

Modeling collective axon growth from in vivo data reveals the importance of physical axon-axon interactions

Agustina Razetti, Caroline Medioni, Grégoire Malandain, Florence Besse, Xavier Descombes

► To cite this version:

Agustina Razetti, Caroline Medioni, Grégoire Malandain, Florence Besse, Xavier Descombes. Modeling collective axon growth from in vivo data reveals the importance of physical axon-axon interactions. Cell biology of the neuron: Polarity, plasticity and regeneration, EMBO, May 2017, Heraklion, Greece. hal-01483505

HAL Id: hal-01483505 https://hal.archives-ouvertes.fr/hal-01483505

Submitted on 6 Mar 2017

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers. L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.

Modeling collective axon growth from *in vivo* data reveals the importance of physical axon-axon interactions

<u>Agustina Razetti¹, Caroline Medioni², Grégoire Malandain³, Florence Besse² and Xavier Descombes^{3*}</u>

¹. Université Côte d'Azur (UCA), University of Nice Sophia Antipolis (UNS), , France.

². Université Côte d'Azur (UCA), Institute of Biology Valrose (IBV), Centre National de la Recherche Scientifique (CNRS)-Unité Mixte de Recherche (UMR) 7277, Institut National de la Santé et de la Recherche Médicale (INSERM)-UMR1091, Nice, France.

³. Université Côte d'Azur (UCA), INRIA, France.

*All the authors belong to the Morpheme Team INRIA I3S IBV

Neurite extension is essential to establish complex neuronal circuits during brain development. In particular, neurons extend long cytoplasmic projections, the axons, in a crowded environment to reach target territories and connect to specific partners. Much work has been performed on cultured isolated neurons, focusing on the response of axon growing ends to chemical guidance cues. However, the cellular mechanisms involved in the growth of axon groups in their natural physical environment (here, the brain), are still poorly understood. Our objective is to shed light on collective axon growth and branching processes in a complex environment, with the help of mathematical modeling.

To study these mechanisms in the context of a living organism, we use *Drosophila* mushroom body gamma neurons as a paradigm. This population of neurons represents a good model for collective axon growth, as their adult axonal processes grow synchronously, in a relatively short time scale (around 15-20 hours), in a constraint environment (i.e. medial lobe of the Mushroom body inside the central brain). To feed our mathematical model we use an average pattern of medial lobe obtained from biological samples, and a database composed of confocal images of individual wild-type and mutants gamma neurons labeled with GFP.

Growth of individual axons is modeled by a Gaussian Markov chain in a 3D space, which depends on two main parameters estimated from real data: i- axon rigidity and ii- attraction to the target field. Furthermore, we hypothesize that axon tips pause when encountering a mechanical obstacle (i.e. other neurons or the lobe limits), and associate this behavior with the birth of long terminal branches. We show that the proposed mechanistic branch generation process is plausible. More importantly, our model predicts that branch formation in response to mechanical interactions enhances the probability that axons reach their final destination at the population level, and that axon density influences arborization patterns. Indeed, about 40% of simulated axons do not grow properly in the absence of branching in a wild-type environment, a result that is validated by biological data obtained with mutant neurons, in which axon growth defects are associated with defective branching.

To further validate the model hypotheses concerning branch formation, we are currently collecting and analyzing live-imaging data sets from brains with GFP-labeled growing axons.

We hope that combining cell biology, imaging and mathematical modeling will help us better understand the process of axon growth at the population level, in both normal and pathological contexts.