Radboud University Nijmegen

PDF hosted at the Radboud Repository of the Radboud University Nijmegen

The following full text is a publisher's version.

For additional information about this publication click this link. http://hdl.handle.net/2066/81743

Please be advised that this information was generated on 2017-12-06 and may be subject to change.

Lecture presentation

Open Access

Realistic mean field forward predictions for the integration of co-registered EEG/fMRI

Ingo Bojak*, Thom F Oostendorp, Andrew T Reid and Rolf Kötter

Address: Donders Institute, Centre for Neuroscience, Radboud University Medical Centre, Nijmegen, Netherlands Email: Ingo Bojak* - i.bojak@donders.ru.nl * Corresponding author

from Eighteenth Annual Computational Neuroscience Meeting: CNS*2009 Berlin, Germany. 18–23 July 2009

Published: 13 July 2009 BMC Neuroscience 2009, 10(Suppl 1):L2 doi:10.1186/1471-2202-10-S1-L2

This abstract is available from: http://www.biomedcentral.com/1471-2202/10/S1/L2 © 2009 Bojak et al; licensee BioMed Central Ltd.

Brain activity can be measured non-invasively with functional imaging techniques. Each pixel in such an image represents a neural mass of about 10⁵ to 10⁷ neurons. Mean field models (MFMs) approximate their activity by averaging out neural variability while retaining salient underlying features, like neurotransmitter kinetics. However, MFMs incorporating the regional variability, realistic geometry and connectivity of cortex have so far appeared intractable. This lack of biological realism has led to a focus on gross temporal features of the EEG. We address these impediments and showcase a "proof of principle" forward prediction of co-registered EEG/fMRI for a fullsize human cortex in a realistic head model with anatomical connectivity, see figure 1.

MFMs usually assume homogeneous neural masses, isotropic long-range connectivity and simplistic signal expression to allow rapid computation with partial differential equations. But these approximations are insufficient in particular for the high spatial resolution obtained with fMRI, since different cortical areas vary in their architectonic and dynamical properties, have complex connectivity, and can contribute non-trivially to the measured signal. Our code instead supports the local variation of model parameters and freely chosen connectivity for many thousand triangulation nodes spanning a cortical surface extracted from structural MRI. This allows the introduction of realistic anatomical and physiological parameters for cortical areas and their connectivity, including both intra- and inter-area connections. Proper cortical folding and conduction through a realistic head model is then added to obtain accurate signal expression for a comparison to experimental data. To showcase the synergy of these computational developments, we predict simultaneously EEG and fMRI BOLD responses by adding an established model for neurovascular coupling and convolving "Balloon-Windkessel" hemodynamics. We also incorporate regional connectivity extracted from the CoCoMac database [1].

Importantly, these extensions can be easily adapted according to future insights and data. Furthermore, while our own simulation is based on one specific MFM [2], the computational framework is general and can be applied to models favored by the user. Finally, we provide a brief outlook on improving the integration of multi-modal imaging data through iterative fits of a single underlying MFM in this realistic simulation framework.

References

- Kötter R, Wanke E: Mapping brains without coordinates. Phil Trans R Soc B 2005, 360:751-766.
- 2. Bojak I, Liley DTJ: Modeling the effects of anesthesia on the EEG. Phys Rev E 2005, 71:041902.

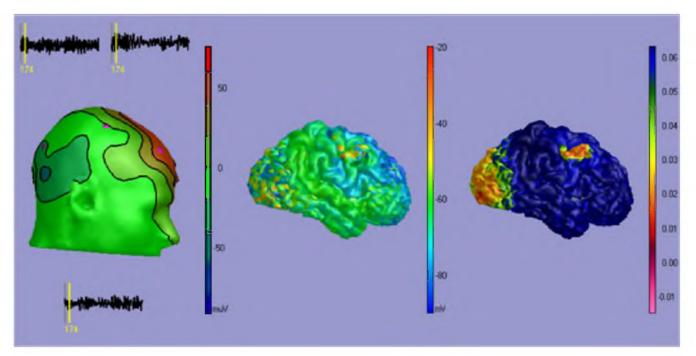


Figure I

MFM prediction of simultaneous **EEG** and **fMRI** with local, contralateral and visual connectivity. Left: EEG scalp potentials. *Middle*: MFM excitatory soma membrane potential h_e . *Right*: fMRI BOLD response.

