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POSTER PRESENTATION

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# Homeostasis in large networks of neurons through the Ising model - do higher order interactions matter?

Dagmara Panas<sup>1\*</sup>, Alessandro Maccione<sup>2</sup>, Luca Berdondini<sup>2</sup>, Matthias H Hennig<sup>2</sup>

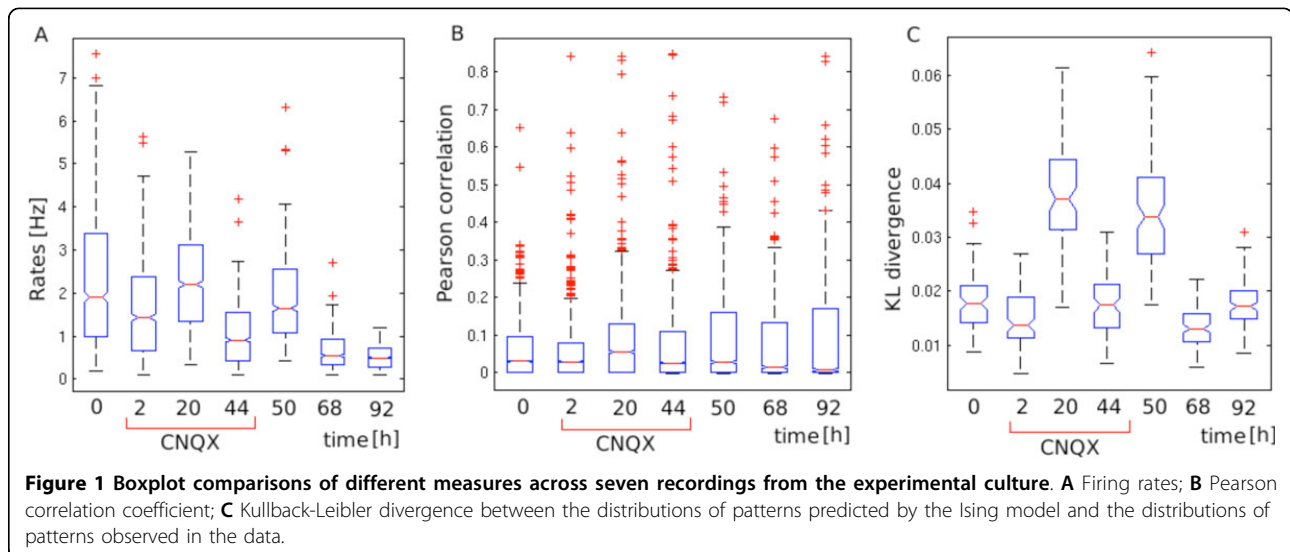
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Homeostatic activity in large networks of neurons is a relatively scanty explored area of neuroscience, both on experimental and computational level [1]. With recent advance in recording techniques, the lack of experimental data is gradually ceasing to be the limitation. New multi-electrode arrays (MEA) allow for monitoring cultures of thousands of neurons over many days with high spatial resolution [2]. However, the interpretation of multi-neuron recordings is not straightforward and requires methods going beyond the simplest descriptive statistics.

Here we explore a novel approach to analyzing multi-unit neuronal activity recorded over a five day homeostatic

experiment by employing the Ising model [3,4]. This statistical model explains the probability of multi-neuron spike patterns solely on the basis of firing rates and correlations, assuming an otherwise minimally structured distribution. Its application to a variety of recordings has helped re-evaluate the importance of neural interactions in shaping the global activity [3,4]. In addition, due to the models minimal structure, the quality of the fits can be treated as an indicator of higher-order interactions in the activity [4].

We compare the Ising model fits in the same preparation over several recordings: before, during and after CNQX application. We find that, in addition to the changes in



\* Correspondence: D.Panas@sms.ed.ac.uk

<sup>1</sup>Institute for Adaptive and Neural Computation, The University of Edinburgh, Edinburgh, EH8 9AB, UK

Full list of author information is available at the end of the article

firing rates and correlations, also the quality of the fits changes significantly across recordings (Figure 1). However, while firing rates and correlations do not appear to stabilize to a baseline level, the quality of the model fit does (Figure 1). Altogether this indicates that changes to first and second order statistics cannot explain the homeostatic changes in activity; and that higher order interactions might be a significant component of homeostatic compensation. Whether homeostatic maintenance of a complex higher-order dynamics is an effect of interplay of simple mechanisms or a global homeostatic set-point remains to be investigated.

#### Author details

<sup>1</sup>Institute for Adaptive and Neural Computation, The University of Edinburgh, Edinburgh, EH8 9AB, UK. <sup>2</sup>Department of Neuroscience and Brain Technologies, Italian Institute of Technology, 16163 Genova, Italy.

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