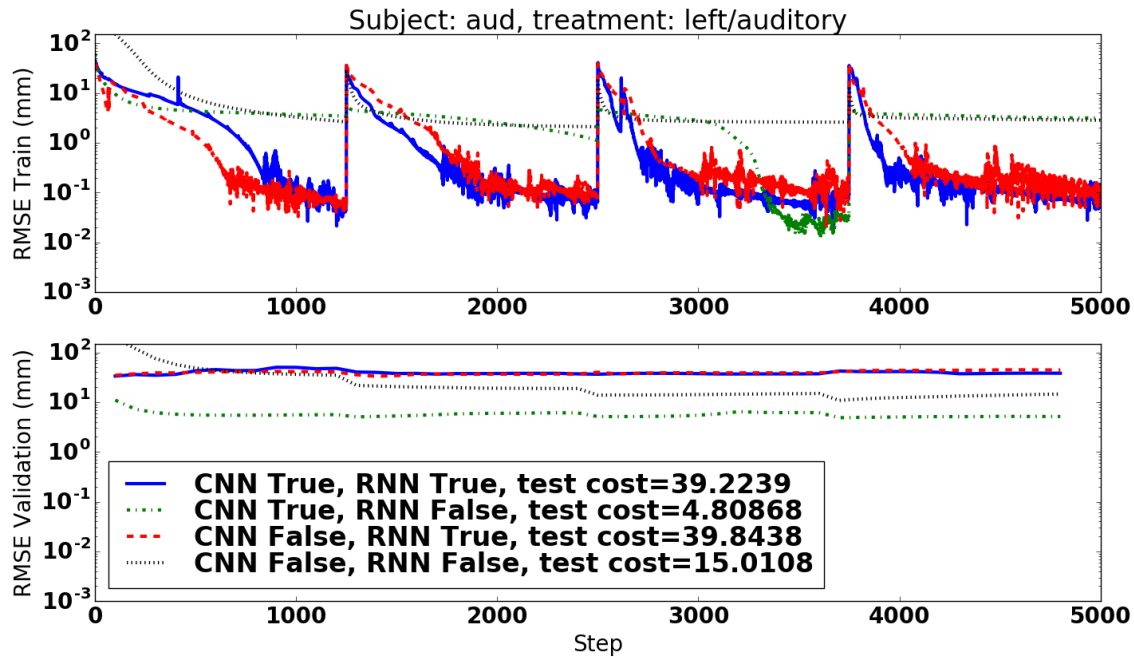


Fig. 12. Training/validation results for auditory stimulus dataset, only considering the left/auditory treatment (1 dipole).

Training and validation curves are given in Figs. 6, 7, and 8, along with test dataset cost, for the auditory dataset and the faces dataset, respectively. Mean and standard deviation of test cost are given in Table II. For both datasets, CNN and MLP networks outperform the networks including an RNN. While they are capable of achieving low training cost, they tend to overfit (the validation error remains flat), and thus do poorly on the test set (test cost of around 45 mm for the faces dataset, 30 mm for the auditory dataset). For reference, the human head is about 75 mm in diameter. On the other hand, simpler networks did better, with the CNN only network performing best, as measured by the RMSE on the test data set (cost of 5.1 mm for the auditory set, and 1.9 mm for the faces subject 7 dataset). This indicates a high degree of accuracy in locating the peak neuronal current, since this error in the spatial location of the dipole is near the minimum grid size use by MNE; in other words, it is at the level of the discretization error of our truth measurements. MNE relies on a voxel grid generated from the MRI to locate neurons, and so its level of discretization is the accuracy limit. Fewer parameters allowed for better generalization, and the CNN layer allows us to preserve spatial information critical for inversion. For all data sets, CNN-only networks and MLP networks had steadily

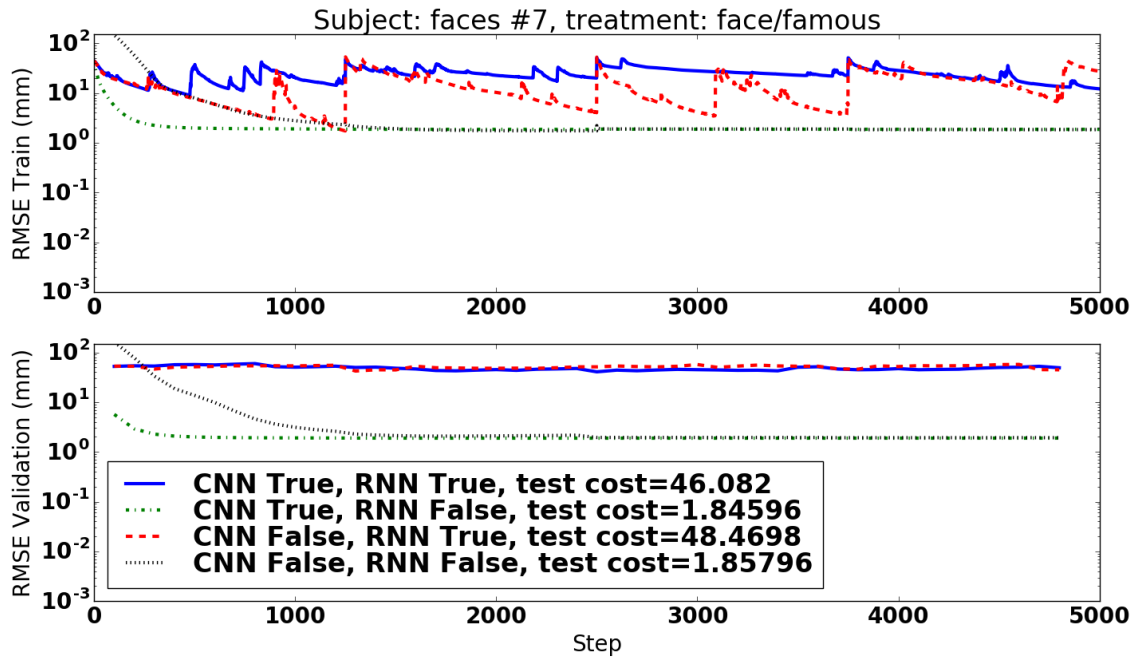


Fig. 13. Training/validation results for faces stimulus dataset on subject #7, only considering the famous face treatment (1 dipole).

declining training and validation errors, indicating better accuracy and generalization.

The poor generalization of the RNN networks led us to suspect that it may be too sensitive to noise in the location of the peak dipole. In other words, since we were using the MNE estimate of the dipole currents as truth, and these have some inherent error due to sensor noise and inversion error, the performance of the network to estimate the location may be poor. To test this, instead of comparing the neural network estimate to the peak dipole, we instead tried comparing it to the top 100 dipoles (which is approximately the top 5%-10%), and picking the closest one for calculating the cost. This has the effect of averaging out error in the dipoles estimated by MNE. Plots showing the training curves for this top-100 approach are shown in Figs. 9, 10, and 11. Clearly, this reduces the test, training, and validation costs, but the performance ranking of the four networks remains the same (from best to worst: CNN, MLP, CNN+RNN, and RNN), and the training and validation error curves show similar trends as in the single dipole case.

Finally, we attempted to test the networks' ability on datasets limited to a single treatment. In each dataset, there were several treatments - each treatment is just a different type of stimulus, meant to provoke a response in a different part of the brain. In the auditory dataset, the treatments

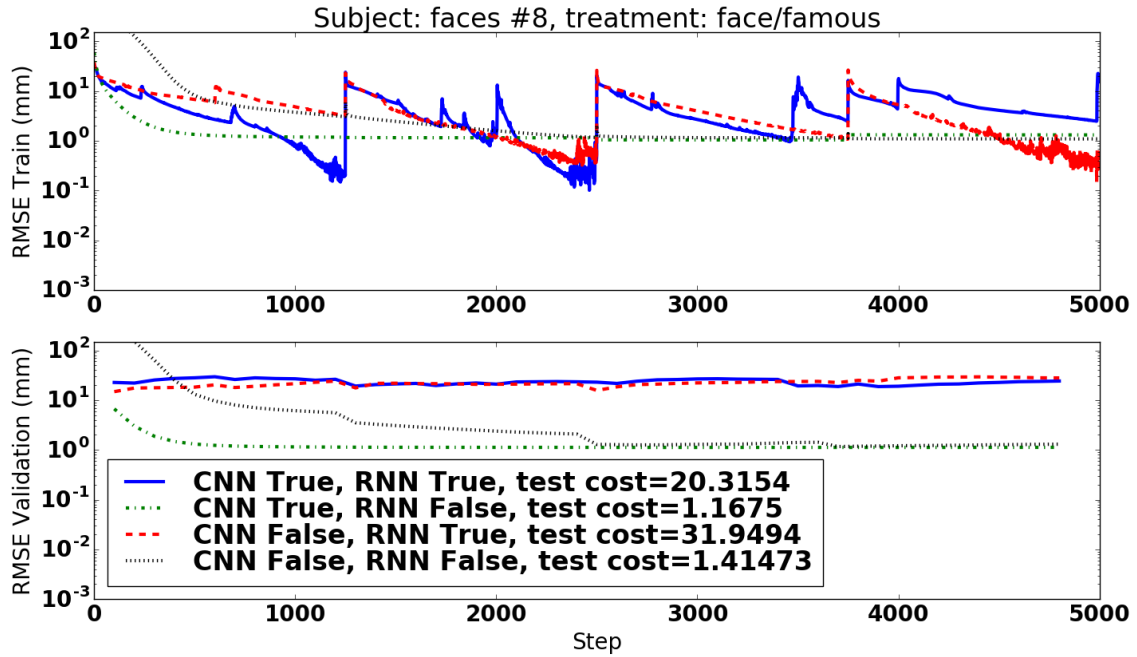


Fig. 14. Training/validation results for faces stimulus dataset on subject #8, only considering the famous face treatment (1 dipole).

were checkerboard patterns presented into the left and right visual field, interspersed by tones to the left or right ear; in the faces dataset, the treatments were pictures of famous, unfamiliar, or scrambled faces. Our thinking was that we could train a different network for each type of treatment, then aggregate the networks, to provide a single best estimate. However, this did not have the intended effect. There is not much difference in the training curves or in the test errors when compared to the results for mixing all treatments together; see Figs. 12 (left-auditory stimulus), 13 (famous faces stimulus), and 14 (famous faces stimulus) for an example.

IV. CONCLUSION

We have presented a new technique for rapid inversion of MEG/EEG data, using neural networks to estimate the location of the peak dipole, rather than the current distribution over the cortex. While training the network can take a long time, once the network parameters are known, applying the network to another set of inputs provides results in seconds, rather than the minutes required by MNE or other inversion techniques. This opens up the possibility of real-time neuronal diagnostics in a clinical setting. In examining four possible network architectures,

it was found that the CNN networks perform best in determining the location of the peak dipole at each timestep. More complicated network architectures, or those without the CNN layers, performed worse, due to overfitting and loss of spatial information. Treating the EEG and MEG data as a two-channel image is an effective method for retaining spatial field patterns, and is easily extensible to situations with more available sensors, such as NIRS. Future work may investigate using sequence-to-sequence neural machine translation to interpret timeseries.

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