

## SODIUM TRANSIENTS IN AN ASTROCYTE

K. Lenk<sup>1</sup>, A. Tervonen<sup>2</sup>, J. Hyttinen<sup>2</sup>

<sup>1</sup>Institute for Neural Engineering, Graz University of Technology, Austria

<sup>2</sup>BioMediTech, Faculty of Medicine and Health Technology, Tampere University, Finland

kerstin.lenk@tugraz.at

**Abstract**— Astrocytes, a type of glial cells in the brain, are actively involved in neuronal information processing and memory formation. An astrocyte participates in neuronal activity by receiving neurotransmitters, e.g., glutamate, from an adjacent pre-synapse. This leads to the propagation of astrocytic intra- and intercellular calcium waves. In this study, we extend the computational model presented by Oschmann et al. (2017) to a finite element model of single astrocytes, including a realistic 2D geometry and inositol 1,4,5-trisphosphate ( $IP_3$ ) and ion diffusion. Here, we conduct a parameter exploration of the sodium-calcium exchanger (NCX) and sodium-potassium-ATPase (NKA) rate constants. The calcium concentration directly depends on the amplitude of the glutamate stimulus. However, the sodium concentration instead relates only to the sodium-potassium pump activity.

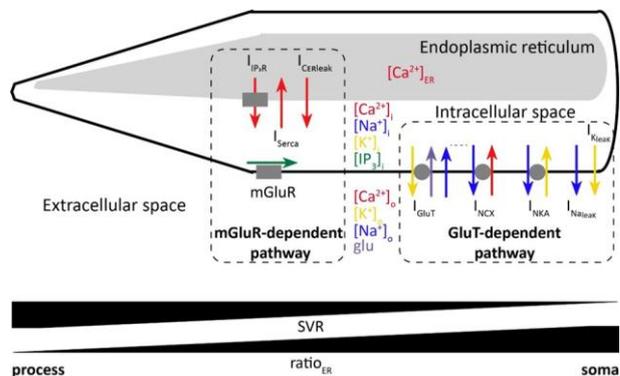
**Keywords**— Astrocyte, simulation, sodium, calcium, diffusion

### Introduction

Astrocytes are a glial cell type that plays a major role in the ion and energy homeostasis of the brain. Astrocytes and neurons are in a ratio of 1:2 in rodents and even 1:1 in humans. A single astrocyte can connect between 270 000 and 2 million neuronal synapses [1]. Thus, astrocytes can functionally modulate neuronal activity and complex mammalian behavior. In a landmark paper [2], mice with transplanted human astrocytes showed improved long-term potentiation and performance in learning tasks, emphasizing the importance of human astrocytes for the unique human cognitive abilities. Experimental findings strongly indicate that astrocytes might be involved in various pathologies such as epilepsy, depression, Alzheimer's, and Huntington's disease [3], [4]. Computational modeling can aid experimental work by forming a theoretical framework to characterize the anatomy and function of neurons and astrocytes [5]. Thus, a well-planned theoretical study can guide experimentalists towards the most relevant experiments, and in this way, save time and resources. In our review paper [5], we summarized astrocyte models from subcellular to network level that have started to emerge.

Oschmann et al. [6] introduced an astrocyte model including two glutamate pathways which both induce calcium ( $Ca^{2+}$ ) elevations (Fig. 1): activation of (1) the metabotropic glutamate receptors (mGluRs) and (2)

the glutamate transporters (mGluT) at the plasma membrane. The mGluR pathway includes the uptake of glutamate from the extracellular space. Subsequently, inositol 1,4,5-trisphosphate ( $IP_3$ ) followed by a  $Ca^{2+}$  release from the endoplasmic reticulum (ER) into the cytosol through  $IP_3$  receptor channels. The increasing  $Ca^{2+}$  levels lead to an opening of further  $IP_3$  channels and a  $Ca^{2+}$ -induced  $Ca^{2+}$  release (CICR). The second pathway includes the mGluTs, which transports potassium ( $K^+$ ) out and sodium ( $Na^+$ ) and glutamate into the cell. Furthermore, the sodium-calcium exchanger (NCX) exchanges  $Na^+$  and  $Ca^{2+}$  ions. The  $Na^+$ - $K^+$ -ATPase (NKA) transports  $Na^+$  ions out and  $K^+$  ions into the cell. Additionally, leak currents for  $Na^+$  and  $K^+$  are modeled. The novelty of the model is the combination of those two pathways and their dependence on the surface-to-volume ratio (SVR) and ER amount in the respective cell section. The SVR is large in perisynaptic astrocytic processes that connect to the neurons and small in the soma of the astrocyte. The ER ratio behaves the opposite.



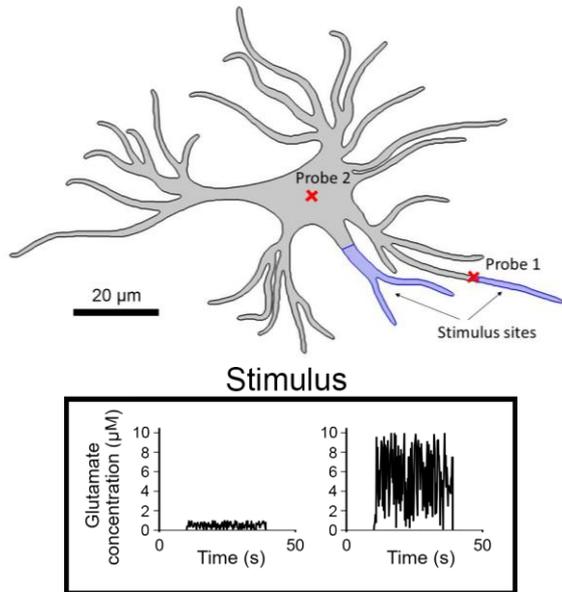
**Figure 1:** Scheme of the calcium dynamics within an astrocyte, including the mGluR- and the GluT-dependent pathway (adapted from [6]).

In this study, we combined the model by Oschmann et al. with a whole-cell astrocyte morphology [7]. Based on the width of a cell section, we automatically calculated the SVR. Furthermore, we added  $IP_3$  and ion ( $Ca^{2+}$ ,  $K^+$ , and  $Na^+$ ) diffusion to the model.

### Methods

We used the model and its parameters as described in Oschmann et al. [7] with the following additions: diffusion of  $IP_3$ ,  $Ca^{2+}$ ,  $K^+$ , and  $Na^+$  with coefficients 300, 13, 1 960, and 1 330  $\mu m^2/s$ , respectively. Our model was implemented as a finite-

element method with a realistic geometry (Fig. 2 top) in COMSOL Multiphysics version 5.6. The geometry is implemented in 2D as a tridomain model (i.e., ER, cytosol, and extracellular space). The SVR in the geometry required by the model was based on the width of the processes.



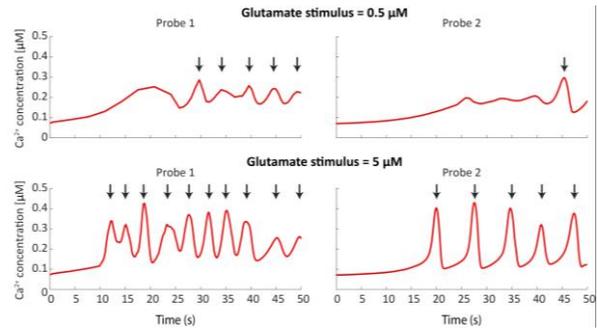
**Figure 2:** Top: Geometry of a single astrocyte with an indication of the stimulus sides (blue) and the locations of the two measurement probes (red crosses). Bottom: Random (uniform distribution) stimuli from 10 to 40 s with either 0.5 μM or 5 μM mean glutamate concentration.

We conducted a parameter exploration for the maximum pump current of NCX,  $I_{NCXmax}$ , with the values  $10^{-6}$ ,  $10^{-5}$ ,  $10^{-4}$ ,  $10^{-3}$ , 0.01, and 0.1  $\text{pA}/\mu\text{m}^2$ , and for the maximal pump activity of NKA,  $I_{NKAmx}$ , with the values  $10^{-6}$ , 0.1, 0.5, 1, 1.52, and 5  $\text{pA}/\mu\text{m}^2$ . Similar parameter ranges have also been explored in Oschmann et al. [6]. Furthermore, we applied two different random glutamate stimuli with a mean of around 0.5 and 5 μM (Fig. 2 bottom) to two processes (Fig. 2 top).

For the different  $I_{NCXmax}$  and  $I_{NKAmx}$  values and the two glutamate input values, we measured the  $\text{Ca}^{2+}$  and  $\text{Na}^+$  concentrations in one of the stimulated processes (Probe 1) and the soma of the astrocyte (Probe 2). Moreover, we counted the number of  $\text{Ca}^{2+}$  events (i.e.,  $\text{Ca}^{2+}$  spikes with a full width at half maximum of maximally five seconds). We also determined the maximum values of the  $\text{Na}^+$  concentration during the stimulation time as a function of the pump currents and the glutamate stimulus.

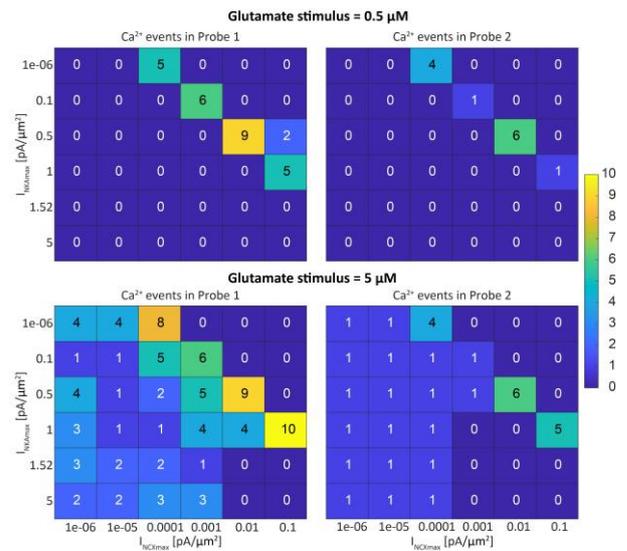
## Results

Fig. 3 shows the  $\text{Ca}^{2+}$  concentrations in Probe 1 and 2 for the two applied glutamate concentrations. The higher stimulus led to more distinct  $\text{Ca}^{2+}$  spikes.



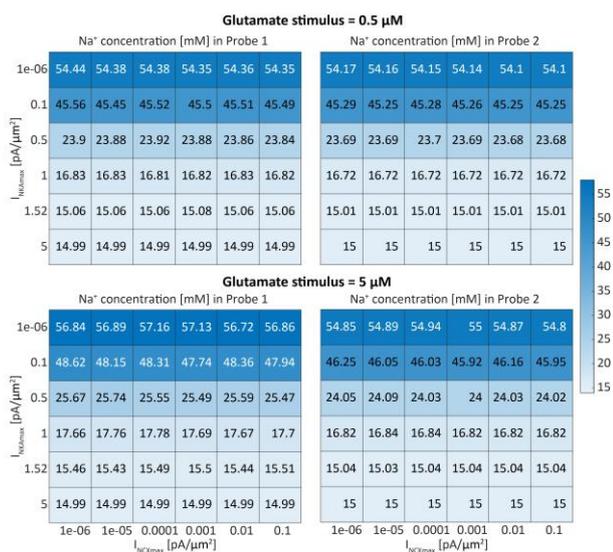
**Figure 3:**  $\text{Ca}^{2+}$  concentrations after stimulating the astrocyte with a glutamate concentration of 0.5 or 5 μM and applying  $I_{NCXmax} = 0.1 \text{ pA}/\mu\text{m}^2$  and  $I_{NKAmx} = 1 \text{ pA}/\mu\text{m}^2$ . Rows: Glutamate stimulus of 0.5 (top) or 5 μM (bottom). Columns:  $\text{Ca}^{2+}$  concentrations in Probe 1 (left) or 2 (right) during the stimulation time. The arrows indicate the  $\text{Ca}^{2+}$  peaks. The x-axis depicts the time in seconds.

The maximum number of  $\text{Ca}^{2+}$  events was yielded with a glutamate stimulus of 5 μM and  $I_{NCXmax} = 0.1 \text{ pA}/\mu\text{m}^2$  and  $I_{NKAmx} = 1 \text{ pA}/\mu\text{m}^2$  (Fig. 4). As expected, the number of spikes was higher for a higher glutamate stimulus. For smaller  $I_{NKAmx}$  values and higher  $I_{NCXmax}$  values, the most number of  $\text{Ca}^{2+}$  oscillations were detectable.



**Figure 4:** Number of  $\text{Ca}^{2+}$  events after stimulating the astrocyte with a glutamate concentration of 0.5 or 5 μM and different values for  $I_{NCXmax}$  and  $I_{NKAmx}$ . Rows: Glutamate stimulus of 0.5 (top) or 5 μM (bottom). Columns: Number of  $\text{Ca}^{2+}$  events (color bar) in Probe 1 (left) or 2 (right) during the stimulation time.

The maximum  $\text{Na}^+$  concentration was clearly dependent on the values of  $I_{NKAmx}$  rather than on the values of  $I_{NCXmax}$  and the glutamate concentration (Fig. 5). The lower the values of  $I_{NKAmx}$  was the higher the  $\text{Na}^+$  concentration.



**Figure 5:** Na<sup>+</sup> concentrations after stimulating the astrocyte with a glutamate concentration of 0.5 or 5 μM and different values for  $I_{NCXmax}$  and  $I_{NKAmx}$ . Rows: Glutamate stimulus of 0.5 (top) or 5 μM (bottom). Columns: maximum Na<sup>+</sup> concentrations (color bar) in Probe 1 (left) or 2 (right) during the stimulation time.

## Discussion

Our computational model investigates the intracellular calcium and sodium concentrations in dependence on the glutamate stimulus as well as the sodium-calcium exchanger, NCX, and the sodium-potassium-ATPase, NKA, pump rates. Therefore, we have extended the model by Oschmann et al. [7] by a realistic geometry and IP<sub>3</sub> and ion diffusion.

There is a complex interplay between the calcium and sodium dynamics in the astrocyte. Calcium signals and oscillations may be triggered by activated mGluRs [8], [9] as well as by the NCX in reverse mode [10] and the inhibition of NKA [11]. In a study by Ziemens et al. [12], the authors used the equation of the NCX model as also used in [7]. They found that after glutamate-induced stimulation, the NCX switches from an inward mode to an outward/ reverse mode, which is connected to a calcium influx to the astrocyte. Furthermore, the model suggests that the calcium increase only weakly antagonizes the NCX in the processes but not in the soma.

Our study suggests that the intracellular sodium concentration is mainly driven by the NKA. A blockage of the NKA transporter by ouabain led to an increase of cytosolic sodium concentrations and thus seems to be the primary cause for sodium elevations in the astrocyte [10].

To conclude, our simulations showed that the calcium levels depend on the given glutamate concentrations and a balance of the sodium-calcium exchanger and the sodium-potassium-ATPase pump rates.

The sodium levels rather relate only to the sodium-potassium pump activity.

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