

# 1 Calcium Microdomain Formation at the Perisynaptic Cradle due to 2 NCX Reversal: A Computational Study

3 John Joseph Wade<sup>1\*</sup>, Kevin Breslin<sup>1</sup>, KongFatt Wong-Lin<sup>2</sup>, Jim Harkin<sup>1</sup>, Bronac Flanagan<sup>1</sup>,  
4 Harm Van Zalinge<sup>3</sup>, Steve Hall<sup>3</sup>, Mark Dallas<sup>4</sup>, Angela Bithell<sup>4</sup>, Alexei Verkhratsky<sup>5,6</sup>, Liam  
5 McDaid<sup>1</sup>

6 <sup>1</sup>Computational Neuroscience and Neural Engineering (CNET) Research Team, Intelligent Systems  
7 Research Centre, Ulster University, Derry, United Kingdom.

8 <sup>2</sup>Neural Systems and Neurotechnology Research Team, Intelligent Systems Research Centre, Ulster  
9 University, Derry, United Kingdom.

10 <sup>3</sup>Department of Electrical Engineering and Electronics, University of Liverpool, Liverpool, United  
11 Kingdom.

12 <sup>4</sup>Reading School of Pharmacy, University of Reading, Reading, United Kingdom

13 <sup>5</sup>Faculty of Biology, Medicine and Health, University of Manchester, Manchester, United Kingdom.

14 <sup>6</sup>Achucarro Center for Neuroscience, IKERBASQUE, Basque Foundation for Science, 48011 Bilbao,  
15 Spain

## 16 \* Correspondence:

17 John Joseph Wade  
18 jj.wade @ ulster.ac.uk

19 **Keywords:** perisynaptic cradle, calcium microdomains, astrocytic process, Na<sup>+</sup>/Ca<sup>2+</sup> exchange,  
20 compartment model, glutamate transport, sodium dynamics

## 21 Abstract

22 It has recently been proposed using a multi-compartmental mathematical model that negatively fixed  
23 charged membrane-associated sites constrain the flow of cations in perisynaptic astroglial processes.  
24 This restricted movement of ions between the perisynaptic cradle (PsC), principal astroglial  
25 processes and the astrocyte soma gives rise to potassium (K<sup>+</sup>) and sodium (Na<sup>+</sup>) microdomains at the  
26 PsC. The present paper extends the above model to demonstrate that the formation of an Na<sup>+</sup>  
27 microdomain can reverse the Na<sup>+</sup>/Ca<sup>2+</sup> exchanger (NCX) thus providing an additional source of  
28 calcium (Ca<sup>2+</sup>) at the PsC. Results presented clearly show that reversal of the Na<sup>+</sup>/Ca<sup>2+</sup> exchanger is  
29 instigated by a glutamate transporter coupled increase in concentration of cytoplasmic [Na<sup>+</sup>]<sub>i</sub> at the  
30 PsC, which and instigates Ca<sup>2+</sup> influx through the NCX. As the flow of Ca<sup>2+</sup> along the astrocyte  
31 process and away from the PsC is also constrained by Ca<sup>2+</sup> binding proteins, then a Ca<sup>2+</sup>  
32 microdomain forms at the PsC. The paper also serves to demonstrate that the EAAT, NKA and NCX  
33 represent the minimal requirement necessary and sufficient for the development of a Ca<sup>2+</sup>  
34 microdomain and that these mechanisms directly link neuronal activity and glutamate release to the  
35 formation of localised Na<sup>+</sup> and Ca<sup>2+</sup> microdomains signals at the PsC. This local source of Ca<sup>2+</sup> can  
36 provide a previously underexplored form of astroglial Ca<sup>2+</sup> signaling.

37

38 **1 Introduction**

39 The concept of astroglial ionic excitability was established in 1990s following the discovery of  
 40 calcium ion ( $\text{Ca}^{2+}$ ) signaling and propagating  $\text{Ca}^{2+}$  waves in astrocytes in vitro and in situ (Cornell-  
 41 Bell et al., 1990; Cornell-Bell and Finkbeiner, 1991; Dani et al., 1992; Verkhratsky et al., 1998).  
 42 Initially, astroglial excitability has been thought to be mediated by  $\text{Ca}^{2+}$  ions; subsequent years have  
 43 demonstrated signaling roles for sodium ( $\text{Na}^+$ ) (Kirischuk et al., 2012; Parpura and Verkhratsky,  
 44 2012; Rose and Chatton, 2016; Rose and Verkhratsky, 2016) and chloride ( $\text{Cl}^-$ ) (Wilson and Mongin,  
 45 2018) while a signaling role for potassium ( $\text{K}^+$ ) has also begun to be considered (Breslin et al., 2018),  
 46 see also (Verkhratsky and Nedergaard, 2018). Ionic signaling in astrocytes serves several  
 47 physiological roles, in particular, coupling neuronal activity with astroglial homeostatic response  
 48 within the confines of the astroglial cradle that enwraps at least 50% of all synapses in the central  
 49 nervous system (Verkhratsky and Nedergaard, 2014; Rose and Verkhratsky, 2016; Verkhratsky and  
 50 Nedergaard, 2018). Astrocytes express elaborate molecular machinery controlling sodium ( $\text{Na}^+$ )  
 51 homeostasis and allowing transient intracellular  $\text{Na}^+$  increases in response to physiological activity in  
 52 neuronal networks (Kirischuk et al., 2012; Rose and Verkhratsky, 2016). The  $\text{Na}^+$  signals localized to  
 53 perisynaptic astroglial processes regulate the activity of numerous plasmalemmal transporters  
 54 responsible for a glutamine-glutamate (GABA) astroglial-neuronal shuttle, for  $\text{K}^+$  buffering, for  
 55 regulation of pH and for cellular metabolism as well as for secretion of reactive oxygen species  
 56 scavengers and various neuroactive molecules (Rose and Verkhratsky, 2016; Verkhratsky and  
 57 Nedergaard, 2016; Verkhratsky and Nedergaard, 2018). Cytoplasmic  $\text{Na}^+$  dynamics are therefore  
 58 directly linked to the functional activity of astrocytes and represent a mechanism for fast and local  
 59 signaling at the single synapse/perisynaptic process level. Although cytosolic  $\text{Na}^+$  has emerged as a  
 60 prominent ion at the interface between signaling and metabolic pathways (Chatton et al., 2016), the  
 61 spatiotemporal organization of cytosolic  $\text{Na}^+$  dynamics is far from being fully characterized.  
 62 Additionally, pathways that serve to trigger  $\text{Na}^+$  and  $\text{K}^+$  microdomains need to be resolved.

63 Computational modelling of astrocyte-neurone interaction is essential for the understanding of the  
 64 transport processes between cells. However, models of ionic signaling in astrocytes still have some  
 65 way to go if they are to fully capture the complex morphology of astrocytic processes seen in vivo.  
 66 Some of these models focus on changes of ionic fluxes crossing the astrocyte membrane controlled  
 67 by pumps and exchangers, for example  $\text{Na}^+/\text{K}^+$  pump (NKA) and  $\text{Na}^+/\text{Ca}^{2+}$  exchanger, (NCX) and  
 68  $\text{K}^+$ ,  $\text{Na}^+$  and  $\text{Cl}^-$  channels as well as axial fluxes of ions in the intracellular space (Halnes et al., 2013).  
 69 This electro-diffusive model allows channels to be distributed in different ways while gating  
 70 parameters can be altered to match experimental data. Intercellular  $\text{Ca}^{2+}$  waves and oscillations in  
 71 astrocytes (Pasti et al., 1997; Wade et al., 2011; Wade et al., 2012; Naeem et al., 2015) have also  
 72 been modelled to identify potential mechanisms of neurone to astrocyte intracellular intercellular  
 73 signaling.

74 **1.1 Astrocytic  $\text{Na}^+$  Signaling**

75 The major contributors to astrocyte  $\text{Na}^+$  homeostasis and signaling are the NKA, NCX and the  
 76 sodium-dependent glutamate transporters (EAAT1 and EAAT2). The NKA regulates resting  $\text{Na}^+$   
 77 concentration in the cytosol ( $[\text{Na}^+]_i$ ) and expels excess  $\text{Na}^+$  that enters astrocytes during periods of  
 78 neuronal activity, whereas glutamate transporters are responsible for the bulk of  $\text{Na}^+$  influx into  
 79 astrocytes accompanying glutamatergic synaptic transmission (Rose and Verkhratsky, 2016; Rose  
 80 and Karus, 2013; Kirischuk et al. 2007). The NKA, NCX and glutamate transporters co-localize in

81 astrocytic perisynaptic processes (Minelli et al., 2007; Melone et al., 2018), suggesting their  
82 functional coupling.

83 All three subunits of NCX (NCX1/SLC8A1, NCX2/SLC8A2 and NCX3/SLC8A3) are expressed in  
84 astroglia, with NCX1/SLC8A1 being the predominant isoform (Verkhatsky and Nedergaard, 2018;  
85 Pappalardo et al., 2014). The stoichiometry of astroglial NCX is  $3\text{Na}^+ : 1\text{Ca}^{2+}$ , and hence the  
86 equilibrium potential can be calculated from Nernst equation:  $E_{\text{NCX}} = (nE_{\text{Na}} - 2E_{\text{Ca}})/(n - 2)$  where  $n$  is  
87 a stoichiometry of  $\text{Na}^+$ , and  $E_{\text{Na}}$  and  $E_{\text{Ca}}$  are equilibrium potentials of  $\text{Na}^+$  and  $\text{Ca}^{2+}$ , respectively.  
88 Assuming  $[\text{Ca}^{2+}]_i$  of 50 - 80 nM and  $[\text{Na}^+]_i$  of 15 mM, the  $E_{\text{NCX}}$  could be as negative as  $\sim -85$  to  $-90$   
89 mV, in a similar range to measured resting membrane potential values of astrocytes (Verkhatsky and  
90 Nedergaard, 2018). As a result the NCX is prone to fluctuate between forward and reverse transport  
91 depending on actual changes in  $[\text{Ca}^{2+}]_i$  and  $[\text{Na}^+]_i$  and the astroglial membrane potential ( $V_m$ ).  
92 Conceptually, depolarization or an increase in  $[\text{Na}^+]_i$  will favor NCX operation in the reverse mode,  
93 whereas an increase in  $[\text{Ca}^{2+}]_i$  promotes the forward mode of the exchanger. In this way NCX can  
94 regulate both  $\text{Ca}^{2+}$  and  $\text{Na}^+$  signals, being relevant in shaping ionic signals in astroglial PsCs.

95 Recently we presented a new hypothesis (Breslin et al., 2018), which addressed ionic dynamics in  
96 thin ( $< 100$  nm) perisynaptic processes whereby negatively charged lipids form deep potential wells  
97 near the dipole heads restricting the flow of cations along the process. The negative ion foci serve to  
98 form “traps” that attract free cytosolic cations forcing them to hop from trap to trap, thus restricting  
99 ion propagation along the process and isolating them from the soma. This “ionic retention by traps”  
100 can potentially explain the generation of the transient  $\text{Na}^+$  and  $\text{K}^+$  microdomains at PsCs. The present  
101 paper aims to further develop this model with the inclusion of  $\text{Ca}^{2+}$  dynamics at the PsC. We stress  
102 here that while this model is not the focus of this paper, it is necessary to include it as it replicates  
103 closely the experimentally observed  $\text{Na}^+$  microdomain in astroglial processes (Langer and Rose,  
104 2009). Furthermore, the model allows us to test the hypothesis that reversal of the NCX, due to the  
105  $\text{Na}^+$  microdomain, leads to  $\text{Ca}^{2+}$  influx and this, coupled with  $\text{Ca}^{2+}$  retention along the thin astrocyte  
106 process, provides for a plausible mechanism for the generation of a  $\text{Ca}^{2+}$  microdomain at the PsC.  
107 Moreover, as this source of  $\text{Ca}^{2+}$  is remote from any endoplasmic reticulum (ER) mediated  $\text{Ca}^{2+}$   
108 release, we propose that this local source of  $\text{Ca}^{2+}$  may provide a previously under-explored form of  
109 astrocyte  $\text{Ca}^{2+}$  signaling.

## 110 2 Model

111 It was proposed in our earlier paper (Breslin et al., 2018) that  $\text{Na}^+$  and  $\text{K}^+$  retention occurs in thin  
112 astrocyte processes. In this paper we consider  $\text{Ca}^{2+}$  retention along the thin processes. This  $\text{Ca}^{2+}$   
113 retention is captured using an extended form of the multi-compartmental mathematical model  
114 described in (Breslin et al., 2018). The model consists of a single synapse surrounded by an  
115 astrocytic PsC. Due to the complexity of neuronal/astrocytic morphology (Patrushev et al., 2013;  
116 Lushnikova et al., 2000; Witcher et al., 2007; Xu-Friedman et al., 2001), this simplified cylindrical  
117 compartmental model (Breslin et al., 2018) simplifies the highly complex structures and the  
118 associated computational overheads, whilst retaining sufficient functionality to produce meaningful  
119 observations. All necessary dimensional details of the model can be found in Table 1. Figure 1  
120 provides details of the multi-compartmental model described by (Breslin et al., 2018).

121 In the previous study (Breslin et al., 2018),  $\text{K}^+$  and  $\text{Na}^+$  were modelled to determine their influence  
122 on ion retention in the thin astrocyte process. In the current study, we adapt this model to include  
123  $\text{Ca}^{2+}$  dynamics within the astrocyte PsC and extracellular space. From Figure 2, it can be seen that the

124 synapse and PsC contain a number of ionic channels, exchangers and pumps to provide homeostasis  
125 and dynamic exchange of ions between the two cells and extracellular space.

126 The neurone exchanges  $K^+$  and  $Na^+$  with the PsECS via a voltage-gated potassium channel ( $K_{Neu}$ ),  
127 voltage-gated sodium channel ( $Na_{Neu}$ ), a sodium potassium pump ( $NKA_{Neu}$ ), a potassium background  
128 channel ( $K_B$  on the synapse) and a sodium background channel ( $Na_B$  on the synapse). The astrocyte  
129 exchanges  $Na^+$ ,  $K^+$ , and  $Ca^{2+}$  with the PsECS via a background sodium transport ( $Na_B$  on the  
130 astrocyte), potassium background transport ( $K_B$  on the astrocyte), potassium inwardly rectifying  
131 channel ( $K_{ir}$ ), sodium-potassium-ATPase ( $NKA$ ), a glutamate-sodium-potassium-proton antiporter  
132 ( $EAAT1/2$ ), and the  $NCX$ . In both the astrocyte and neurone models, we define  $Na^+$  and  $K^+$   
133 background channels; although modelled as a single transport channel for each ion, these background  
134 channels represent a lumped model for  $Na^+$  and  $K^+$  transport, taking into account a multitude of  
135 influx and efflux pathways.  $I_{KPF}$ ,  $I_{NaPF}$  and  $I_{CaPF}$  model the flow of  $K^+$ ,  $Na^+$  and  $Ca^{2+}$  ions along the  
136 astrocytic process to the soma.  $I_{KECSL}$  and  $I_{NaECSL}$  models the  $K^+$  and  $Na^+$  ionic flow generated by  $K^+$   
137 and  $Na^+$  leaking from the PsECS to the GECS. The mathematical descriptions of ionic exchanges  
138 between a neurone and astrocyte at the PsC are now presented. For a more detailed description refer  
139 to Breslin et al. (2018).

## 140 2.1 Astrocyte Model

### 141 2.1.1 Membrane potential and ion concentrations

142 In this model the astrocyte resting membrane potential is taken as  $\sim -80$  mV, which is widely reported  
143 as the resting membrane potential for astrocytes (Verkhatsky and Nedergaard, 2018). We also  
144 assume that isopotentiality is preserved in the PsC as is the case for the astroglial syncytium (Kofuji  
145 and Newman, 2004), therefore the astrocyte membrane potential remains fixed at  $\sim -80$  mV.

146 The perisynaptic model described by (Breslin et al., 2018) comprises five compartments, namely  
147 PsC, PsECS, GECS, and the astrocyte process and soma. The astrocyte process is modelled as a long  
148 thin cylindrical channel that restricts the flow of cations along the process within the channel due to  
149 ion retention. In this work, each of these compartments contains three ionic concentrations,  $K^+$ ,  $Na^+$ ,  
150 and  $Ca^{2+}$ . All channels, exchangers and transporters permeable to these ions reside on the PsC. The  
151 kinetic equations for the changes of ionic concentration of each of these ions is given below. Note:  
152  $z_x F Vol_y$  is used to convert the total ionic current of ion  $x$  into a concentration for the volume  $y$ , where  
153  $z_x$  is the valency of ion  $x$ ,  $F$  is Faradays constant and  $Vol_y$  is the volume of compartment  $y$ . All initial  
154 conditions and parameters for this model are described in Tables 2 and 3 respectively. The change in  
155 PsC  $K^+$  concentration ( $[K^+]_{PsC}$ ) in the PsC is given by:

$$156 \quad \frac{d[K^+]_{PsC}}{dt} = - \left( \frac{I_{Kir} + I_{KNKA} + I_{KEAAT} + I_{KPF}}{z_K F Vol_{PsC}} \right)$$

157 where  $I_{Kir}$  is the  $K_{ir}$  channel current,  $I_{KNKA}$  is the  $K^+$  current through the astrocyte  $NKA$ ,  $I_{KEAAT}$  is the  
158  $K^+$  current created by the glutamate transporter and  $I_{KPF}$  is the  $K^+$  current flowing along the astrocyte  
159 process.  $K^+$  changes in the PsECS ( $[K^+]_{PsECS}$ ) is given by:

$$160 \quad \frac{d[K^+]_{PsECS}}{dt} = - \left( \frac{I_{KECSL} + I_{KNeu} - I_{Km}}{z_K F Vol_{PsECS}} \right)$$

161 where  $I_{KECSL}$  is current due to  $K^+$  leakage from the PsECS to the GECS,  $I_{KNeu}$  is the  $K^+$  current from  
 162 the neurone and  $I_{Km}$  is the total  $K^+$  current flowing through the astrocyte membrane.  $K^+$  is held  
 163 constant at baseline in the GECS and astrocyte soma compartments.

164 Changes in the PsC  $Na^+$  concentration ( $[Na^+]_{PsC}$ ) is given by:

$$165 \quad \frac{d[Na^+]_{PsC}}{dt} = - \left( \frac{I_{NaB} + I_{NaNKA} + I_{NaEAAT} + I_{NaNCX} + I_{NaPF}}{z_{Na}FV_{ol}_{PsC}} \right)$$

166 where  $I_{NaB}$  is a current due to  $Na^+$  influx across the membrane via  $Na^+$  permeable ion channels, this is  
 167 referred to as background  $Na^+$  channel (Breslin et al., 2018),  $I_{NaNKA}$  is the  $Na^+$  dependent current  
 168 component of the astrocyte NKA,  $I_{NaEAAT}$  is the  $Na^+$  current component of the glutamate transporter,  
 169  $I_{NaNCX}$  is the  $Na^+$  current component of the NCX and  $I_{NaPF}$  is the  $Na^+$  current flowing in the astrocyte  
 170 process.  $[Na^+]$  changes in the PsECS ( $[Na^+]_{PsECS}$ ) is given by:

$$171 \quad \frac{d[Na^+]_{PsECS}}{dt} = - \left( \frac{I_{NaECSL} + I_{NaNeu} - I_{Nam}}{z_KFV_{ol}_{PsECS}} \right)$$

172 where  $I_{NaECSL}$  is current due to  $Na^+$  leakage from the PsECS to the GECS,  $I_{NaNeu}$  is the  $Na^+$  current  
 173 from the neurone and  $I_{Nam}$  is the total  $Na^+$  current flowing through the astrocyte membrane.  $Na^+$  is  
 174 held constant at baseline in the GECS and astrocyte soma compartments.

175 Changes in the PsC  $Ca^{2+}$  concentration ( $[Ca^{2+}]_{PsC}$ ) is given by:

$$176 \quad \frac{d[Ca^{2+}]_{PsC}}{dt} = - \left( \frac{I_{CaNCX} + I_{CaPF}}{z_{Ca}FV_{ol}_{PsC}} \right)$$

177 where  $I_{CaNCX}$  is the  $Ca^{2+}$  dependent current component of the NCX, and  $I_{CaPF}$  is the  $Ca^{2+}$  current  
 178 flowing in the astrocyte process.  $[Ca^{2+}]$  changes in all other compartments are not considered and  
 179 remain constant at baseline.

### 180 **2.1.2 Glutamate Transporter (EAAT1/2)**

181 Glutamate released into the extracellular PsECS in the course of neurotransmission is assumed here  
 182 to be entirely removed by astrocytic EAAT1/2. A transport cycle involves the co-transport of 3  $Na^+$   
 183 and 1  $H^+$  with 1 glutamate and counter-transport of 1  $K^+$  (Grewer et al., 2014; Murphy-Royal et al.,  
 184 2015). EAAT1/2 proteins are trafficked to the plasma membrane to facilitate the rapid removal  
 185 (~3ms) of glutamate from the cleft. Glutamate bound to these proteins is then transported to the  
 186 astrocytic cytosol over a longer period: in this work a complete transport cycle is assumed to be 30  
 187 ms (Otis and Kavanaugh, 2000; Zhou and Danbolt, 2013). Existing EAAT mathematical  
 188 formulations do not capture adequately this rapid binding and slow release function and therefore in  
 189 this work we adopt a different approach. To model the stoichiometry and cycle rate we assume that  
 190 initially a release of glutamate instantaneously binds to membrane-bound proteins and thereafter the  
 191 flux of  $Na^+$  through the EAAT transporter pore follows an exponentially decaying rate given by:

$$192 \quad \frac{dJ_{NaEAAT}}{dt} = - \frac{J_{NaEAAT}(t)}{\tau} + J_0\delta(t - tsp)$$

193 where  $J_{NaEAAT}$  is the flux rate of  $Na^+$  through the EAAT1/2,  $J_0$  is the max flux rate through the  
 194 transporter,  $\delta$  is the Dirac Delta function,  $t$  is time, and  $tsp$  is the previous neuronal spike time. In our

195 model we view the membrane as a capacitor charged with bound glutamate and  $J_{NaEAAT}$  as a  
196 discharging flux.

197 The  $Na^+$  current through the transporter can be calculated by:

$$198 \quad I_{NaEAAT} = \frac{-J_{NaEAAT} z_{Na} F Vol_{PsECS}}{SA_{PsC}}$$

199 where  $J_{NaEAAT}$  is the  $Na^+$  flux through the EAAT co-transporter,  $z_{Na}$  is the valency of  $Na^+$ ,  $F$  is the  
200 Faradays Constant,  $Vol_{PsECS}$  is the volume of the perisynaptic ECS and  $SA_{PsC}$  is the surface area of  
201 the PsC.

202 The associated  $K^+$  current through the transporter is given by:

$$203 \quad I_{KEAAT} = \frac{-I_{NaEAAT}}{3}$$

204 Note that we are not considering glutamate transport to the astrocytic cytosol as we are only  
205 interested in  $Na^+$ ,  $K^+$  and  $Ca^{2+}$  dynamics. Additionally, our model for EAAT1/2 transport would need  
206 further consideration to include the dependency of fluxes on intra and extra cellular ionic  
207 concentrations.

### 208 **2.1.3 Sodium Calcium Exchanger (NCX)**

209 The NCX is a reversible antiporter which uses the electrochemical gradient of  $Na^+$  to exchange 3  $Na^+$   
210 ions for 1  $Ca^{2+}$  ion across the membrane. Depending on the membrane potential and transmembrane  
211  $Na^+$  gradient, the transporter operates either in forward mode ( $Na^+$  is transported into the cell while  
212  $Ca^{2+}$  is extruded) or in the reversed mode (providing for influx of  $Ca^{2+}$  and efflux of  $Na^+$ ) (Jeffs et al.,  
213 2007).

214 The  $Na^+$  current component of the transporter is given by (Gabbiani and Cox, 2010):

$$215 \quad I_{NaNCX} = \left( \bar{I}_{NCX} \left( \frac{[Na^+]_{PsC}}{[Na^+]_{PsECS}} \right)^3 e^{\frac{\gamma FV}{RT}} - \left( \frac{[Ca^{2+}]_{PsC}}{[Ca^{2+}]_{PsECS}} \right) e^{\frac{(\gamma - 1)FV}{RT}} \right) SA_{PsC}$$

216 where  $\bar{I}_{NCX}$  is the NCX exchanger conductance and  $\gamma$  is a partition parameter.

217 The  $Ca^{2+}$  current component is given by:

$$218 \quad I_{CaNCX} = -2 \left( \frac{I_{NaNCX}}{3} \right)$$

### 219 **2.1.4 Leakage from perisynaptic ECS to global ECS**

220 The diffusion of  $K^+$  and  $Na^+$  between the PsECS and the GECS is modelled as a simple  
221 electrochemical gradient controlled channel in which a zero extracellular potential is assumed and is  
222 given by:

$$223 \quad I_{iECSL} = g_{iECS} E_{iECS} SA_{ECSL}$$

224 where  $i$  is the ion under consideration,  $g_{iECS}$  is the conductance of the channel,  $SA_{ECSL}$  is the surface  
 225 area between the PsECS and the GECS, and  $E_{iECS}$  is the Nernst like potential of the channel given by:

$$226 \quad E_{iECS} = \frac{RT}{F} \ln \left( \frac{[i^+]_{PsECS}}{[i^+]_{GECS}} \right)$$

### 227 2.1.5 Astrocyte Process Ionic Transport Model

228 Breslin et al. (2018) proposed that ion retention within thin astrocyte processes can give rise to the  
 229 formation of  $K^+$  and  $Na^+$  microdomains at the PsC. This localisation of astroglial ionic microdomains  
 230 arises because in thin processes, surface conduction dominates over volume conduction, and because  
 231 membrane lipids are negatively charged, deep potential wells form near the dipole heads restricting the  
 232 flow of cations along the process. Therefore, cations must hop from well to well which restricts ion  
 233 conduction along the membrane. This hopping effectively semi-isolates the PsC from the astrocytic  
 234 main body allowing the formation of  $K^+$  and  $Na^+$  microdomains at the PsC under different conditions.

235 Breslin et al. (2018) proposed that the current flow  $I_{iPF}$  (see figure 2) through the thin process, due to  
 236 ionic hopping can be represented as:

$$237 \quad I_{iPF} = K_i \frac{V_A - V_m - V_r}{l} \exp \left[ - \frac{Q_i \left( \phi_w - \sqrt{\frac{Q_i(V_A - V_m - V_r)}{l\pi\epsilon}} \right)}{k_B T} \right] CSA_P$$

238 where  $K$  is a constant which represents mobility and concentration of mobile ions,  $V_m$  is the resting  
 239 membrane potential of the astrocyte,  $\phi_w$  is the well activation energy or potential barrier to ion flow,  $l$   
 240 is the length of the process,  $Q$  is the charge on a single ion taken as the charge on an electron,  $T$  is the  
 241 absolute temperature,  $CSA_P$  is the cross-sectional area of the process,  $\epsilon$  is the dynamic permittivity and  
 242 is given by  $\epsilon = \epsilon_0 \epsilon_r$ , where  $\epsilon_0$  is the absolute permittivity and  $\epsilon_r$  is the relative permittivity of the  
 243 cytoplasm, and  $k_B$  is the Boltzmann constant.

244 The concentrations of  $K^+$ ,  $Na^+$ , and  $Ca^{2+}$  in the astrocyte soma are held constant but will be  
 245 continuously changing at the PsC thus establishing a dynamic concentration gradient associated with  
 246 these cations. Consequently, we formulate a Nernst-like reversal potential for  $Na^+$ ,  $K^+$  and  $Ca^{2+}$   
 247 between the astrocyte soma (AS) and the PsC as:

$$248 \quad V_r = \frac{RT}{F} \ln \left( \frac{[i]_{AS}}{[i]_{PsC}} \right)$$

249 where  $i$  is the ion under consideration. A more in-depth discussion and assumptions of the full ionic  
 250 transport mechanism along thin astrocyte processes is given in (Breslin et al., 2018).

## 251 2.2 Neurone model

252 The neuronal model utilized in this work consists of the biophysical Hodgkin and Huxley (HH) type  
 253 model described in (Breslin et al., 2018) with the addition of a voltage-gated  $Na^+$  channel,  $K^+$   
 254 background channel and an  $Na^+$  background channel. These background channels reflect implicit  
 255 influx/efflux pathways, necessary for the system to ensure dynamic equilibrium. All parameter values  
 256 for the neurone model are described in Table 4. For reasons of simplicity, the internal concentrations

257 of neurone  $\text{Na}^+$  and  $\text{K}^+$  remain constant. Whilst we recognize that the neuron NKA is driven by  
 258 internal  $\text{Na}^+$  and astrocyte NKA is driven by external  $\text{K}^+$ , as we are not considering internal neurone  
 259  $\text{Na}^+$  concentration change, we apply the same NKA model for each, altering the maximum pump  
 260 rates accordingly.

### 261 2.2.1 Voltage-Gated Neuronal Sodium Channel ( $\text{Na}_{\text{Neu}}$ )

262 The HH model simulates current flow of  $\text{Na}^+$  through a voltage-gated channel, therefore the current  
 263 flow of  $\text{Na}^+$  from the neurone can be modelled as:

$$264 \quad I_{\text{NaNeu}} = -g_{\text{NaNeu}}m^3(V_{\text{Neu}} - E_{\text{NaNeu}})SA_{\text{Syn}}$$

265 where  $g_{\text{NaNeu}}$  is the maximum  $\text{Na}^+$  channel conductance,  $E_{\text{NaNeu}}$  is the reversal potential of the sodium  
 266 channel,  $V_{\text{Neu}}$  is the membrane voltage of the neurone and  $SA_{\text{Syn}}$  is the surface area of the synapse.

### 267 2.2.2 Neuronal Background ion channels ( $\text{Na}_B / \text{K}_B$ )

268 In this model, there are two individual background ion channels for  $\text{Na}^+$  and  $\text{K}^+$ . These channels use  
 269 the electrochemical gradient between the PsC and ECS, resulting in an influx of  $\text{Na}^+$  and efflux of  $\text{K}^+$   
 270 under normal physiological conditions. They were modelled as a simple passive electrochemical  
 271 gradient dependent channel given by (Steratt et al., 2011):

$$272 \quad I_{iB\text{Neu}} = g_{iB\text{Neu}}(V_{\text{neu}} - E_i)SA_{\text{Syn}}$$

273 where  $i$  is the ion under consideration,  $g_{iB\text{Neu}}$  is the channel conductance. Note: the value of  $g_{iB\text{Neu}}$  is  
 274 chosen in such a way that the total flux of ion  $i = 0$  at steady state.  $V_{\text{neu}}$  is the neurone membrane  
 275 voltage,  $SA_{\text{Syn}}$  is the surface area of the neuronal synapse, and  $E_i$  is the channel Nernst potential and is  
 276 given by:

$$277 \quad E_i = \frac{RT}{F} \ln \left( \frac{[i^+]_{\text{PsECS}}}{[i^+]_{\text{Syn}}} \right)$$

278 Note that the concentrations of  $\text{K}^+$  and  $\text{Na}^+$  within the neuronal synapse are held at baseline.

279 The complete astrocyte/neurone model was implemented using Matlab 2015b 64-bit (Windows  
 280 version) by Mathworks. All simulation results presented in the results section of this paper used the  
 281 forward Euler method of integration with a fixed time step of  $\Delta t = 10\mu\text{s}$ .

## 282 3 Results

283 This section reports the outcomes of a series of simulations that demonstrate the formation of a  $\text{Ca}^{2+}$   
 284 microdomain at the PsC. The simulations show that the  $\text{Ca}^{2+}$  microdomain is a direct result of ion  
 285 retention along the thin astrocyte process. We have shown in a recent paper (Breslin et al., 2018) that  
 286 ion retention underpins  $\text{Na}^+$  and  $\text{K}^+$  microdomain formation at the PsC during physiological neuronal  
 287 excitation. In this work we show that the uptake of  $\text{Na}^+$ , via EAAT channels during neuronal  
 288 stimulus, creates the  $\text{Na}^+$  microdomain in the PsC thereby causing the NCX to reverse with  
 289 subsequent formation of a  $\text{Ca}^{2+}$  microdomain in the PsC.

### 290 3.1 NCX reversal under physiological stimulation



291 To explore the reversal of the astrocyte NCX, a series of simulations were carried out with the  
292 presynaptic neurone stimulated using an external current to produce firing rates of 10 Hz, 20 Hz and  
293 30 Hz respectively. The neuronal stimulus has a duration of ~1 minute where the first 0.1 minute  
294 allows the model to reach a steady state condition and the stimulus ceases after 1 min. This long  
295 stimulus period allows investigation of what effect sustained neural activity has on the  
296 intracellular/extracellular ionic concentrations. In these simulations PsECS  $\text{Ca}^{2+}$  is held constant at  
297 baseline, however  $\text{K}^+$  and  $\text{Na}^+$  are permitted to change via the neurone and astrocyte  $\text{K}^+$  and  $\text{Na}^+$   
298 channels. Each time the neurone spikes, it is assumed that there is 1mM of glutamate released into  
299 the PsC and there are enough EAAT1/2 transports which allow the instantaneous binding of all the  
300 extra glutamate. Therefore, the extracellular glutamate concentration always remains at baseline  
301 concentration. Furthermore, the maximum flux rate of EAAT1/2 ( $J_0$ ) is tuned such that 3 mM of  $\text{Na}^+$   
302 is taken up from the PsECs and 1 mM of  $\text{K}^+$  is released into the PsECS by the EAAT1/2 over a  
303 period of 30 ms. Moreover, the astrocyte membrane voltage is held constant at ~ -80 mV, in line with  
304 reported resting membrane potential for astrocytes, which incidentally is close to the reversal  
305 potential for the NCX. Therefore, during periods of homeostatic rest there is no net flow of  $\text{Ca}^{2+}$  or  
306  $\text{Na}^+$  across the membrane associated with the NCX.

307 The results presented in Figure. 3A show that during periods of neural stimulus,  $\text{K}^+$  ions released by  
308 the neurone, build up in the PsECS and are cleared by the astrocyte, which results in a  $\text{K}^+$   
309 microdomain formation at the PsC (Figure 3B). At the start of the neuronal stimulation, there is a  
310 transient loss of  $\text{Na}^+$  from the PsECS (Figure 3C), due to neuronal depolarization and astrocytic  
311 EAAT transport. Furthermore, the transport of  $\text{Na}^+$  across the astrocyte membrane via  $\text{Na}^+$   
312 background channels and EAAT transporters results in a  $\text{Na}^+$  microdomain formation (Figure 3D).

313 In addition to  $\text{Na}^+$  and  $\text{K}^+$  microdomain formation, it is clear from Figure 3E that a local PsC  $\text{Ca}^{2+}$   
314 microdomain is also formed. This  $\text{Ca}^{2+}$  microdomain is formed even in the absence of an ER: the ER  
315 is widely believed to be essential for astrocyte  $\text{Ca}^{2+}$  dynamics (Verkhratsky et al., 2012). The  
316 microdomain of  $\text{Ca}^{2+}$  is caused by the reversal of the NCX causing  $\text{Ca}^{2+}$  influx in exchange for  
317 astrocytic  $\text{Na}^+$  efflux. This can be seen in Figure 4C where the only  $\text{Ca}^{2+}$  influx pathway in our model  
318 is via the NCX, the complete reversal of the NCX is due to the sudden changes in PsC  $\text{Na}^+$  during  
319 neuronal stimulation onset as the astrocyte membrane voltage is held constant.

320 Figure 4 presents the  $\text{Na}^+$  currents associated with the astrocyte. The main pathway for  $\text{Na}^+$  efflux  
321 from the astrocyte is the NKA (Figure. 4 A) while the main pathway responsible for  $\text{Na}^+$  uptake is the  
322 EAAT (Figure 4B).

323 The transient loss of  $\text{Na}^+$ , observed in the PsECS (Figure 3C), is caused by the large, transient uptake  
324 by EAAT1/2 accompanying glutamate removal following neuronal stimulation, and the EAAT  
325 remains active for approximately 30ms (Figure 4B). This large, transient, EAAT-derived  $\text{Na}^+$  flux,  
326 compared to the smaller, slower NKA and NaB activating transport rates, tips the transport  
327 uptake/release balance in favor of inward transportation and initiates the formation of a  $\text{Na}^+$   
328 microdomain. As the NKA and NaB uptake increases, due to the increased  $\text{Na}^+$  concentration in the  
329 PsC, the influx/efflux,  $\text{Na}^+$  pathways tend towards a state of equilibrium and the microdomain of  $\text{Na}^+$   
330 remains at a stable concentration. It is also noted that as  $\text{Na}^+$  increases in the PsC that the NCX works  
331 in reverse mode to remove  $\text{Na}^+$  from the PsC. Figure 4E and F show the  $\text{Na}^+$  currents along the  
332 process and between PsECS and GECS respectively. Since the efflux of  $\text{Na}^+$  via the process is  
333 several orders of magnitude smaller than the other  $\text{Na}^+$  currents this is the main driving force for the  
334 creation of the  $\text{Na}^+$  microdomain in the PsC. Moreover, as the PsECS  $\text{Na}^+$  concentration changes, the

335 leak current ( $I_{NaECSL}$ ) between the PsECS and GECS attempts to maintain the  $Na^+$  levels in the  
336 PsECS.

337 Figure 5 describes the astrocytic  $Ca^{2+}$  currents; again, it can be seen that during neural stimulation,  
338 the influx of  $Na^+$  cause the NCX to work in reverse mode and therefore there is an influx of  $Ca^{2+}$   
339 (Figure 5A). The only efflux pathway for  $Ca^{2+}$  considered within the model is via the thin astrocyte  
340 process, which is also governed by the well-hopping mechanism described in (Breslin et al., 2018)  
341 (Figure 5B), Therefore the efflux pathway is much less dominant than the influx pathway which  
342 results in a microdomain of  $Ca^{2+}$  forming, as seen in Figure 3E.

343 The  $K^+$  currents behave in a similar manner as previously described in Breslin et al. 2018. NKA and  
344 Kir represent the dominant fluxes of  $K^+$  across the astrocytic membrane; NKA is purely responsible  
345 for  $K^+$  influx whereas Kir is in a constant state of transition between forward ( $K^+$  influx) and reverse  
346 ( $K^+$  efflux) mode during neuronal stimulus (see Figure 6). When neural stimulus ceases, the voltage-  
347 dependent reversal of Kir, along with efflux of  $K^+$  via the  $K^+$  background channel, brings the PsC  
348 levels of  $K^+$  back to the initial resting state.

#### 349 4 Discussion

350 Intracellular ionic signaling represents the substrate for glial excitability (Verkhratsky and  
351 Nedergaard, 2018). These intracellular signaling events are mediated through spatially and  
352 temporally organized fluctuations in the concentration of major ions; there is firm evidence for  
353 physiologically relevant  $Ca^{2+}$  and  $Na^+$  signaling (Rose and Verkhratsky, 2016, Verkhratsky and  
354 Nedergaard, 2018; Verkhratsky et al., 2019),  $Cl^-$  signaling (Wilson and Mongin, 2018; Verkhratsky  
355 et al., 2019) and  $K^+$  signaling (Olsen et al., 2015; Rimmele and Chatton, 2014). The formation of  
356 localized concentration microdomains is critical for ionic signaling spatial fidelity; while the  
357 mechanisms underlying formation of these microdomains remain under debate, a new hypothesis has  
358 already been proposed (Breslin et al., 2018). In the CNS, most excitatory synapses are tightly  
359 enwrapped by perisynaptic astroglial processes forming the synaptic cradle (Reichenbach et al.,  
360 2010; Verkhratsky and Nedergaard, 2014). This structure provides homeostatic control of the  
361 synaptic cleft and therefore requires the ability to generate relevant signals in response to neuronal  
362 activity. The perisynaptic astroglial compartment is devoid of the ER (Reichenbach et al., 2010),  
363 hence excluding the metabotropic pathway for generation of local  $Ca^{2+}$  microdomains. Astrocytes  
364 express glutamatergic ionotropic receptors, although the receptor-mediated current density is rather  
365 low (with whole cell currents rarely exceeding 10 – 100 pA), thus limiting the ionic influx  
366 (Verkhratsky and Burnstock, 2014; Rusakov, 2015). Astroglial perisynaptic membranes also express  
367 high densities of glutamate transporters (EAAT1/2), which couple glutamate transport with  
368 substantial  $Na^+$  influx (Kirischuk et al., 2007; Langer and Rose, 2009). These transporters are co-  
369 localized with NCX (Minelli et al., 2007), which couple  $Na^+$  and  $Ca^{2+}$  fluxes in opposite directions.  
370 Here we applied the reduced model of PsC to test the hypothesis that glutamate transporters and  
371 NCX working together are sufficient to create local  $Ca^{2+}$  microdomains in astroglial perisynaptic  
372 cradles.

373 Our model demonstrates that stimulation of astrocytes with glutamate, mimicking neuronal activity,  
374 generates substantial  $Na^+$  influx, which forms local microdomains due to the previously suggested  
375 mechanism of ion retention, where cation retention in wells dominates over conventional  
376 electrochemical diffusion (Breslin et al., 2018; Wade et al. 2018). Moreover, the generation of  $Ca^{2+}$   
377 microdomains has previously been reported, albeit not at the fine process level, but the underlying  
378 ionic fluxes (and channels/transporters contributing to) have not been examined (Rusakov, 2015).

379 This computational modelling study has identified the molecular targets and their relative  
380 contributions to the formation of a  $\text{Ca}^{2+}$  microdomain in the absence of an ER region. Specifically,  
381 our model predicts that the generation of a  $\text{Na}^+$  microdomain switches the NCX into reverse mode,  
382 which is sufficient to produce relevant focal  $\text{Ca}^{2+}$  signals; while cross-disciplinary research to test  
383 this hypothesis is beyond the scope of this paper, we envisage that this work will instigate such a  
384 project.

385 In essence only the EAAT, NKA and NCX are necessary and sufficient for the development of a  
386  $\text{Ca}^{2+}$  microdomain: note that to avoid rapidly increasing  $\text{Ca}^{2+}$  concentrations within the cradle, with  
387 increasing neuronal frequency, our model would require a  $\text{Ca}^{2+}$  efflux pathway, and a likely  
388 candidate is the PMCA pump. These mechanisms, which do not depend on intracellular sources for  
389  $\text{Ca}^{2+}$ , directly link neuronal activity and glutamate release to the formation of  $\text{Na}^+$  and  $\text{Ca}^{2+}$   
390 microdomains in the perisynaptic astroglial processes, instrumental for generation of astroglial  
391 homeostatic response, which is critical for maintenance of synaptic transmission.

### 392 **5 Conflict of Interest**

393 *The authors declare that the research was conducted in the absence of any commercial or financial*  
394 *relationships that could be construed as a potential conflict of interest.*

### 395 **6 Author Contributions**

396 JW, LM, AV, MD, SH, HVZ, BF and AB contributed to the conception and design of the study. JW  
397 and KB developed the software for simulations. JW and BF created all graphics and data  
398 visualizations. JW, KB, LM, AV, MD, AB wrote the first draft of the manuscript. All authors  
399 contributed to manuscript revision, read and approved the submitted version.

### 400 **7 Funding**

401 The authors received no specific funding for this work.

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508 **9 Data Availability Statement**

509 No datasets were generated or analyzed for this study.

510 **10 Figure Legends**

511 Figure 1. Model Morphology. The model consists of a single synapse enwrapped by a single astrocyte.  
 512 In total there are six compartments, 1) Global Extracellular Space (GECS), 2) Synapse, 3) Perisynaptic  
 513 Extracellular Space (PsECS), 4) Perisynaptic Cradle, 5) Astrocyte Process, and 6) Astrocyte Soma.  
 514 Each compartment is modelled as a cylindrical structure except the GECS and soma, which are deemed  
 515 dimensionless because ionic concentrations remain constant within these compartments (Breslin et al.,  
 516 2018).

517 Figure 2. Ion transport machinery of the perisynaptic cradle and synapse. The model consists of 11  
 518 ionic transports. The synapse contains 5 ionic transports for Na<sup>+</sup> and K<sup>+</sup>; NKA is the sodium/potassium  
 519 pump which extrudes 3 Na<sup>+</sup> ions for every 2 K<sup>+</sup> ions it imports, Na<sub>neu</sub> and K<sub>neu</sub> which are the voltage  
 520 gated sodium and potassium channels of the Hodgkin and Huxley model and Na<sub>B</sub> and K<sub>B</sub> which are

521 lump models of all the other  $\text{Na}^+$  and  $\text{K}^+$  channels. The astrocyte contains 6 channels on the inside  
 522 surface area (facing the synapse) of the perisynaptic cradle;  $\text{Kir}$  is an inward rectifying  $\text{K}^+$  channel,  
 523  $\text{NKA}$  is a sodium potassium pump similar to that found on the synapse,  $\text{EAAT1/2}$  represents the EAAT  
 524 glutamate transport which is sensitive to glutamate in the synaptic cleft. This transport cotransports 3  
 525  $\text{Na}^+$  ions with every glutamate ion and counter transports 1  $\text{K}^+$  ion. Since this work does not consider  
 526 changes in synaptic and perisynaptic glutamate changes the glutamate ion is not shown in the figure.  
 527 Moreover, the EAAT dependency on  $\text{H}^+$  is also ignored. The  $\text{NCX}$  represents the sodium/calcium  
 528 exchanger which exchanges 3  $\text{Na}^+$  ions for every 1  $\text{Ca}^{2+}$  ion across the membrane. At resting conditions  
 529 there is a very small exchange in the forward direction as noted in the diagram. This can be reversed  
 530 under physiological increase of perisynaptic  $\text{Na}^+$  concentration. Finally, the  $\text{Na}_B$  and  $\text{K}_B$  are lumped  
 531 models of all other  $\text{Na}^+$  and  $\text{K}^+$  ionic transports respectively. The currents  $I_{\text{KECSL}}$  and  $I_{\text{NaECSL}}$  represent  
 532 leak currents of  $\text{K}^+$  and  $\text{Na}^+$  respectively from the Perisynaptic Extracellular Space (ECS) to the global  
 533 ECS. It is worth nothing that the currents  $I_{\text{KPF}}$ ,  $I_{\text{NaPF}}$  and  $I_{\text{CaPF}}$  represent the currents of the 3 ions under  
 534 consideration from the perisynaptic cradle along the process. As described in Breslin et al. (2018),  
 535 these currents model the hypothesized well-hopping mechanism which severely restricts current flow  
 536 of these ions along thin processes. This results in the trapping of these ions in the perisynaptic cradle  
 537 as they are imported across the membrane. Since our model assumes a well-mixed compartment, we  
 538 consider these trapped ions as a microdomain formation in the perisynaptic cradle.

539 Figure 3. Astrocyte PsC ion concentrations. (A)  $[\text{K}^+]_{\text{PsECS}}$  transient. (B)  $[\text{K}^+]_{\text{PsC}}$  transient. (C)  $[\text{Na}^+]_{\text{PsECS}}$   
 540 transient. (D)  $[\text{Na}^+]_{\text{PsC}}$  transient. (E)  $[\text{Ca}^{2+}]_{\text{PsC}}$  transient. During periods of neuronal activity, increased  
 541  $[\text{K}^+]_{\text{PsECS}}$  is cleared by the astrocyte leading to a microdomain of  $\text{K}^+$  in the PsC. Moreover, due to the  
 542 influx of  $\text{Na}^+$  predominantly via EAAT channels there is an increase in  $[\text{Na}^+]_{\text{PsC}}$ , resulting in the  
 543 reversal of the NCX and  $\text{Ca}^{2+}$  microdomain formation. The inserts within (A-E) show the fast dynamics  
 544 of the ionic concentration changes within the PsC in response to the neurone activity.

545 Figure 4. Astrocyte  $\text{Na}^+$  currents. (A)  $\text{Na}^+$  NKA current. This current is the main pathway for  $\text{Na}^+$   
 546 efflux from the astrocyte is the NKA while the main pathway for sodium efflux is  $\text{Na}^+$  current through  
 547 the EAAT (see (B)) (C)  $\text{Na}^+$  NCX current. It can be seen that during periods of neurone stimulation,  
 548 the NCX reverses which results in the efflux of  $\text{Na}^+$  from the PsC in response to the increased  $\text{Na}^+$   
 549 uptake via the EAAT. (D) Background  $\text{Na}^+$  current. This represents the uptake of  $\text{Na}^+$  through a lumped  
 550 model of all other  $\text{Na}^+$  channel. The current can be seen to slow down during periods of PsC  $\text{Na}^+$   
 551 increase during neurone stimulus. (E)  $\text{Na}^+$  current along the process. The current along the process is  
 552 governed by the well-hopping mechanism described by Breslin et al. (2018) which severely restricts  
 553 the flow of  $\text{Na}^+$  from the PsC to the soma. This restriction results in a microdomain of  $\text{Na}^+$  forming in  
 554 the PsC compartment as the  $\text{Na}^+$  cannot simply diffuse along the process. (F)  $\text{Na}^+$  current due to the  
 555 leakage of  $\text{Na}^+$  from the PsECS to the GECS which is purely diffusive. The inserts within (A-F) show  
 556 the fast dynamics of the currents in response to neurone activity.

557 Figure 5. Astrocyte  $\text{Ca}^{2+}$  currents. (A) Astrocyte  $\text{Ca}^{2+}$  NCX current. During periods of neural  
 558 stimulation an influx of  $\text{Na}^+$  via the EAAT1/2 cotransporter results in the reversal of the NCX and thus  
 559 there is an influx of  $\text{Ca}^{2+}$  into the PsC. (B)  $\text{Ca}^{2+}$  current along the process. The only efflux pathway  
 560 within the model is along the process. The current along the process is governed by the well-hopping  
 561 mechanism described by Breslin et al. (2018) which severely restricts the flow of  $\text{Ca}^{2+}$  from the PsC to  
 562 the soma. This restriction results in a microdomain of  $\text{Ca}^{2+}$  forming in the PsC compartment. The  
 563 inserts within (A-B) show the fast dynamics of the currents in response to neurone activity.

564 Figure 6. Astrocyte  $\text{K}^+$  currents. (A)  $\text{K}^+$  Kir Current. During periods of neurone stimulation the Kir  
 565 channel will initially uptake  $\text{K}^+$  before releasing it again at a much slower rate (see insert (A)). (B)  $\text{K}^+$

566 NKA current. The rate of NKA increases with neurone stimulation due to the increase of PsECS  $K^+$ .  
 567 This current is mainly responsible for the uptake  $K^+$  in the PsC. (C)  $K^+$  current along the process. Since  
 568 the current along the process is governed by the well hopping mechanism described by Breslin et al.  
 569 (2018) it severely restricts the flow of  $K^+$  from the PsC to the soma. This restriction results in a  
 570 microdomain of  $K^+$  forming in the PsC compartment as the  $K^+$  cannot simply diffuse along the process.  
 571 Furthermore,  $K^+$  is transported across the membrane via background and EAAT  $K^+$  currents (D, E  
 572 respectively) much more slowly as these currents are mainly driven by transient  $K^+$  fluctuations in the  
 573 PsECS due to the neurone stimulation; while these currents have a higher peak magnitude, they are  
 574 much more “spike like in nature” than NKA currents (See inserts B, D, E). (F)  $K^+$  current due to the  
 575 leakage of  $K^+$  from the PsECS to the GECS which is purely diffusive.

576 **11 Tables**

577 **Table 1: Astrocyte Morphology.**

Parameter	Value	Units	Description
<b>Lengths:</b>			
<b>d<sub>IPS</sub></b>	$300 \times 10^{-9}$	m	Perisynaptic internal diameter
<b>d<sub>EPS</sub></b>	$500 \times 10^{-9}$	m	Perisynaptic external diameter
<b>r<sub>IPS</sub></b>	$150 \times 10^{-9}$	m	Perisynaptic internal radius
<b>r<sub>EPS</sub></b>	$250 \times 10^{-9}$	m	Perisynaptic external radius
<b>l<sub>PS</sub></b>	$300 \times 10^{-9}$	m	Perisynaptic length
<b>d<sub>P</sub></b>	$100 \times 10^{-9}$	m	Process diameter
<b>r<sub>P</sub></b>	$50 \times 10^{-9}$	m	Process radius
<b>l<sub>P</sub></b>	$25 \times 10^{-6}$	m	Process length
<b>d<sub>Syn</sub></b>	$270 \times 10^{-9}$	m	Synapse diameter
<b>r<sub>Syn</sub></b>	$135 \times 10^{-9}$	m	Synapse radius
<b>l<sub>syn</sub></b>	$300 \times 10^{-9}$	m	Synapse length
<b>Areas:</b>			



<b>CSA<sub>Ps</sub></b>	$3.5343 \times 10^{-14}$	m <sup>2</sup>	Perisynaptic cross sectional area
<b>SA<sub>Ps</sub></b>	$1.4137 \times 10^{-13}$	m <sup>2</sup>	Perisynaptic surface area
<b>CSA<sub>P</sub></b>	$7.854 \times 10^{-15}$	m <sup>2</sup>	Process cross sectional area
<b>SA<sub>P</sub></b>	$7.854 \times 10^{-12}$	m <sup>2</sup>	Process surface area
<b>CSA<sub>Syn</sub></b>	$2.8628 \times 10^{-14}$	m <sup>2</sup>	Synapse cross sectional area
<b>SA<sub>Syn</sub></b>	$1.2723 \times 10^{-13}$	m <sup>2</sup>	Synapse surface area
<b>SA<sub>PsECS-GECS</sub></b>	$1.5715 \times 10^{-14}$	m <sup>2</sup>	Surface area between PsECS and GECS
<b>Volumes:</b>			
<b>Vol<sub>Ps</sub></b>	$1.8850 \times 10^{-17}$	L	Perisynaptic volume
<b>Vol<sub>P</sub></b>	$1.9635 \times 10^{-16}$	L	Process volume
<b>Vol<sub>Syn</sub></b>	$8.5883 \times 10^{-16}$	L	Synapse volume
<b>Vol<sub>PsECS</sub></b>	$2.0145 \times 10^{-18}$	L	Perisynaptic ECS volume

578 **Table 2: Astrocyte Model Variables.**

<b>Variable</b>	<b>Initial Value</b>	<b>Units</b>	<b>Description</b>
<b>V<sub>A</sub></b>	-0.0807	V	Astrocyte Membrane potential
<b>[K<sup>+</sup>]<sub>PsC</sub></b>	0.1	M	K <sup>+</sup> concentration in the perisynaptic cradle
<b>[Na<sup>+</sup>]<sub>PsC</sub></b>	0.015	M	Na <sup>+</sup> concentration in the perisynaptic cradle
<b>[K<sup>+</sup>]<sub>PsECS</sub></b>	0.004	M	Perisynaptic extracellular K <sup>+</sup> concentration
<b>[Na<sup>+</sup>]<sub>PsECS</sub></b>	0.135	M	Perisynaptic extracellular Na <sup>+</sup> concentration
<b>[Glu]<sub>ECS</sub></b>	$25 \times 10^{-9}$	M	Perisynaptic extracellular Glutamate concentration

580 **Table 3: Astrocyte Model Parameters.**

<b>Parameter</b>	<b>Value</b>	<b>Units</b>	<b>Description</b>
<b>V<sub>m</sub></b>	-0.0807	V	Astrocyte resting membrane potential
<b>φ<sub>w</sub></b>	0.267	eV	Well activation energy
<b>k<sub>B</sub></b>	$1.38 \times 10^{-23}$	J/K	Boltzmann constant
<b>R</b>	8.31	J/mol/K	Gas constant
<b>T</b>	310	K	Temperature
<b>F</b>	96485	C/mol	Faraday constant
<b>Q</b>	$1.6022 \times 10^{-19}$	C	Coulomb
<b>C<sub>m</sub></b>	0.01	F/m <sup>2</sup>	Membrane capacitance
<b>g<sub>Kir</sub></b>	144	S/m <sup>2</sup>	K <sub>ir</sub> channel conductance
<b>g<sub>K</sub></b>	17.9364	S/m <sup>2</sup>	K <sup>+</sup> background transport conductance
<b>g<sub>Na</sub></b>	0.9761	S/m <sup>2</sup>	Na <sup>+</sup> background transport conductance
<b>K<sub>K</sub></b>	0.018	S/m	K <sup>+</sup> Poole-Frenkel channel constant
<b>K<sub>Na</sub></b>	0.018	S/m	Na <sup>+</sup> Poole-Frenkel channel constant
<b>PNKA<sub>max</sub></b>	$0.1 \times 10^{-5}$	mol/m <sup>2</sup>	Maximum NKA-ATPase Pump Rate
<b>K<sub>Nai</sub></b>	$10 \times 10^{-3}$	M	Na <sup>+</sup> threshold for NKA-ATPase
<b>K<sub>KE</sub></b>	$1.5 \times 10^{-3}$	M	K <sup>+</sup> threshold for NKA-ATPase
<b>z<sub>K</sub></b>	1		K <sup>+</sup> Valency

## Calcium Microdomains in Astrocyte Processes

$z_{Na}$	1		Na <sup>+</sup> Valency
$z_{Ca}$	2		Calcium Valency
$[H^+]_{PsC}$	$60 \times 10^{-9}$	M	H <sup>+</sup> Concentration in the perisynaptic cradle
$[Glu]_{PsC}$	$1.5 \times 10^{-3}$	M	Glutamate Concentration in the perisynaptic cradle
$[K^+]_{AS}$	0.1	M	K <sup>+</sup> Concentration in the astrocyte soma
$[Na^+]_{AS}$	0.015	M	Na <sup>+</sup> Concentration in the astrocyte soma
$[Ca^{2+}]_{AS}$	$100 \times 10^{-9}$	M	Ca <sup>2+</sup> Concentration in the astrocyte soma
$[H^+]_{PsECS}$	$40 \times 10^{-9}$	M	Perisynaptic extracellular H <sup>+</sup> concentration
$[Ca^{2+}]_{PsECS}$	$1.5 \times 10^{-3}$	M	Perisynaptic extracellular Ca <sup>2+</sup> concentration
$[K^+]_{GECS}$	0.004	M	Perisynaptic global ECS K <sup>+</sup> concentration
$[Na^+]_{GECS}$	0.135	M	Perisynaptic global ECS Na <sup>+</sup> concentration
$[Ca^{2+}]_{GECS}$	$1.5 \times 10^{-3}$	M	Perisynaptic global ECS Ca <sup>2+</sup> concentration
$[Ca^{2+}]_{PsC}$	$100 \times 10^{-9}$	M	Ca <sup>2+</sup> concentration in the perisynaptic cradle
$[Ca^{2+}]_{PsECS}$	$1.5 \times 10^{-3}$	M	Perisynaptic extracellular Ca <sup>2+</sup> concentration
$\epsilon_0$	$8.85 \times 10^{-12}$	F/m	Vacuum permittivity
$\epsilon_r$	0.82	F/m	Relative permittivity of brain tissue
$g_{ECS}$	3.3	S/m <sup>2</sup>	Perisynaptic ECS leak conductance
$\alpha_{EAAT}$	0.0032	A/m <sup>2</sup>	Glutamate transport fitting parameter
$\beta_{EAAT}$	28.8	mV <sup>-1</sup>	Glutamate transport fitting parameter

$r_g$	$5 \times 10^{-7}$	$M^{-1}$	Slope of glutamate uptake
$s_g$	$9 \times 10^{-6}$	M	Threshold for glutamate uptake
$\bar{I}_{NCX}$	1	A/m <sup>2</sup>	NCX Partition Parameter
$\gamma$	0.5		
$J_0$	0.06	M/s	Maximum EAAT1/2 Flux rate

581 **Table 4: Neurone Parameters.**

Parameter	Value	Units	Description
$P_{NKAmaxNeu}$	$-3.7863 \times 10^{-8}$	mol/m <sup>2</sup>	Maximum NKA-ATPase Pump Rate
$K_{NaNeu}$	$10 \times 10^{-3}$	M	Na <sup>+</sup> threshold for NKA-ATPase
$K_{KNeu}$	$1.5 \times 10^{-3}$	M	K <sup>+</sup> threshold for NKA-ATPase
$[Na^+]_{Syn}$	0.015	M	Na <sup>+</sup> concentration in the synapse
$[K^+]_{Syn}$	0.1	M	K <sup>+</sup> concentration in the synapse
$g_{KNeu}$	360	S/m <sup>2</sup>	Maximum K <sup>+</sup> channel conductance
$g_{NaNeu}$	1200	S/m <sup>2</sup>	Maximum Na <sup>+</sup> channel conductance
$g_{LNeu}$	3	S/m <sup>2</sup>	Maximum leak channel conductance
$g_{KBNeu}$	1.0522	S/m <sup>2</sup>	K <sup>+</sup> Background channel conductance
$g_{NaBNeu}$	2.3217	S/m <sup>2</sup>	Na <sup>+</sup> Background channel conductance
$E_{KNeu}$	-0.12	V	K <sup>+</sup> channel reversal potential
$E_{NaNeu}$	0.115	V	Na <sup>+</sup> channel reversal potential
$E_{LNeu}$	0.010613	V	Leak channel reversal potential

$C_m$	0.01	F/m <sup>2</sup>	Membrane capacitance
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