




Sinaps: A Python library to simulate voltage dynamic and ionic electrodiffusion in neurons

Nicolas Galtier² and Claire Guerrier¹

¹ CNRS, LJAD, Université Côte d'Azur ² Independent Researcher

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Software

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Summary

How neuronal dendrites collect, process and transmit information? What is the role of neurons specific geometry on neuronal activity? The old postulate that dendrites serve mainly to connect neurons and to convey information, with no specific role in cognitive processes, is currently challenged by the emergence of novel experimental techniques (M & M, 2005 ; R & R, 2018). Hence, the role of the dendritic tree in transforming synaptic input into neuronal output is now a leading question in developmental neuroscience. In particular, using genetically-encoded Ca^{2+} - indicators, state-of-the-art techniques have been developed to track the calcium dynamics within the entire dendritic tree, at a high spacial and temporal scales (K et al., 2020). Tracking neuronal activity through calcium dynamics is nevertheless ambitious. Calcium concentration fluctuations are known to reflect neuronal activity in a very complex way, as ions can flow from many sources that interact non-linearly with each other. There is thus an enhanced need for modeling to resolve what can be seen as an inverse problem: finding from experimental recordings the markers of synaptic activity, and distinguish in the calcium signals the different calcium sources.

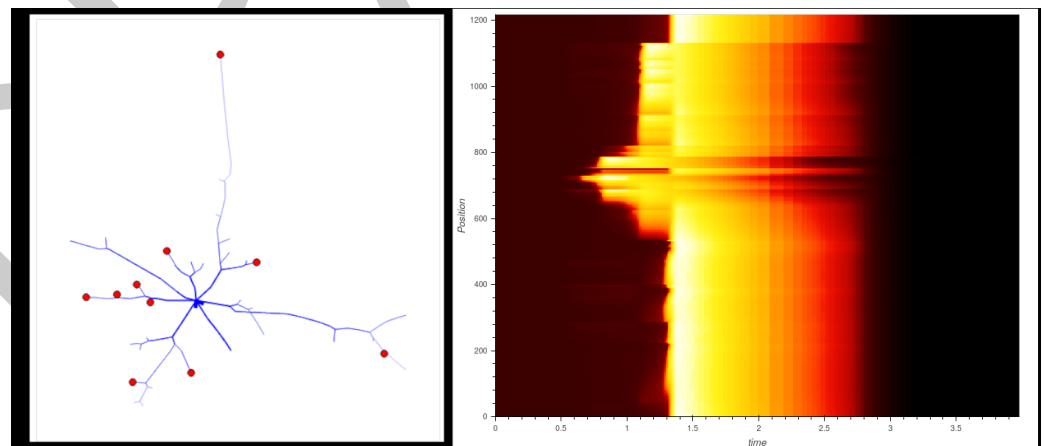


Figure 1: Left: a neuron complete geometry created using Sinaps. Right: Voltage propagation in a dendritic tree.

Statement of need

Sinaps is an easy-to-use Python library to simulate voltage propagation, ionic electrodiffusion and chemical reactions in neurons. It is based on the Cable equation for voltage propagation, coupled to the Nernst-Planck equation for ionic electrodiffusion. This library has been

24 designed for neuroscience laboratories using both an experimental and a modeling approach.
25 It includes the code to simulate voltage dynamic and ionic electrodiffusion, Hodgkin-Huxley
26 type membrane channels, and chemical reactions. Templates to code custom reaction-diffusion
27 mechanisms, as well as specific membrane channels are provided. We also provide the pos-
28 sibility to load a full morphometric geometry from data following neuromorpho.org file type
29 (swc file) [Figure 1](#).

30 Numerous softwares has been designed to realize simulations of voltage propagation in neurons
31 ([JM & D, 1998](#) ; [T & M, 2006](#)). While most of those softwares are using the simple Cable
32 theory model, and are designed toward neuronal networks simulation, our Python library is
33 designed to realize fast simulation of both voltage and ionic dynamics, taking into account
34 electrodiffusion of ions at a fine spatial scale. We also choose to realize the code in Python,
35 which has the advantage of having a code fully transparent with easy access to all the variables.
36 The class structure renders the code easily editable. Hence, our library provides an easy way to
37 simulate voltage and ionic dynamics, at the spacial scale reached by morphometric techniques,
38 and at a temporal scales not yet available for *in vivo* imaging of the full neuronal scale.

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