

MACACC – Modeling the Activity in the Cortex and Analysing the Cortical Neural Code.

ALCHEMY (INRIA); CORTEX (INRIA); INCM (CNRS); LJAD (U Nice-CNRS); ODYSSEE (INRIA).

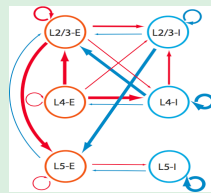
Neuronal information processing is related to the brain bio-electrical activity. Current neuro imaging techniques allow the measurement of this bio-electrical activity at different scales of time and space, from neurons to the brain as a whole (e.g. LFP, ECoG, EEG, MEG). But the analysis of data coming from these measurements requires the parallel development of suitable models. Namely, these models have to be, on one hand, close enough to phenomenology, taking into account the various types of bio-electrical activity and their scales relations, in order to propose a coherent representation of information processing in the brain (from neurons to neuronal populations, cortical columns, brain area, etc). On the other hand, these models must be well posed and analytically tractable. This requires a constant interaction between neurobiology, modeling and mathematics. In this spirit, this project aims to tackle the following questions.

Mesoscopic modeling of cortical columns, bifurcations, and imaging.

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Goals. Brain imaging techniques, like optical imaging or MEG-EEG, require a modeling of cortical brain activity at a spatial scale of order 0.1-1 mm². The goal of this project is to propose a mesoscopic model of the measured biological signal, at the space scale of a cortical column, and to analyse this model dynamics, using analytical methods and numerical simulations. These theoretical predictions will be then compared to the cortical activity of the visual system (area V1-V2), measured by optical imaging and MEG-EEG, in order to better understand the link between what is measured and the underlying cortical activity.

Methods. We want to characterize the activity of P neuronal populations in a cortical column, including non stationary dynamics and local field potential fluctuations. This is achieved by using dynamic mean-field methods and bifurcations analysis. This is a first step towards a mathematical and numerical characterization of local field potentials, exhibited by a realistic model of a cortical column with a connectivity scheme based on anatomical data, at a scale corresponding precisely to the resolution of optical imaging or MEG-EEG.



Then, we will compare the dynamical behaviour predicted by our analysis to real experimental data of the cortical activity of the visual system (V1/V2). This will help us to constrain the model parameters and structure in order to fit the real data.

Results.

1) Mean-field equations. Considering the evolution of an assembly of N neurons belonging to P populations where neurons are described by stochastic differential equations and are coupled by random synaptic connections, we have shown that the average activity of a population is described, when N is large, by explicit and well posed functional equations. Moreover, we provide a constructive and numerically tractable method for effectively computing their unique solution. These results shed new light on such neural mass models as the one of Jansen and Rit: their dynamics only approximate the much richer dynamics that emerge from our analysis because their approach neglects the random fluctuations around the mean values and their correlations [1].

2) Bifurcation analysis. Assuming that orientation columns in the visual cortex are coupled with symmetries allows an analysis of bifurcations via the study of spontaneous symmetries breaking. As a preliminary (master) work we have reconsidered the seminal work of Bressloff et al [2] about visual hallucinations using more recent techniques in bifurcation analysis of systems with symmetries [3].

Microscopic equations

$$d\tilde{V}^{(N)}(t) = (L^{(N)}(t)\tilde{V}^{(N)}(t) + \tilde{I}^{(N)} + \tilde{I}^{(N)}) dt + A^{(N)}(t) \cdot dW_t^{(N)}$$

$$\tilde{V}^{(N)} = \begin{pmatrix} 0 \\ 0 \\ 0 \\ 0 \\ 0 \\ 0 \end{pmatrix} \text{ and } \tilde{I}^{(N)} = \begin{pmatrix} 0 \\ 0 \\ 0 \\ 0 \\ 0 \\ 0 \end{pmatrix}$$

Mesoscopic equations

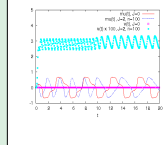
Effective interaction process

$$\begin{cases} \mathbb{E}[V_{\alpha}^X(t)] = \bar{J}_{\alpha\beta} m_{\alpha\beta}^X(t) \text{ where } m_{\alpha\beta}^X(t) \stackrel{\text{def}}{=} \mathbb{E}[S_{\alpha\beta}(X_{\beta}(t))]; \\ \text{Cov}(V_{\alpha}^X(t), V_{\alpha'}^X(s)) = \sigma_{\alpha\alpha'}^X \Delta_{\alpha\alpha'}^X(t, s) \text{ where} \\ \Delta_{\alpha\alpha'}^X(t, s) \stackrel{\text{def}}{=} \mathbb{E}[S_{\alpha\beta}(t)S_{\alpha'\beta}(s)]; \\ \text{Cov}(V_{\alpha}^X(t), V_{\alpha'}^Y(s)) = 0 \text{ if } \alpha \neq \alpha' \text{ or } \beta \neq \beta'. \end{cases}$$

Dynamic mean-field equations.

$$d\tilde{V}(t) = (L(t)\tilde{V}(t) + \tilde{I}^V + \tilde{I}(t)) dt + \Lambda(t) \cdot dW_t$$

Example: two populations model.



Evolution of the mean $\mu(t)$ and variance $v(t)$ for the mean field of population 1, over a time window [0,20]. Though Gaussian fluctuations are small, they have a strong influence on $\mu(t)$.

Next steps. On the basis of these theoretical results we intend to analyse what are the biological parameters compatible with experimental measurements in optical imaging and MEG-EEG. This will allow to determine precise constraints that must be introduced in neural masses models, for example the functional distinction between excitatory and inhibitory neurons, the dendritic and axonal activity, and the cortex laminar structure. As an important application, in the continuity of F. Grimbert's thesis [4], such models of cortical activity are prototypes for biological sources of voltage sensitive dye (VSD) imaging, at a mesoscopic scale corresponding to one pixel of optical imaging (~50µm). In a recent paper S. Chemla et al [5] have shown, using numerical simulations on an accurate model of cortical column, that the contribution of inhibitory cells, spiking activity and deep layers to the VSD signal is stronger than initially thought. It would be interesting to compare these results to a bifurcation analysis of a mesoscopic mean-field model.

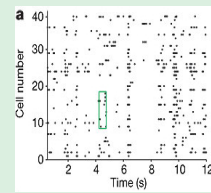
References.

- [1] O. Faugeras, J. Touboul, B. Cessac, «A constructive mean field analysis of multi-population neural networks with random synaptic weights and stochastic inputs», submitted to Frontiers in Neuroscience
- [2] P.C. Bressloff, J.D. Cowan, M. Golubitsky, P.J. Thomas, M.C. Wiener, «What geometric visual hallucinations tell us about the visual cortex?», Neural Comput. 14, 472-491 (2002).
- [3] O. Kodih, «Formation spontanée de structures dans le cortex visuel: un problème de bifurcations avec symétries», Master report (F. Chossat, B. Cessac, supervisors).
- [4] François Grimbert, PhD thesis, «Mesoscopic models of cortical structures», (2008).
- [5] S. Chemla, T. Vieville, F. Chavane, «Biophysical cortical column model for optical signal analysis», NeuroComp 2008.

Statistical analysis of spike trains

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Goals. Information transport by neurons is mediated by spikes trains. How information is encoded remains however an open issue. Thus, the analysis of spike trains obtained from in vivo or in vitro experimental data, requires suitable statistical models. Our goal is to propose a generic method to construct optimized statistical models from empirical data, using an approach combining mathematical modelling and analysis from in vivo experiments, together with numerical treatments. Beyond theoretical aspects, this approach is intended to provide new algorithms to improve spike trains analysis.



Methods. We want to characterize the spike dynamics of neural models, close enough to real neurons so that this analysis provides relevant issues in theoretical and experimental neuroscience. Using methods from ergodic theory and statistical physics we propose to approximate the empirical measure generated by a spike train by a Gibbs distribution, whose form is imposed by a variational principle. A special emphasis is put on the effects of synaptic plasticity, which leads to specific forms of spike probability distributions.

These theoretical foundations provide new algorithms for the analysis of real data provided by our biologist and neurophysiologist partners, taking a step further in the characterization of spike trains obtained from in vivo and in vitro data with the scope of providing a software allowing an automatic treatment of experimental data.

Results.

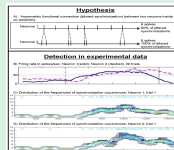
1) Neuron dynamics. We have made the first complete characterization of conductance based Integrate and Fire models dynamics with discrete time [1,2].

2) Gibbs measures. In these models, we have shown that the spikes train statistics, given the average value C_{α} of some specified observables ϕ_{α} , can be best approximated by a Gibbs measure with a specific potential [2], [3].

3) Synaptic plasticity. In this context, a general class of synaptic plasticity rules can be formulated as a variational principle. The spikes trains produced after adaptation are described by a Gibbs distribution that we have explicitly written down [3,4].

For the analysis of spike trains coming from numerical models or real data, we have developed numerical codes available at <http://enas.gforge.inria.fr>.

Next steps. Three complementary directions are investigated.



1) Analysis of experimental data. We are currently working on data obtained by INCM in experiments with monkeys. Recent investigations [5] suggest that two synchronized neurons may be asymmetrically connected to each other in functional terms, even with a similar firing rate and a common synchronized state (A&B). This implies that each neuron is able to join a synchronized assembly with its own dynamics, that fluctuates from trial to trial (C&D). We analyse this issue with our statistical tools [6].

2) Plasticity. Our results on synaptic plasticity open up the possibility of solving variational problems using spiking neural networks after a suitable training. Also, the role of specific neural architectures constrained by plasticity mechanisms will be investigated, in the continuity of [7] with possible extensions to intrinsic plasticity [8].

3) Application to vision tasks. Spikes appearing in vision processing can be precisely reproduced thanks to simulators [9]. Modeling further the vision system in a spiking architecture can also be considered for higher level tasks. For example, we showed in [10] that action recognition could be performed thanks to a spiking model of cortical layers V1-MT. Future work will focus on proposing plausible spiking models of vision-related cortical layers and see how the induced spiking activity can be analyzed for a vision task.

References.

- [1] B. Cessac, «A discrete time neural network model with spiking neurons. Rigorous results.», J. Math. Biol., 56, 3, 311-345 (2008).
- [2] B. Cessac, T. Vieville, «On Dynamics of Integrate-and-Fire Neural Networks with Adaptive Conductances.», Front. in Neurosci., 2008, Vol. 2, 2.
- [3] B. Cessac, H. Rostro, J.C. Vazquez, T. Vieville, «Statistics of spikes trains, synaptic plasticity and Gibbs distributions», NeuroComp 2008.
- [4] B. Cessac, H. Rostro, J.C. Vazquez, T. Vieville, «To which extend is the "neural code" a metric?», NeuroComp 2008.
- [5] F. Grammont, in preparation.
- [6] F. Grammont, B. Cessac, T. Vieville, in preparation.
- [7] B. Sini, H. Berry, B. Cessac, B. Delord, M. Quoy, «A mathematical analysis of the effects of Hebbian learning rules" to app. in Neural Comput.»,
- [8] J. Naukš, S. Geret, H. Berry, S. Mathon, J. Piaz, B. Delord, «A formalization of the computational impact of intrinsic plasticity», NeuroComp2008
- [9] A. Woherer and P. Kornprobst, «Virtual Retina: A biological retina model and simulator, with contrast gain control.», J. Comput. Neurosci. (2008).
- [10] M.-J. Escobar, G. S. Masson, T. Vieville and P. Kornprobst, «Action Recognition Using a Bio-Inspired Feedforward Spiking Network», in prep.

Conclusions. The present work relies on an approach combining theoretical methods from physics and mathematics, numerical treatments, and biological modeling / experimentation. At the present stage the two directions developed concerns different scales in the brain, but this separation is only apparent. As possible extensions we plan to study mean field methods for spiking neuron models and apply Gibbs statistical methods for the analysis of data at the scale of cortical columns.