

# ARBOR AN INTRODUCTION

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## **Zooming In**

From Points to Dendrites

#### Def.: Point

The (adult) human eye has a resolution of roughly  $30\mu m$ .

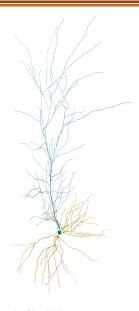
Thus going from a point to a typical neuron — a pyramidal cell — with dimensions of  $10\mu m$  (soma) –  $1\mu m$  (axon), we need to magnify by  $3-30\times$ .

At this level, we can model the electrical processes within a cell and resolve the distribution of dynamics across the cell's surface.

Cells will still communicate using action potentials.



#### **^**=arbor≡



- Design goals:
  Usability, performance portability, and strong separation of concerns.
- Made to exploit Multicore, SIMD, and GPU
- Scales extremely well: 768 nodes × 4 A100 on JUWELS booster. 70 Million cells.
- HPB since 2016 by CSCS and FZJ
- C++17 and Python3
- Linux and MacOS
- FOSS with a permissive BSD3 license
- Modern dev cycle in the open:
  Git, Code Review, CI, tests, sanitizers, ...

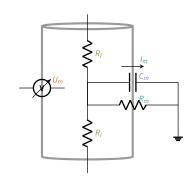


## The Bio-Physical Model

Consider a cell as a bi-lipid membrane suffused with ion channels, gating proteins, separating charged solutions of ions.

$$C_m \partial_t \underline{U_m} = \partial_x \left( \frac{1}{R_l} \partial_x \underline{U_m} \right) + I_m$$

- $C_m$ : Membrane capacitance.
- I<sub>m</sub>: Transmembrane current.
- *U<sub>m</sub>*: Membrane potential.
- R<sub>l</sub>: axial resistance.



The interplay of ion channels, in particular their dependence on the the membrane potential, creates action potentials and dynamics of the cells.

### **Performance**

