

Poster presentation

Open Access

## Computational tools for assessing the properties of 2D neural cell cultures

Jugoslava Acimovic\*, Heidi Teppola, Jyrki Selinummi and Marja-Leena Linne

Address: Department of Signal Processing, Tampere University of Technology, P.O. Box 553, FI-33101 Tampere, Finland

Email: Jugoslava Acimovic\* - jugoslava.acimovic@tut.fi

\* 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):P170 doi:10.1186/1471-2202-10-S1-P170

This abstract is available from: <http://www.biomedcentral.com/1471-2202/10/S1/P170>

© 2009 Acimovic et al; licensee BioMed Central Ltd.

Neurons cultured in vitro provide a particularly promising experimental system for the analysis of properties, such as information coding, transmission, and learning, that are conventionally associated with biological neural networks. In these systems, isolated cells are placed on top of a recording plate (microelectrode array, MEA), where they spontaneously develop a random connectivity structure. Typical cultures consist of several thousands of neurons and the connectivity density varies from very low at the beginning of an experimental trial to high in mature cultures. In the absence of external stimuli, a culture exhibits a typical pattern of spontaneous activity, alternating intervals of slow spiking and bursting with the transition intervals of increasing activity. Spontaneous activity recorded in the cultures of rat cortical cells is described in [1,2] and an explanation of the phenomena is proposed in [3]. The behavior in the presence of external stimuli is also reported in the literature, for example, the adaptation exhibited in the presence of frequent and rare stimuli is assessed experimentally and through a computational model in [4]. The present work is related to the previously reported study [3] in which an image-processing algorithm is used to detect some structural parameters of cell cultures. A typical result from this study is illustrated in Figure 1. The original image of cultured cells on top of recording plate is shown in panel A, one of its segments in B, and the result of the applied algorithm in C. The blue pattern on panel C corresponds to cells. This approach, in general, enables automated estimation of parameters like the number of cells, or the average density of connections

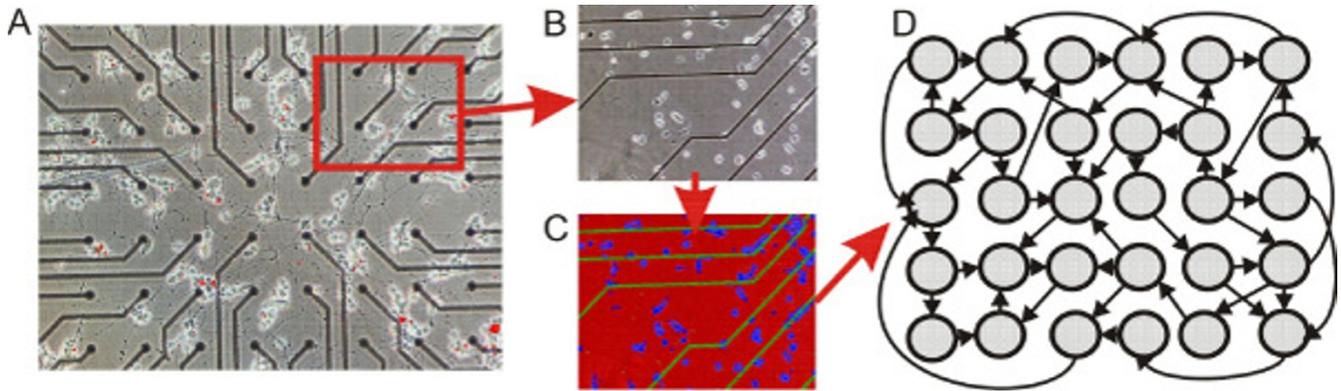
between the cells. Here, we propose a computational model based on the study in [3]. The neural network model is composed of leaky integrate-and-fire neurons, connected in a recurrent network as shown in panel D. The network is fed with the quantitative information about the structure of the cell cultures. Such model, although approximate, captures well the essential properties of the topologies observed in cultures. The presented model is used to reproduce and analyze network behavior observed in the absence of external stimuli. The structural parameters are estimated in different phases of development to closely relate them to the observed behavior. The relation between the network topology and behavior is systematically examined throughout this study.

### Acknowledgements

This work was supported by Academy of Finland.

### References

1. Wagenaar DA, Pine J, Potter SM: **An extremely rich repertoire of bursting patterns during the development of cortical cultures.** *BMC Neurosci* 2006, **7**:11.
2. Eckmann J-P, Jacobi S, Marom S, Moses E, Zbiden C: **Leader neurons in population bursts of 2D living neural networks.** *New J Phys* 2008, **10**:015011.
3. Teppola H, Sarkanen J-R, Selinummi J, Pettinen A, Jalonen TO, Linne M-L: **The effects of different MEA coating agents on growth and differentiation of human SH-SY5Y neuroblastoma cells.** *Proc. MEA meeting, 6th Int Meeting on Substrate-Integrated MEA. Reutlinger, Germany, 8–11 July 2008* **5**:105-106.
4. Wallach A, Eytan D, Marom S, Meir R: **Selective Adaptation in Networks of Heterogeneous Populations: Model, Simulation, and Experiment.** *PLOS Comput Biol* 2008, **4**:e29.



**Figure 1**

**From cell cultures to models of neural networks.** The technique for pattern recognition from images of cell cultures (reported in [1]) is used to assess the structural parameters incorporated in the computational model.

Publish with **BioMed Central** and every scientist can read your work free of charge

*"BioMed Central will be the most significant development for disseminating the results of biomedical research in our lifetime."*

Sir Paul Nurse, Cancer Research UK

Your research papers will be:

- available free of charge to the entire biomedical community
- peer reviewed and published immediately upon acceptance
- cited in PubMed and archived on PubMed Central
- yours — you keep the copyright

Submit your manuscript here:  
[http://www.biomedcentral.com/info/publishing\\_adv.asp](http://www.biomedcentral.com/info/publishing_adv.asp)

