

POSTER PRESENTATION

Open Access

Nonlinear dynamics of large-scale activity in “networks of networks”

Fereshteh Lagzi^{1*}, Fatihcan M Atay², Stefan Rotter¹

From Twenty Second Annual Computational Neuroscience Meeting: CNS*2013
Paris, France. 13-18 July 2013

As a first step toward understanding the macro-dynamics of brain-like systems, we study the large-scale dynamics of balanced random networks of excitatory and inhibitory integrate-and-fire neurons. Based on the dynamical equations of the model, a mean field approach was previously employed to reduce the dimensionality of the network dynamics [1,2]. Here, we analyze the joint activity dynamics of excitatory and inhibitory populations employing a pair of mutually interacting nonlinear differential equations. In absence of a voltage leak for individual neurons, and for negligible synaptic transmission delay, these equations take the form of Lotka-Volterra equations. These have been used to describe predator-prey systems, corresponding to excitatory and inhibitory populations of neurons in our case. For non-zero identical synaptic transmission delay, we obtain Lotka-Volterra equations with delay. We try to infer the parameters for the non-autonomous differential equations given a dataset from numerical simulations of such a network. Moreover, we attempt to analytically constrain the parameters and compare them with their statistical estimators. Using simulation results, the significance of the nonlinear dynamics becomes obvious in the vector field of excitatory-inhibitory activity, which corresponds nicely with the vector field of the analytical equations.

We have analyzed the stability of the network considering two bifurcation parameters: the relative strength of recurrent inhibition, “ g ”, which controls the balance between excitation and inhibition in the network, and the intensity of external input to the network, “ η ”. We have found out that for a value of “ g ” that keeps the exact balance between excitation and inhibition, a bifurcation from unstable to stable network dynamics takes

place. This bifurcation separates Synchronous Regular (SR) from Asynchronous Irregular (AI) activity of the network, similar to what was found in a previous study on the same network using a Fokker-Planck approach [3]. The influence of synaptic delays on the reduced dynamics of the network is currently under study.

It has been shown that Lotka-Volterra equations are capable of representing switching dynamics between different states of neural networks [2,4]. Our analysis represents a first step toward analyzing the dynamics of more complex “networks of networks” that are implicated in various cognitive abilities of the brain.

Acknowledgements

This work is supported by the German Federal Ministry of Education and Research (BMBF; grant 01GQ0420 to BCCN Freiburg), the German Research Foundation (DFG; grant EXC 1086 to the Cluster of Excellence BrainLinks-BrainTools) and the European Commission (FP7-ICT grant 318723 to MatheMACS).

Author details

¹Bernstein Center Freiburg & Faculty of Biology, University of Freiburg, Germany.
²Max Planck Institute for Mathematics in the Sciences Leipzig, Germany.

Published: 8 July 2013

References

1. Cardanobile S, Rotter S: **Multiplicatively interacting point processes and applications to neural modeling.** *Journal of Computational Neuroscience* 2010, **28**(2):267-284.
2. Cardanobile S, Rotter S: **Emergent properties of interacting populations of spiking neurons.** *Frontiers in Computational Neuroscience* 2011, **5**:59.
3. Brunel N: **Dynamics of sparsely connected networks of excitatory and inhibitory spiking neurons.** *Journal of Computational Neuroscience* 2000, **8**(3):183-208.
4. Bick C, Rabinovich M: **On the occurrence of stable heteroclinic channels in Lotka-Volterra models.** *Dynamical Systems* 2010, **25**:97-110.

doi:10.1186/1471-2202-14-S1-P331

Cite this article as: Lagzi et al.: Nonlinear dynamics of large-scale activity in “networks of networks”. *BMC Neuroscience* 2013 **14**(Suppl 1): P331.

* Correspondence: fereshteh.lagzi@bcf.uni-freiburg.de

¹Bernstein Center Freiburg & Faculty of Biology, University of Freiburg, Germany
Full list of author information is available at the end of the article