Neurovascular Coupling Equations for Code version 3.3

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Todo list

| | re is an inconsistency here as β_{Glu} cannot be both s^{-1} and $\mu M s^{-1}$ since Glu_{switch} is non-dimensional we use calcium (Ca ²⁺)-calmodulin complex at all? | |
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Introduction

Version 3.3 is written in Python (as opposed to Matlab) and now incorporates the ability to model somatosensory cortex stimulation as well as interneuron stimulation within the same code. In addition to version 3.1 we have added the simulation of the enzyme GABA-T which degrades GABA to succinic semialdehyde and glutamate [23?]

The single neurovascular unit (NVU) model originally developed by Farr and David [5] and later extended by Dormanns et al. [4], Dormanns et al. [3], Mathias et al. [15], Kenny et al. [9], Mathias et al. [14] and most recently by Kenny (unpublished) contains 55 ordinary differential equations (ODEs) plus a large number of algebraic variables and parameters. The equations and parameters are divided into sections corresponding to different compartments and pathways of the model. Figure 1 shows the current status of the new more simplified neuron/full model (see below for details).

The difficulty in analysing the single neuron model used by [15] to produce a viable simulation of the Berwick mouse experimental data **refrerence Berwick results here** (HbT/O/R) has encouraged us to look at both neural mass models (which essentially produce oscillations modelling α and β waves \Rightarrow EEG data) and the work of Wilson and Cowan (especially their seminal 1972 paper see Biophys Jour 12 (1-24), 1972). We concentrate here on the Wilson Cowan model (designated CW hereafter) for the time being. The derivation of the model is reproduced here for clarity and to ensure a thorough understanding of the model and its appropriateness. We define the following

- E(t) = proportion of excitatory cells firing per unit time at time t
- I(t) = proportion of inhibitory cells firing per unit time at time t

E(t)=I(t)=0 is considered the resting state and note that both E(t) and I(t) can become negative at some time.

Suppose refractory period of excitable cells is r msecs, then the proportion of cells that are in a refractory state is given by

$$\int_{t-r}^{r} E(\tau)d\tau \tag{1}$$

thus cells which are sensitive are defined as

$$1 - \int_{t-r}^{r} E(\tau)d\tau \tag{2}$$

Similarly for I(t). We define the expected proportions of either E(t) or I(t) (subpopulations) receiving at least threshold excitation per unit time as a function of the average levels of excitation within the subpopulations by $\mathfrak{S}_e(x)$ and $\mathfrak{S}_i(x)$. Suppose a distribution function \mathfrak{D} of individual thresholds for the subpopulation then

$$\mathfrak{S}(x) = \int_0^{x(t)} \mathfrak{D}(\theta) d\theta \tag{3}$$

where x(t) is the average excitation OR

$$\mathfrak{S}(x) = \int_{\frac{\theta}{x(t)}}^{\infty} \mathfrak{C}(\omega) d\omega \tag{4}$$

where $\mathfrak{C}(\omega)$ is a distributions of synapses per cell. Both definitions produce (if $\mathfrak{D}, \mathfrak{C}$ are monotonic) an $]\mathfrak{S}(x)$ which is sigmoidal in shape. We can write this as

$$\mathfrak{S}(x) = \frac{1}{1 + exp[-\gamma(x - \delta)]} \tag{5}$$

Assuming individual cells sum their inputs and the delay from stimulation is $\alpha(t)$ then the average level of excitation generated will be

$$\int_{-\infty}^{t} \alpha(t-\tau)[c_1 E(\tau) - c_2 I(\tau) + P(\tau)]d\tau$$
(6)

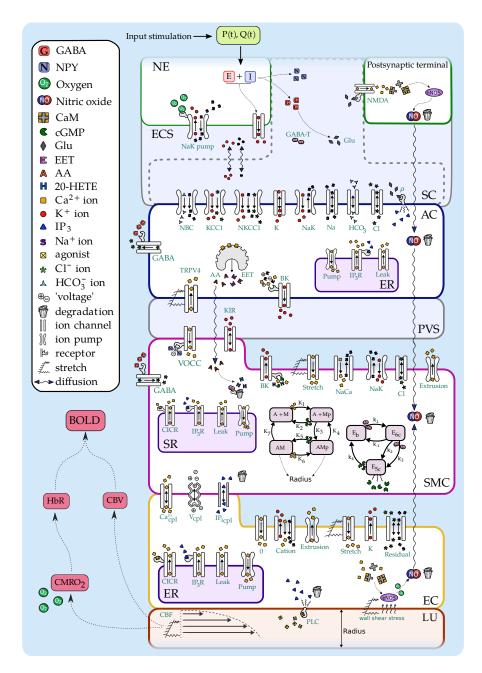


Figure 1: FULL NVU 3.3 including BOLD pathway and Wilson and Cowan neuron model

 c_1 and c_2 represent the average number of excitatory/inhibitory synapses per cell and $P(\tau)$ is the external input to the excitatory subpopulation. By assuming a "coarse grain" variable defined by

$$\bar{f}(t) = \frac{1}{s} \int_{t-s}^{t} f(\tau) d\tau \tag{7}$$

and assuming that at time $t+\tau$ the dynamics of the localised populations are given by

$$E(t+\tau) = \left[1 - \int_{t-r}^{t} E(t')dt'\right] \mathfrak{S}_{e}(x_{e}(t))$$
(8)

with x(t) defined previously, and

$$I(t+\tau) = \left[1 - \int_{t-r}^{t} I(t')dt'\right] \mathfrak{S}_{i}(x_{e}(t))$$
(9)

Using Taylor series expansion about τ =0 gives

$$\tau \frac{dE}{dt} = -\bar{E} + (1 - r\bar{E})\mathfrak{S}_e \left[kc_1\bar{E} - c_2\bar{I} + kP(t) \right]$$

$$\tau' \frac{dI}{dt} = -\bar{I} + (1 - r\bar{I})\mathfrak{S}_i \left[k'c_3\bar{E} - c_4\bar{I} + k'Q(t) \right]$$
 (10)

Q(t) is the stimulation of the inhibitory cells and

$$\int_{t-r}^{t} E(t')dt' = r\bar{E}$$

$$\int_{-\infty}^{t} \alpha(t-t')E(t')dt' = k\bar{E}$$
(11)

With some redefinitions we have the final odes

$$\tau_e \frac{dE}{dt} = -E + (k_e - r_e E)\mathfrak{S}_e \left[c_1 E - c_2 I + P(t) \right]$$

$$\tau_i \frac{dI}{dt} = -\bar{I} + (k_i - r_i I)\mathfrak{S}_i \left[c_3 E - c_4 I + Q(t) \right]$$
(12)

We should note the following . The state E=I=0 should be stable and be a steady state solution to equations for P=Q=0. \mathfrak{S}_e and \mathfrak{S}_i are transformed so that $\mathfrak{S}_e=\mathfrak{S}_i=0$. This gives

$$\mathfrak{S}(x) = \frac{1}{1 + exp[-\alpha(x - \delta)]} - \frac{1}{1 + exp(\alpha\delta)}$$
 (13)

But the maximum values of $\mathfrak{S}_{e,i}$ will be less than zero and hence k_e and k_i are the max values. as defined

$$\lim_{x \to \infty} \mathfrak{S}(x) = 1 - \frac{1}{1 + exp(\alpha \delta)} \tag{14}$$

The isoclines for the equations suggest that \exists 3 or possibly 5 steady states. In our case we wish to progress from E=I=0 when P=Q=0 to some stable state with P,Q, \neq 0 and for the system to return to E=I=0. This is not trivial (see Wilson and Cowan paper). However using the following parameters we have been able to produce a reasonable profile for E(t) and I(t)

Table 1

| Table 1 | | |
|------------|----------|--|
| Parameter | Value | |
| c_1 | 12 | |
| c_2 | 10 | |
| c_3 | 13 | |
| c_4 | 11 | |
| a_e | 1.2 | |
| θ_e | 2.8 | |
| a_i | 1.0 | |
| $	heta_i$ | 4.0 | |
| r_e | 1.0 | |
| r_i | 4.0 | |
| $	au_e$ | 10 msecs | |
| $	au_i$ | 10 msecs | |
| | | |

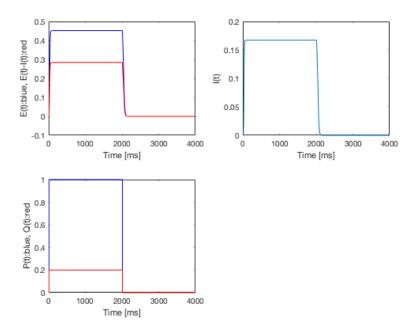


Figure 2: Upper left: E(t) and E(t)-I(t), Upper right: I(t). Lower left P(t) and Q(t)

 $k_e = \mathfrak{S}_e(10)$ and $k_i = \mathfrak{S}_i(10)$ are used to produce maximum values of the sigmoid function. Both k_e and k_i are \sharp 1. We should note here that due to the small values of the characteristic time scales τ_e and τ_1 the output for E and I are essnetially 'square pulses' if the functions P and Q are also 'square pulses' and the stimulation period is much larger than τ_e or τ_i . Figure shows the results of E(t) and I(t) for a 2 second stimulation

Note that the horizontal axes are in milliseconds. In order to develop the neuron model further we need to ensure that the more simple model replicates (adequately) the results of the more complex one. It turns out (see below the section in determining the ATP-ase pump) that in order to develop this simple model all that is needed are four outputs, viz. K^+ in the ECS, Na^+ in the soma and dendrite and the time derivative of the K^+ in the ECS.

We show the output of the full single neuron model for 2 (Figure 3) and 16 second (Figure 4) stimuli. K^+ in the synaptic cleft is determined from the derivative of the potassium (K^+) in the ECS. It is important to make sure that the resulting K^+ replicates adequately the result of Ostby [17], since we know this works!!

It can be seen that the profiles for K^+ and both sodium (Na⁺) could be modelled using a differential equation of the form

$$\frac{d\phi_i(t)}{dt} + \alpha_i \phi(t) = f_i(t) \tag{15}$$

Furthermore $f_i(t)$ could be found using the profiles of E(t) and I(t). Such that

$$f_i(t) = f_i(E(t) - I(t)) \tag{16}$$

The general solution looks like

$$\phi_i(t) = \frac{f_i(t)}{\alpha_i} \left(1 - exp(-\alpha_i t) \right) \tag{17}$$

or more succintly

$$\frac{d\phi_i(t)}{dt} + \beta\phi = \alpha\beta \tag{18}$$

with solution

$$\phi_i(t) = \alpha \left(1 - \exp(-\beta t) \right) \tag{19}$$

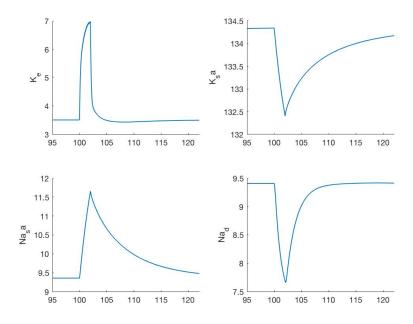


Figure 3: Time-dependent profiles of a 2 sec stimulus for K^+ in the ECS and the synapse of the neuron (top row) and Na^+ in the soma and dendrite (bottom row). Simulation from the single neuron model

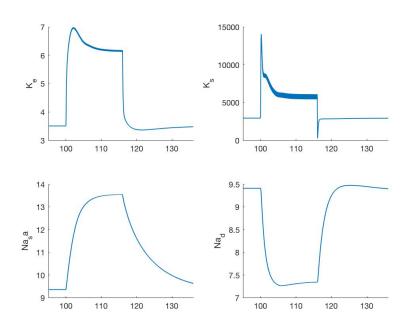


Figure 4: Time-dependent profiles of a 16 sec stimulus for K^+ in the ECS and the synapse of the neuron (top row) and Na^+ in the soma and dendrite (bottom row). Simulation from the single neuron model

We are now in a position to develop a simple neuron model. Some experiments have provided information with which we can establish adequate values for α and β . We assume that the forcing function (RHS of ode) is a 'square wave'. We further assume that the stimulus is of the order of 0.022 mA cm⁻². On this basis the functions modelling K⁺ and Na⁺

are given by

$$[K_e^+] (t) \simeq 3.5 + 2.7(1 - exp(-1.5t))$$

$$[Na_d^+] (t) \simeq 9.42 - 2.12(1 - exp(-0.7t))$$

$$[K_{sa}^+] (t) \simeq 9.37 + 4.23(1 - exp(-0.65t))$$

$$(20)$$

We now use the values of the parameters listed in Table 1 above and those in equation 20 since these replicate the original single neuron model output and find the solution to the time-dependent profiles for K_e^+, Na_{sa}^+ and Na_d^+ using various values of P(t) to signify a varying stimulus magnitude and the forcing function given by $\delta(E(t)-I(t))$ where for a stimulus of 0.022 mA cm^{-2} the forcing function is of unit magnitude. This value of 0.022 mA cm^{-2} was the original value when using the Mathias neuron model. There is a clear bifurcation at P(t) \simeq 0.4 . This is expected as Wilson and Cowan show that the steady state values traverse from stable through unstable to stable again as P(t) moves from 0.3 to higher values. As shown in Figure 5. We assume WLOG that p(t)=0.5 correpsonds to a stimulus of 22 mA cm². Varying the value of P(t) therefore enables the model to vary the stimulus input magnitude.

In the Ostby model the input to the NVU system was essentially a flux of K⁺ into the synaptic cleft. In the Mathias

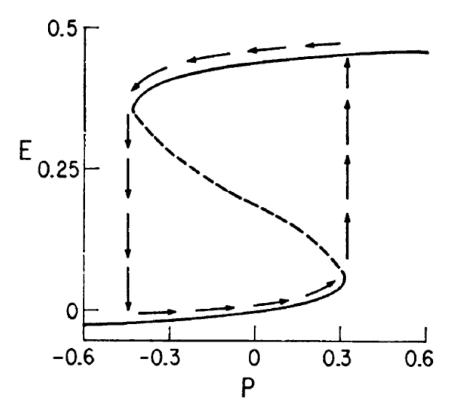


Figure 5: E vs P (taken from Wilson and Cowan paper)

neuron model this was achieved by using the derivative of the ECS K^+ multiplied by a constant to take into account the small syaptic volume compared to that of the ECS. We therefore need to determine the time dependent value of $\frac{dK_e^+}{dt}$. This is trivial as it is just a rearrangement of the ode given in equation . It is written as

$$\frac{dK_e^+}{dt} = -\beta K_e^+ + \alpha\beta \tag{21}$$

Figure 6 shows the derivative $\frac{dK_e^+}{dt}$ as a function of time. This should be compared to the Ostby input as shown in Figure 7. Note that the time scales are different but the essential form is similar. And the integrals of the positive value of $\frac{dK_e^+}{dt}$ is equal to the negative value of $\frac{dK_e^+}{dt}$ as is the case with Ostby.

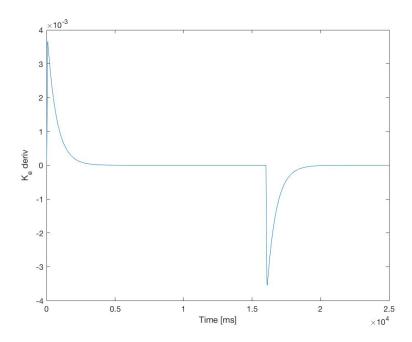


Figure 6: $\frac{dK_e^+}{dt}$ as a function of time using a stimulation of 16 seconds. re-arrangement of basic K^+ ode, see equation .

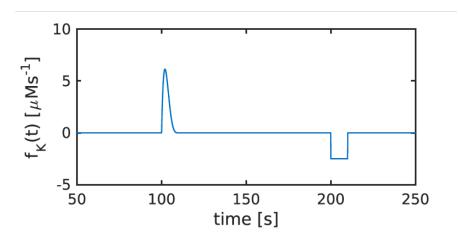


Figure 7: Ostby input (replicating a neuron activation) over 200 seconds, compare with Figure 6

1 Results from Wilson and Cowan model

We are now in a position to compare the new neuron model developed from Wilson and Cowan with experimental results from [30]. Firstly Figure 8 shows the excitatory and inhibitory inputs E(t) and I(t) resp. All figures show the old model in red and new model in blue. Figure 9 shows the flux of K^+ into the synaptic cleft for the old neuron model [15] (red) compared with the new model (blue). Figure 10 shows the K^+ in the ECS for both models (colours as noted above). Figure 1 shows the K^+ in the synaptic cleft. Figure 12 shows the resulting radius (red: old model, blue: new model) Here we see that the radius now has the correct 'plateau' profile. The rate of increase at the start of the atimulation is the same as the old model. A comparison can be done with the results of [30]. Figure 13 provides the time-dependent profiles of Δ CBF for both Zheng and the Wilson/Cowan model. The value of the K^+ in the ECS has been adjusted to make the maximum height of the Δ CBF profile fit to the experiment. We note a number of issues

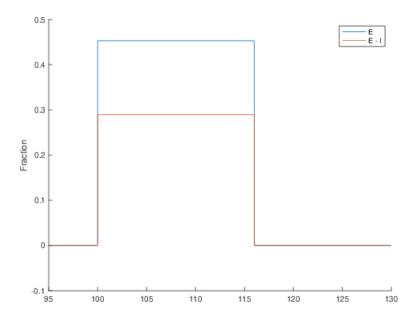


Figure 8: E(t) and I(t) for a 16 second stimulation

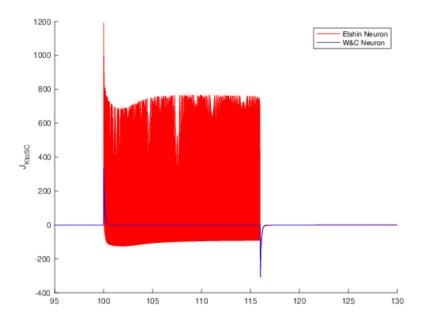


Figure 9: Flux of K+into the synaptic cleft (old model red, new model blue)

- \bullet The initial rate of increase of $\triangle CBF$ is lower than the Zheng result from the experiment
- \bullet The decay of the ΔCBF is also too long.
- the model profile does not exhibit a 'dip' in the middle of the stimulus.

The Δ CBF rates in increase and decay are functions of the radius profile. We are able to change the initial rate and final decay by altering the smooth muscle mechanics. This will be done at a later stage. The 'dip' however can be

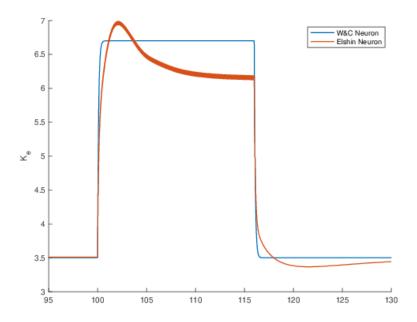


Figure 10: K+ in the ECS (red: old model. Blue: new model)

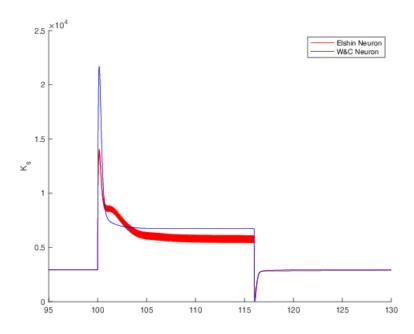


Figure 11: K+(t) in the synaptic cleft (red: old model, blue: new model)

simulated by investigating the profile of the neuronal input. We show below how this may be achieved.

1.1 Models of Murine whisker stimulation

Chapter 11 of the book Mathematical Foundations of Neuroscience by Ermentrout and Terman provides a model of the excitatory and inhibitory neurons resulting from thalamic input as a way of describing the stimulation of the somoato-

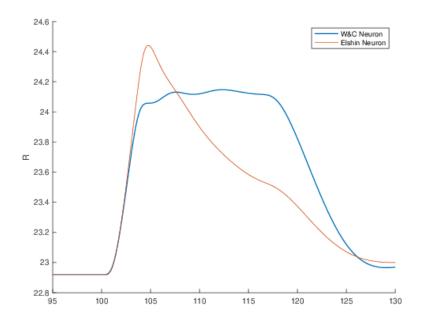


Figure 12: Radius R(t) for a 16 second stimulus (red: old model, blue: new model)

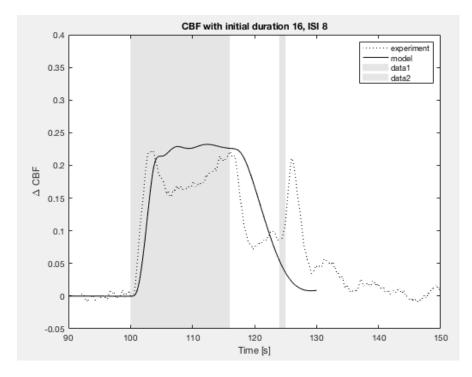


Figure 13: Comparison of old neuron model with new model for ΔCBF

sensory cortex via the whisker pad. Within the barrel cortex of the rat there exist strong recurrent excitatory and inhibitory networks. These connected networks receive stimulus from the thalamus (T(t)). Each neuronal type inhibits the other and themselves.

The experiments seem to show that it is not the peak value of the thalamic input but the time rate of change of the

stimulation. The chapter provides an example whose equations are given below.

$$\tau_{e} \frac{dE}{dt} = -E + F_{e} \left(w_{ee} E - w_{ie} + w_{te} T(t) \right)$$

$$\tau_{i} \frac{dI}{dt} = -I + F_{i} \left(w_{ei} E - w_{ii} + w_{ti} T(t) \right)$$
(22)

$$\tau_i \frac{dI}{dt} = -I + F_i \left(w_{ei} E - w_{ii} + w_{ti} T(t) \right) \tag{23}$$

Table 1 (parameters for barrel cortex model)

| w_{ee} | 42 | |
|----------|-------|---------|
| w_{ie} | 24 | |
| ei | 42 | |
| w_{ii} | 18 | Along v |
| $	au_e$ | 5 | Along v |
| $	au_i$ | 15 | |
| w_{te} | 53.43 | |
| w_{ti} | 68.4 | |

vith the gain functions written as

$$F_e(x) = \frac{5.12}{\left(1 + exp\left(-\frac{(x-15)}{4 \cdot 16}\right)\right)}$$
(24)

$$F_e(x) = \frac{5.12}{\left(1 + exp\left(-\frac{(x-15)}{4.16}\right)\right)}$$

$$F_i(x) = \frac{11.61}{\left(1 + exp\left(-\frac{(x-15)}{3.94}\right)\right)}$$
(24)

We should note finally that Ermentrout and Terman show that the factor $(1 - r_e E)$ makes little difference to the output so it is assumed that both r_e and $r_i = 0$.

To round out the model a thalamic input is needed. Since the rate of increase of input is the signifying factor in whether the firing rate increases a function is needed which simulates a relatively constant max value but a variation in the time taken to reach that value.

Using the thalamic model parameter values and an input (triangular oscillations) similar to that derived from an experiment by Berwick et al Figure 14 shows a comparison between |E(t) - I(t)|, K_e , Na_{sa} as a function of T (thalamic input) as can be seen the averaged value follows closely that of te oscillatory response. We therefore conclude that using averaged values is best and saves on compute time.

Interneuron Modeling

As part of a collaboration with the Berwick group in Sheffield UK it was decided to create a model of an inhibitory interneuron with γ-aminobutyric acid (GABA) and neuropeptide Y (NPY) to induce dilation when the neuronal vascular response is mediated by GABA but provides constriction with NPY. This is clearly a balance of two competeing pathways. To produce a viable model it was agreed that the model should be compared with experimental data from Berwick et al.

An interneuron is a specialised type of neuron whose primary role is to form a connection between other types of neurons. A large majority of interneurons of the central nervous system are of the inhibitory type. In contrast to excitatory neurons, inhibitory cortical interneurons characteristically release the neuro-transmitter GABA [8].

GABA is the main inhibitory neurotransmitter in the mammalian cortex [19]. Every third chemical synapse in the brain uses neurotransmitter GABA as an integral part of the neurotransmission process [20]. In $GABA_A$ receptors, binding

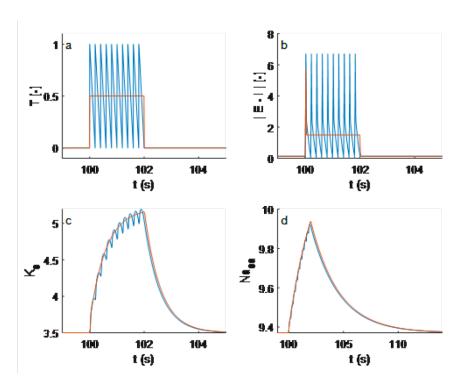


Figure 14: Comparison of WC model resulst for oscillatory thalamic input compared to an averaged value.

of GABA molecules in the extracellular part of the receptor triggers the opening of a chlorine (Cl⁻) ion selective pore, where the channel has a reversal potential of about -75 mV in neurons [?]. These receptors are also found on astrocytes [20] and smooth muscle [16].

Losi et al. [13] found that $GABA_A$ receptors on astrocytes are similar in many, though not all, aspects to those expressed by neurons. One difference is that activation of astrocytic $GABA_A$ receptors leads to a depolarising current in mature astrocytes, as opposed to mature neurons, due to the Na⁺/K⁺/Cl⁻ cotransporter (NKCC1) expression and activity that maintains a larger intracellular Cl⁻ concentration in astrocytes.

Anenberg et al. [2] found that optogenetic stimulation of GABAergic neurons (inhibitory interneurons) can lead to a net increase in cerebral blood flow (CBF). Whereas Uhlirova et al. [25] found that blood vessels in the brain will only constrict in response to inhibitory nerve cells. They identified NPY as a signal that triggers the constriction of the blood vessels. This signaling molecule is majorly expressed in interneurons and is released by a specific subtype of inhibitory nerve cell. NPY binds to a receptor protein on the SMCs and potentiates vasoconstriction by promoting Ca²⁺ entry into SMCs via voltage operated Ca²⁺ channels (VOCCs) [27, 29].

The neurotransmitter GABA (released in conjunction with NPY in inhibitory interneurons) mainly degrades to succinic semialdehyde and glutamate [23?]. This degradation is catalysed by the enzyme γ -aminobutyric acid transaminase (GABA-T), and increased GABA-T activity in the brain diminishes the GABA concentration [7]. Multiple experiments have shown that NO suppresses the activity of GABA-T and increases GABA, whereas a lack of NO (achieved through NO inhibitors such as N(ω)-nitro-L-arginine methyl ester (L-NAME)) increases GABA-T activity and reduces GABA concentration [7, 18, 24, 26].

In the experiments of Paul and Jayakumar [18] they found that the glutamate concentration stayed constant after administering L-NAME (and hence decreasing the NO concentration). They also found that the GABA concentration halved when L-NAME was introduced. The increase in GABA-T activity due to L-NAME is harder to determine due to the large variation in experiments, with increases ranging from 100% to 800% [26].

2.1 GABA and NPY pathways

A schematic of these two pathways is shown in Figure 15. The NPY pathway involves only the SMC. An increase in NPY concentration causes the VOCCs to open, allowing an influx of Ca²⁺ into the SMC leading to vasoconstriction [27, 29]. In contrast, the GABA pathway involves multiple cells and the primary effect is through GABA dependent Cl⁻ channels.

On the neuron, GABA opens Cl⁻ channels allowing an influx of Cl⁻ that hyperpolarises the neuron and inhibits any action potentials [22]. GABA also opens Cl⁻ channels on the astrocyte, allowing an efflux of chlorine from the astrocyte (the Cl⁻ travels out of the cell rather than inwards due to the NKCC1 cotransporter expression and activity that maintains a larger intracellular Cl⁻ concentration in astrocytes). Finally GABA opens Cl⁻ channels on the SMC, allowing an influx of Cl⁻ that hyperpolarises the SMC and closes the VOCCs; hence the Ca²⁺ concentration in the SMC decreases causing vasodilation [22].

GABA also degrades to glutamate in a reaction catalysed by the enzyme GABA-T (where GABA-T activity is inhibited by NO) [?]. Glutamate increases neuronal NO synthase (nNOS) production in the neuron which increases NO concentration and leads to vasodilation (note that the dynamics are slow and NO only has a minor effect over short time periods). Glutamate also leads to an increase in astrocytic inositol trisphosphate (IP₃) concentration via receptors on the astrocytic process. IP₃ causes a release of Ca²⁺ from internal stores leading to an increase in astrocytic Ca²⁺ and arachidonic acid (AA). The AA diffuses to the SMC and increases 20- hydroxyeicosatetraenoic acid (20-HETE) concentration which closes the big potassium (BK) channel. This leads to cell depolarisation, opening the VOCCs and leading to vasoconstriction.

2.2 Interneuron Model

In order to model these two pathways we use the NVU model as a base. A schematic of the previous excitatory neuron model is shown in Figure 1 whereas the new interneuron model is shown in Figure 16. The GABA and NPY concentrations are time dependent input functions that can be switched on or off as needed, where during stimulation the concentrations go from zero to some maximal value. The GABA concentration is determined with an ODE that is comprised of a degradation term (proportional to GABA-T activity) and a time dependent input simulating GABA release from the neuron during neuronal stimulation. The GABA-T activity is modelled as a sigmoid function that decreases with neuronal NO concentration.

The glutamate concentration is determined with an ODE and increases either due to vesicle release from an excitatory neuron or due to the degradation of GABA in an inhibitory neuron.

Two GABA dependent Cl^- channels are added, with one on the astrocyte and one on the SMC. These channels have a reversal potential of -75 mV and a channel conductance dependent on GABA; when GABA = 0, conductance = 0 and when GABA is at some maximal value, the conductance is also some maximal value G_{GABA} . At the moment this maximum is taken as $G_{GABA} = 0.3 \times G_{Cl,i}$ where $G_{Cl,i}$ is the conductance of the SMC Cl^- leak channel (value chosen to fit with Berwick experimental data). There is no GABA dependent Cl^- channel on the neuron as we don't have a neuronal membrane potential or Cl^- equation. Neuronal inhibition is simply modelled by a lack of neuronal stimulation.

The conductance of the VOCC in the SMC is multiplied by a NPY dependent function such that when NPY= 0, the VOCC conductance g_{VOCC} is equal to the regular conductance $G_{Ca,i}$ of the VOCC, and when NPY is at some maximal value g_{VOCC} is equal to $1.05 \times G_{Ca,i}$ (value chosen to fit with Berwick experimental data).

Extra reference: Adamchik et al. [1] formed a 2 variable model of a homogeneous neural population describing the behaviour of interneurons with tonic GABA conductance under the action of non-stationary ambient GABA (however

Interneuron model: GABA and NPY pathways

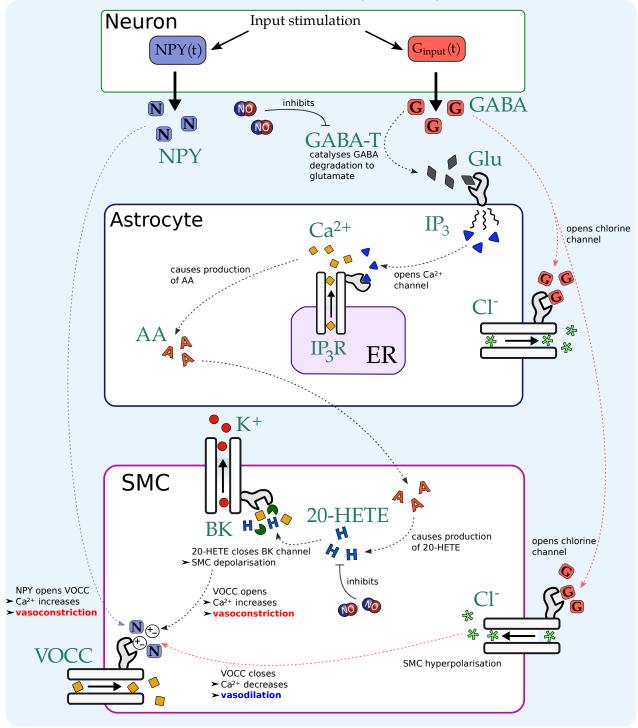


Figure 15: Schematic diagram of the GABA and NPY pathways in an interneuron model. Neuronal stimulation causes a release of NPY and GABA from the neuron. NPY opens the VOCC causing vasoconstriction. GABA opens Cl⁻ channels on the SMC causing cell hyperpolarisation and hence closes the VOCC leading to vasodilation. GABA also degrades to glutamate leading to an increase in astrocytic Ca²⁺ and AA. The AA diffuses to the SMC and increases 20-HETE concentration which closes the BK channel. This leads to cell depolarisation, opening the VOCC and leading to vasoconstriction.

they focused on the emergence of relaxation oscillations).

2.3 Interneuron model equations and parameters

The equations are given below and the parameters can be found in Table 1.

2.4 GABA

The model parameters used in the following equations are chosen to fit with the experimental findings of Paul and Jayakumar [18] and Vega Rasgado et al. [26] so that when L-NAME is administered we obtain a decrease of 50% in GABA concentration and increase of 100% in GABA-T activity.

The GABA concentration is nondimensionalised with respect to some maximal value so that at rest the concentration is zero and during regular neuronal stimulation the nondimensional concentration $GABA_N$ is 1. The ODE for $GABA_N$ is given by

$$\frac{dGABA_N}{dt} = -\kappa_{GABA} GABA_N + G_{input}(t)$$
 (26)

where the first term on the RHS models GABA degradation and $G_{input}(t)$ is a time dependent input function simulating an input of GABA during neuronal stimulation modelled via a heaviside function:

$$G_{input}(t) = \begin{cases} 1 & t_0 < t < t_0 + \Delta t \\ 0 & \text{otherwise} \end{cases}$$
 (27)

where t_0 is the beginning of stimulation and Δt is the length of stimulation. The rate at which GABA degrades is given by κ_{GABA} and is proportional to the GABA-T activity $(G_{T,act})$, where GABA-T is the enzyme that catalyses the degradation reaction:

$$\kappa_{GABA} = p_1 G_{T.act} \tag{28}$$

where $p_1 = 1 \text{ s}^{-1}$ is a scaling parameter to obtain the correct units (s⁻¹). The GABA-T activity is inhibited via NO and modelled with sigmoidal function:

$$G_{T,act} = G_{T,min} + (G_{T,max} - G_{T,min}) \frac{1}{1 + \left[\left(\frac{NO_i - NO_{rest}}{R_{NO}} \right) \right]}$$
(29)

where $G_{T,min}=1$ is the minimum activity (i.e. when NO_n is at normal resting value, $NO_{rest}\approx 0.02047~\mu\mathrm{M}$ and $R_{NO}\approx 0.02\mu\mathrm{M}$), $G_{T,max}=2$ is the maximum activity (i.e. when $NO_n\approx 0~\mu\mathrm{M}$), and $p_2=200~\mu\mathrm{M}^{-1}$ is a scaling parameter. Note that $G_{T,act}$ is nondimensional and normalised w.r.t. some minimal value.

The glutamate concentration can either be increased due to neuronal stimulation as in an excitatory neuron (modelled as a glutamate release when extracellular K^+ concentration K_e is above some threshold) or increased from the degradation of GABA in an inhibitory interneuron (as GABA degrades to succinic semialdehyde and glutamate when catalysed by GABA-T). The nondimensional glutamate concentration Glu (normalised w.r.t. some maximal value) is modelled by the following equation:

$$\frac{dGlu}{dt} = f(K_e) + g(GABA) - p_1 Glu \tag{30}$$

where $f(K_e)$ is a function that models the neuronal release of glutamate and f(GABA) is a function that models the production of glutamate from the degradation of GABA. These two functions are given by

$$f(K_e) = \frac{p_1}{2} \left(1 + \tanh \left[\frac{K_e - Ke_{switch}}{Glu_{slope}} \right] \right)$$
 (31)

Interneuron model Input stimulation \longrightarrow NPY(t) G GABA NE NPY NPY Oxygen Mitric oxide CaM **€** cGMP Glu E EET AA **ℍ** 20-HETE Ca²⁺ ion K^{+} ion ▲ IP₃ **100** Na⁺ ion agonist Cl⁻ ion △ HCO₃ ion ⊕⊝ 'voltage' degradation ion channel PVS ion pump receptor *** ₹ stretch **←** diffusion BOLD SR SMC HbR CBV ER LU

Figure 16: Interneuron model (compare with Figure 1)

Radius

where $Ke_{switch} = 5$ mM is the concentration above which glutamate is released from the neuron and $Glu_{slope} = 0.1$ mM is the slope of the sigmoidal (both parameters taken from the previous version of the model [14]), and

$$g(GABA) = \kappa_{GABA} GABA \tag{32}$$

where κ_{GABA} is the degradation rate of GABA (see Equation 28). At rest both $f(K_e)$ and f(GABA) are zero and so the glutamate concentration tends to zero.

The GABA dependent Cl⁻ channel fluxes on the astrocyte and SMC are added to the differential equations for v_k and v_i respectively and are given by

$$J_{GABA,k} = g_{GABA}(v_k - E_{GABA}) \tag{33}$$

$$J_{GABA,i} = g_{GABA}(v_i - E_{GABA}) \tag{34}$$

where v_k is the astrocytic membrane potential in mV, v_i is the SMC membrane potential in mV, $E_{GABA} = -75$ mV is the reversal potential of the Cl⁻ channels, and the GABA dependent conductance of the channels is given by

$$g_{GABA} = \frac{G_{GABA}}{2} \left(1 + \tanh\left(\frac{GABA_N - g_{mid}}{g_{slope}}\right) \right)$$
 (35)

where $g_{mid}=0.8$ is the midpoint of the sigmoidal, g_{slope} is the slope of the sigmoidal (both model estimates), and G_{GABA} is the maximal conductance of the channel given by $0.3\times G_{Cl,i}$ where $G_{Cl,i}=1.34\times 10^{-6}~\mu\mathrm{M}~\mathrm{mV}^{-1}~\mathrm{ms}^{-1}$ is the conductance of the SMC Cl⁻ leak channel taken from the NVU model. This is a model estimate and can be changed by fitting to data. Hence when $GABA_N=0$, $g_{GABA}=0$ and when $GABA_N=1$, $g_{GABA}=G_{GABA}$.

2.5 NPY

NPY is given by a time dependent input function (heaviside function):

$$NPY_N(t) = \begin{cases} 1 & t_0 < t < t_0 + \Delta t \\ 0 & \text{otherwise} \end{cases}$$
 (36)

where t_0 is the beginning of stimulation and Δt is the length of stimulation. Note that the NPY concentration is nondimensionalised with respect to its maximal value (currently unknown), i.e. $NPY_N = NPY/NPY_{max}$ so that during the stimulation period $NPY_N = NPY_{max}/NPY_{max} = 1$.

Finally the flux of Ca²⁺ through the VOCC channel on the SMC is given by

$$J_{VOCC,i} = g_{VOCC} \frac{v_i - v_{Ca1}}{1 + \exp\left(\frac{-(v_i - v_{Ca2})}{R_C a}\right)}$$
(37)

where $v_{Ca1} = 100 \text{ mV}$ is the reversal potential, $v_{Ca2} = -24 \text{ mV}$ is the half-point of the VOCC activation sigmoidal, and $R_{Ca} = 8.5 \text{ mV}$ is the maximum slope of the sigmoidal (all taken from the NVU model). The NPY conductance is given by

$$g_{VOCC} = G_{Ca,i} \left(1 + \frac{N_{inc}}{2} \tanh \left(\frac{NPY_N - N_{mid}}{N_{slone}} \right) \right)$$
 (38)

where $G_{Ca,i}=1.29\times 10^{-6}~\mu\mathrm{M}~\mathrm{mV}^{-1}~\mathrm{ms}^{-1}$ is the base conductance of the VOCC (taken from the NVU model), $N_{inc}=0.05$ is the proportional increase of the conductance from baseline due to NPY (i.e. 0.05 means a 5% increase from $G_{Ca,i}$ when NPY is released, model estimate), $N_{mid}=0.6$ is the midpoint of the sigmoidal, and $N_{slope}=0.1$ is the slope of the sigmoidal (both model estimates). Hence when $NPY_N=0$, $g_{VOCC}=G_{Ca,i}$ and when $NPY_N=1$, $g_{VOCC}=1.05\times G_{Ca,i}$.

| Parameter | Description | Value | Reference |
|-------------|---|--|-----------|
| p_1 | Scaling parameter for unit conversion | $1 \mathrm{s}^{-1}$ | |
| $G_{T,min}$ | Minimum GABA-T activity | 1 | [26] |
| $G_{T,max}$ | Maximum GABA-T activity | 2 | [26] |
| E_{GABA} | Reversal potential of the GABA dependent Cl ⁻ channels | $-75~\mathrm{mV}$ | [?] |
| G_{GABA} | Conductance of GABA dependent Cl ⁻ channel | $4.02 	imes 10^{-7} \; \mu \mathrm{M \; mV^{-1} \; ms^{-1}}$ | [11] |
| g_{mid} | Midpoint of the g_{GABA} sigmoidal | 0.8 | M. E. |
| g_{slope} | Slope of the g_{GABA} sigmoidal | 0.1 | M. E. |
| N_{inc} | proportional increase in VOCC conductance due to NPY | 0.05 | M. E. |
| N_{mid} | Midpoint of the g_{VOCC} sigmoidal | 0.6 | M. E. |
| N_{slope} | Slope of the g_{VOCC} sigmoidal | 0.1 | M. E. |

Table 1: New parameters of the GABA and NPY model.

We should note that version 3.3 now includes the GABA and NPY pathways as part of the somatosensory cortex simulations and is a function of the inhibitory output I(t).

$$\frac{dGABA}{dt} = -\kappa_{GABA} * (GABA - GABA_{base}) + \alpha_{GABA} \frac{(I_t - I_{min})}{(I_{relative} - I_{min})}$$

$$\frac{dNPY}{dt} = -\beta_{NPY}(NPY - NPY_{base}) + \alpha_{NPY} \frac{(I_t - I_{min})}{(I_{relative} - I_{min})}$$

$$\frac{dGlu}{dt} = -\beta_{Glu}Glu + Glu_{Switch}(f(K_e) + g(GABA))$$

$$f(K_e) = \frac{\beta_{Glu}}{2} \left[1 + tanh \left(\frac{K_e - K_{e,switch}}{Glu_{slope}} \right) \right]$$

$$g(GABA) = \kappa_{GABA}GABA$$
(39)

here is an inconsistency here as β_{Glu} cannot be both s^{-1} and μMs^{-1} since Glu_{switch} is non-dimensional

This allows investigation into the influence (or otherwise) of GABA and NPY during "normal" somatosensory coretx stimulation. We do not yet have any experimental evidence that this actually occurs.

| Parameter | Description | Value | |
|-----------------------|-------------|--------------------------------|--|
| α_{GABA} | | 2.6×10^{-3} μ M | |
| β_{GABA} | | 4.2×10^{-3} s^{-1} | |
| α_{NPY} | | 2.6×10^{-3} μ M | |
| β_{NPY} | | 4.2×10^{-3} s^{-1} | |
| β_{Glu} | | 4.2×10^{-3} s^{-1} | |
| $K_{e,switch}$ 5.0 mM | | 5.0 mM | |
| Glu_{slope} | | | |
| $GABA_{base}$ | | $0.0~\mu\mathrm{M}$ | |
| NPY_{base} | | $0.0 \mu \mathbf{M}$ | |
| I_{min} | | 0.225 These depend on whether | |
| | | the model uses thalamic or so- | |
| | | matosonsory input | |
| $I_{relative}$ | | 1.512 as above | |

3 Equations and Parameters for the Full NVU Model

The following parameters are given for ordinary neurovascular coupling (NVC) conditions.

| Parameter | Description | Value |
|----------------|--|---|
| \overline{F} | Faraday's constant | 96.485 C mmol ⁻¹ |
| R_g | Gas constant | $8.315 \text{ J mol}^{-1} \text{ K}^{-1}$ |
| T | Temperature constant | 300 K |
| ϕ | R_gT/F characteristic voltage | 26.7 mV |
| z_K | Ionic valence for K ⁺ | 1 |
| z_{Na} | Ionic valence for Na ⁺ | 1 |
| z_{Cl} | Ionic valence for Cl ⁻ | -1 |
| z_{NBC} | Effective valence of the NBC cotransporter complex | -1 |
| z_{Ca} | Ionic valence for Ca ²⁺ | 2 |
| γ_v | Change in membrane potential by a scaling factor | 1970 mV μ M $^{-1}$ |

3.1 Some Notes on Basic Hematology

In order to properly compare with some of the experimental results from both Sheffield (Jason Berwick's group) and others we look at some parameters corresponding to hæmoglobin dynamics. hæmoglobin concentration in blood

• Adult Male :- 135-175 gL^{-1}

• Adult Female :- 122-150 gL^{-1}

• Child :- 100-140 gL^{-1}

Molecular weight of hæmoglobin = $64450 \ gmol^{-1}$.

• Adult Male :- 2.1-2.7 mM

• adult Female :- 1.9-2.3 mM Child :- 1.55-2.2 mM

hæmoglobin, [H] decreases from the large arteries to the cerebral vasculature in a ratio of 0.69 (see Wyatt et al 1990, Jour Appli., Physiology, **68**, 1086-1091). With [H]= 2.5 mM in a rta then $[H]_{brain} = [H]_b = 2.5 \times 0.69 = 1.725 mM$. Using the Hill equation and a blood saturation of 0.75 we have that

$$[O_2]b = [O_2]_{plasma} + \frac{[H]bPO}{1 + \frac{\alpha P_{50}}{[O_2]_{plasma}}}$$
(40)

 $P_{50} = P_{O_2}$ at which [H] is 50 % saturated.

(see Valabrague et al , JCBFM, 23:536-545, 2003). Solving the Hilll equation for $[O_2]_{plasma} = 0.053 mM$. (See Hudetz, 1999, Brain Res. 817:75-83) the ratio of oxygnen concentration in cerebral tissue to plasma is approximately 0.2 = g.

• Average normal CBV: 3.5 %

• Gray Matter: 4.7 %

• White Matter: 2.6 %

• Basal Ganglia: 3.9 %

Hence

- $[HbT]_{tissue} = 1.725mM \times 3.5\% = 60\mu M.$
- Gray Matter $[HbT]_{tissue} = 1.725 mM \times 4.7\% = 82 \mu M$.
- Assume $[HbT]_{tissue} = 75 \mu M$.

Here we use the formulation of Buxton for CMRO2 as the definitions seem to be non-aligned. CMRO2 is NOT the rate at which the neuron consumes O2 but the product of the flow in the capillary, the oxygen extraction fraction and the basal oxygen concentration in the precapillary arteriole. In addition the flux J_{vasc} is now formed from the oxygen extraction fraction, CBF and the basal oxygen concentration. We define below the new extraction fraction using the analysis of Buxton and Frank and use this in the equations for HbR etc.

Extraction fraction

Assumptions

- At rest all brain capillaries are perfused. No capillary recruitment
- All O2 leaving the capillary are used by the cell's metabolism. i.e. 100 % efficiency
- exchange of O2 between plasma and RBCs is rapid and hence in equilibrium
- O2 has a probability k, of beinf extracted for metabolism. In the simple model k is proportional to the capillary wall permeability P.

Using the above assumptions then defining $C_p(t)$ as the plasma oxygen concentration, $C_T(t)$ the total oxygen concentration in the capillary then

$$\frac{dC_T(t)}{dt} = -kC_p(t) \tag{41}$$

with $C_T(0) = C_a$. The extraction fraction is defined as

$$E_t(t) = \frac{C_T(0) - C_T(t)}{C_T(0)} \tag{42}$$

Suppose that $r = \frac{C_p(t)}{C_T(t)} = \text{constant}$ and with all capillaries having the same transit time then the solution to equation (3.1) is,

$$E = 1 - e^{rkt} \tag{43}$$

Assuming that the volume of the capillary bed remains constant then the characteristic time through the capillary bed is inversely proportional to the flow velocity, $f_{in} = f$.

$$E_0 = 1 - e^{rk/f_0}$$

$$k = \frac{\ln(1 - E_0)^{f_0}}{r}$$

$$E(f) = 1 - (1 - E_0)^{f_0/f}$$
(44)

In this case f is defined to be cerebral blood flow, CBF. However the probability of oxygen crossing the BBB is proportional to the concentration gradient of oxygen in the plasma to that in the cerebral tissue = 0.2, hence we redevelop

the Buston extraction fraction equaiton to include this.

$$g = \frac{[O_2]_{tissue}}{[O_2]_{plasma}}$$

$$E(f) = 1 - (1 - E_0)^{\frac{f_0}{f(1-g)}}$$
(45)

In the model we use equation (3.1) since $E(f_{in})$ defined in equation (3.1) is not unity.

3.2 Neuron and Extracellular Space

For the simple neuron model we need to list the inputs to the astrocyte initially to ensure that all required ion concentrations are provided as time-dependent functions. The astrocyte requires concentrations of

- Glutamate
- K⁺
- Na⁺
- HCO₃
- Cl⁻

3.3 Input to the model

The input to the neuron model is in two parts. Firstly that of the activation of excitatory neurons (E(t)) defined as P(t) and secondly the activation of inhibitory neurons (I(t)) defined as Q(t). These are given in the first case as rectangular functions.

$$P(t) = \begin{cases} P_{strength} & \text{for } t_0 \le t \le t_f \\ 0 & otherwise \end{cases}$$
(46)

and

$$Q(t) = \begin{cases} Q_{strength} & \text{for } t_0 \le t \le t_f \\ 0 & otherwise \end{cases}$$
(47)

3.4 ODEs

see Table 4.2 for parameter definitions The membrane potential of the soma/axon (v_{sa}) and dendrite (v_d) (mV):

$$\frac{dE}{dt} = \frac{1}{\tau_E} \left\{ -E(t) + (k_e - r_e E(t)) S_e(c_1 E(t) - c_2 I(t) + P(t)) \right\}$$
(48)

$$\frac{dI}{dt} = \frac{1}{\tau_I} \left\{ -I(t) + (k_i - r_i I(t)) S_i(c_3 E(t) - c_4 I(t) + Q(t)) \right\}$$
(49)

here the functions S_e and S_i are given by

$$S_n(x) = \frac{1}{1 + exp[-a_n(x - \theta_n)]} - \frac{1}{1 + exp(a_n\theta_n)} \quad n \in e, i$$
 (50)

$$k_n = \max\left[S_n(x)\right] \quad n \in e, i \tag{51}$$

Table 4.2

| Parameter | Description | Value | |
|-----------|--|-------|--|
| a_e | sigmoid coefficient | 1.2 | |
| $	heta_e$ | sigmoid coefficient | 2.8 | |
| $	au_e$ | characteristic time for excitatory neurons | 3 ms | |
| a_i | sigmoid coefficient | 1.0 | |
| $	heta_i$ | sigmoid coefficient | 4.0 | |
| $	au_1$ | characteristic time for inhibitory neurons | 3 ms | |
| r_e | coefficient | 1 | |
| r_i | coefficient | 1 | |
| c_1 | coefficient | 12 | |
| c_2 | coefficient | 10 | |
| c_3 | coefficient | 13 | |
| c_4 | coefficient | 11 | |

Table 4: Nominal parameters of the neuron populations.

We should note here that the nominal parameters given in Table 4.2 could change depending on the experimental stimulation. This variation has yet to be fully investigated and described. To find the value of $k_{e,i}$ we use x = 10 such that $k_n = S_n(x)$ and this gives a maximum value.

The K⁺ in the extracellular space (ECS) and Na⁺ ion concentrations in the soma and dendrite (mM):

$$\frac{dNa_{s}a}{dt} = -\beta_{Na_{sa}}(Na_{sa} - Na_{sabase}) + \alpha_{Na_{sa}}\beta_{Na_{sa}} \frac{[|E(t) - I(t)|]}{EI_{0}}$$
(52)

$$\frac{dNa_d}{dt} = -\beta_{Na_d}(Na_d - Na_{dbase}) + \alpha_{Na_d}\beta_{Na_d} \frac{[|E(t) - I(t)|]}{EI_0}$$
 (53)

$$\frac{dK_e}{dt} = -\beta_{Ke}(K_e - K_{base}) + \alpha_{Ke}\beta_{Ke} \frac{[\mid E(t) - I(t) \mid]}{EI_0}$$
(54)

Table 3.4

| Parameter | Description | Value | |
|--------------------|-------------|---------------------|--|
| α_{Ke} | | 2.1 (this can vary) | |
| β_{Ke} | | 4.2e-3 | |
| $\alpha_{Na_{sa}}$ | | 4.23 | |
| $\beta_{Na_{sa}}$ | | 0.39e-3 | |
| α_{Na_d} | | -2.12 | |
| β_{Na_d} | | 0.75e-3 | |
| K_{base} | | 3.5 | |
| Na_{sabase} | | 9.37 | |
| Na_{dbase} | | 9.42 | |

Table 5: Nominal parameters of the neuron populations.

AGAIN we should note here that the nominal parameters given in Table 3.4 could change depending on the experimental stimulation. This variation has yet to be fully investigated and described.

3.5 Algebraic Variables

The oxygen consumption ode is the same as the old Kager neuron model.

$$\frac{dO_2}{dt} = J_{O_{2_{vascular}}} - J_{O_{2_{background}}} - J_{O_{2_{pump}}} \tag{55}$$

In the full model this PUMP is defined as $J_{O_{2pump}}$ which is function of the ECS K^+ and the soma or dendrite concentration of Na⁺. This works for the new model as all we need to know is the concentrations of ECS K^+ and the soma or dendrite concentration of Na⁺ and we have that from the solution of the 'simple' odes.

The cerebral blood flow (-):(assuming Poiseuille Flow)

$$CBF = CBF_{init} \frac{R^4}{R_{init}^4}$$
 (56)

The vascular supply of oxygen (mM s⁻¹) is defined by the amount of oxygen extracted and passed across the blood brain barrier, since CBF is defined in terms of mMs^{-1} :

$$J_{O_2 \ vascular} = CBF \frac{E(t)}{E_0} \tag{57}$$

The background oxygen consumption (mM s^{-1}):

$$J_{O_2 \ background} = J_0 P_{O_2} (1 - \gamma_{O_2}) \tag{58}$$

The normalised pump rate (-):

$$P_{O2} = \frac{J_{pump2}(O2) - J_{pump2}(0)}{J_{pump2}(O2_o) - J_{pump2}(0)}$$
(59)

We can simplify this considerably

$$J_{pump2}(O2) = 2\left[1 + \frac{O2_o}{(1 - \alpha_{O2})O2 + \alpha_{O2}O2_o}\right]^{-1}$$

$$J_{pump2}(O2_o) = 1$$

$$J_{pump2}(0) = 2\left[1 + \frac{1}{\alpha_{O2}}\right]^{-1} = \frac{2\alpha_{O2}}{1 + \alpha_{O2}}$$
(60)

so

$$J_{pump2}(O2_o) - J_{pump2}(0) = 1 - \frac{2\alpha_{O2}}{1 + \alpha_{O2}}$$
$$= \frac{1 - \alpha_{O2}}{1 + \alpha_{O2}}$$
(61)

hence the nomalised pump rate is evaluated as

$$P_{O2} = \frac{J_{pump2}(O2) - \frac{2\alpha_{O2}}{1+\alpha_{O2}}}{\frac{1-\alpha_{O2}}{1+\alpha_{O2}}}$$

$$= \frac{(1+\alpha_{O2})J_{pump2}(O2) - 2\alpha_{O2}}{1-\alpha_{O2}}$$
(62)

finally we have the reduction of oxygen due to the ATP-ase pump in the neuron.

$$J_{O_{2pump}} = J_{0}P_{0_{2}}\gamma_{O_{2}}\frac{((J_{pump1_{sa}} + J_{pump1_{d}})}{(J_{pump1init_{sa}} + J_{pump1init_{d}}))}$$

$$J_{pump1_{sa}} = \left(1 + \left(\frac{K_{e,0}}{K_{e}}\right)\right)^{-2} \left(1 + \frac{Na_{sa,0}}{Na_{sa}}\right)^{-3}$$

$$J_{pump1init_{sa}} = 0.0312$$

$$J_{pump1_{d}} = \left(1 + \left(\frac{K_{e,0}}{K_{e}}\right)\right)^{-2} \left(1 + \left(\frac{Na_{d,0}}{Na_{d}}\right)\right)^{-3}$$

$$J_{pump1init_{d}} = 0.0312$$
(63)

4 BOLD response

4.1 ODEs

The non dimensional cerebral blood volume (CBV) (-):

$$\frac{d\text{CBV}}{dt} = \frac{1}{\tau_{MTT} + \tau_{TAT}} \left(f_{in} - \text{CBV}^{\frac{1}{d}} \right)$$
 (64)

with $f_{in} = \frac{CBF}{CBF_{init}}$

The non dimensional deoxyhemoglobin (HbR) concentration (-):

$$\frac{d\text{HbR}}{dt} = \frac{1}{\tau_{MTT}} \left(\frac{f_{in}E(t)}{E_0} - \frac{\text{HbR}}{\text{CBV}} f_{out} \right)$$
 (65)

$$f_{out} = CBV^{\frac{1}{d}} + \tau_{TAT} \frac{dCBV}{dt} \tag{66}$$

Here f_{out} is the flow out of the veinous balloon.

4.2 Algebraic Variables

The non dimensional normalised total hemoglobin (HbT) concentration (-):

$$HbR_N = \frac{HbR_N}{HbR(0)} \tag{67}$$

$$HbT_{N} = \frac{CBF_{N}HbR_{N}}{CMRO_{2_{N}}}$$

$$(68)$$

where the normalised CBF is given by CBF $_N=$ CBF/CBF $_0$ 0 and CBF $_0$ 0 is the steady state value, similarly for HbR. $CMRO2=f_{in}\frac{E(t)}{E_0}$ and $CMRO2_0=1$, hence

$$HbT_N = CBF_N HbR_N \frac{E_0}{f_{in}E(t)}$$

$$= HbR \frac{E_0}{E(t)}$$
(69)

.

The non dimensional normalised oxyhemoglobin (HbO) concentration (-):

$$HbO_N = HbT_N - HbR_N + 1 (70)$$

The BOLD signal change from its steady state value (-):

$$\Delta BOLD \approx V_0 \left(a_1 \left[1 - \text{HbR}_N \right] + a_2 \left[\text{CBV}_N - 1 \right] \right) \tag{71}$$

| Parameter | Description | Value | |
|-------------|--|-------|--|
| $	au_{MTT}$ | Mean transit time | 3 s | |
| $	au_{TAT}$ | Transient adjustment time constant | 20 s | |
| d | Empirical relation between CBF and CBV | 2.5 | |
| a_1 | Weight for HbR change | 3.4 | |
| a_2 | Weight for CBV change | 1 | |
| V_0 | Resting venous blood volume fraction | 0.03 | |
| E_0 | Baseline oxygen extraction fraction | 0.4 | |

Table 6: Parameters of the BOLD submodel, for references see Mathias et al. [15].

| Parameter | Description | Value |
|-----------------------|---|-----------------------------|
| t_0 | Start time of input current | 0 s |
| t_f | Final time of input current | variable |
| $O2_0$ | Equilibrium tissue oxygen level | 0.01 mM |
| γ_{O2} | Fraction of the total oxygen consumption at steady state | 0.1 |
| J_0 | Equilibrium change in oxygen concentration due to CBF | $0.053~{ m mM~s^{-1}}$ |
| CBF_{init} | Equilibrium CBF | 0.032 |
| $J_{pump1_{sa_0}}$ | Steady state pump rate in the soma/axon | 0.0312 |
| $J_{pump1_{d_0}}$ | Steady state pump rate in the dendrite | 0.0312 |
| R_{init} | Vessel radius when passive and no stress is applied | $20~\mu\mathrm{m}$ |
| α_{O2} | Fraction of oxygen independent adenosine triphosphate (ATP) produc- | 0.05 |
| | tion | |
| $J_{pump2}(0)$ | Pump rate when oxygen concentration is 0 | 0.0952 |
| $J_{pump2}(O2_0)$ | Pump rate when oxygen is at equilibrium | 1 |
| $K_{e,0}$ | Equilibrium K_e | 2.9 mM |
| $Na_{sa,0}$ | Equilibrium Na_{sa} | 10 mM |
| $Na_{d,0}$ | Equilibrium Na_d | 10 mM |
| I_{max} | Maximum rate of Na ⁺ /K ⁺ ATP-ase pump | $0.078~\mathrm{mA~cm^{-2}}$ |

Table 7: Parameters of the neuron and extracellular space submodel, for references see Mathias et al. [15].

5 Synaptic Cleft and Astrocyte

5.1 ODEs

 K^+ concentration in the synaptic cleft (SC) (μ M):

$$\frac{dK_s}{dt} = \frac{1}{VR_{sk}} \left(J_{K_k} - 2J_{NaK_k} - J_{NKCC1_k} - J_{KCC1_k} \right) + J_{KNEtoSC}$$
 (72)

Na⁺ concentration in the SC (μ M):

$$\frac{dNa_s}{dt} = \frac{1}{VR_{sk}} \left(J_{Na_k} + 3 * J_{NaK_k} - J_{NKCC1_k} - J_{NBC_k} \right) - J_{NaNEtoSC}$$
 (73)

 $\ensuremath{\mathrm{HCO_3^-}}$ concentration in the SC (\$\mu{\rm M}\$):

$$\frac{dHCO_{3_s}}{dt} = \frac{1}{VR_{sk}} \left(-2J_{NBC_k} \right) \tag{74}$$

 K^+ concentration in the astrocyte (μM):

$$\frac{dK_k}{dt} = -J_{K_k} + 2J_{NaK_k} + J_{NKCC1_k} + J_{KCC1_k} - J_{BK_k} \tag{75}$$

Na⁺ concentration in the astrocyte (μ M):

$$\frac{dNa_k}{dt} = -J_{Na_k} - 3J_{NaK_k} + J_{NKCC1_k} + J_{NBC_k} \tag{76}$$

 HCO_3^- concentration in the astrocyte (μ M):

$$\frac{dHCO_{3_k}}{dt} = 2J_{NBC_k} \tag{77}$$

Cl⁻ concentration in the astrocyte via electroneutrality (μ M):

$$\frac{dCl_k}{dt} = \frac{dNa_k}{dt} + \frac{dK_k}{dt} - \frac{dHCO_{3_k}}{dt} + 2\frac{dCa_k}{dt}$$
(78)

The astrocytic cytosolic Ca^{2+} concentration (μM):

$$\frac{dCa_k}{dt} = B_{cyt} \left(J_{IP3_k} - J_{pump_k} + J_{ERleak_k} - \frac{J_{TRPV_k}}{r_{buff}} + J_{CICR_k} \right)$$
 (79)

note that astrocyte now has a CICR flux which Figure 16 doesn't show

The astrocytic IP₃ concentration (μ M):

$$\frac{dIP3_k}{dt} = r_h G - k_{deg} IP3_k \tag{80}$$

The astrocytic epoxyeicosatrienoic acid (EET) concentration (μ M):

$$\frac{deet_k}{dt} = V_{eet} \max(Ca_k - c_{k_{min}}, 0) - k_{eet}eet_k$$
(81)

The Ca²⁺ concentration in the astrocytic endoplasmic reticulum (ER) (μ M):

$$\frac{ds_k}{dt} = \frac{-B_{cyt}}{VR_{ERcyt}} \left(J_{IP3_k} - J_{pump_k} + J_{ERleak_k} + J_{CICR_k} \right) \tag{82}$$

Membrane potential of the astrocyte (AC) (mV):

$$\frac{dv_k}{dt} = \gamma_v \left(-J_{BK_k} - J_{K_k} - J_{Cl_k} - J_{NBC_k} - J_{Na_k} - J_{NaK_k} - 2J_{TRPV_k} - J_{GABA,k} \right) \tag{83}$$

The open probability of the BK channel (-):

$$\frac{dw_k}{dt} = \phi_n(w_\infty - w_k) \tag{84}$$

The inactivation variable h_k of the astrocytic IP₃R channel (-):

$$\frac{dh_k}{dt} = k_{on} \left[K_{inh} - \left(Ca_k + K_{inh} \right) h_k \right] \tag{85}$$

The TRPV4 channel open probability

$$\frac{dm_k}{dt} = trpv4switch\frac{(m_{inf} - m_k)}{\tau_{TRPV4}} \tag{86}$$

The concentration of arachidonic acid in the astrocyte AA_k .

$$\frac{dAA_k}{det} = \frac{AA_mAA_{max}}{(AA_m + (Ca_k - Ca_0))^2 \frac{dCa_k}{dt}} + \frac{(AA_i - AA_k)}{\tau_{AA}}$$
(87)

$$\tau_{AA} = \frac{x_{ki}^2}{2D_{AA}} \tag{88}$$

5.2 Algebraic Variables

The glutamate concentration in the SC (μ M):

$$Glu = \frac{Glu_{max}}{2} \left(1 + \tanh\left(\frac{K_e - Ke_{switch}}{Glu_{slope}}\right) \right) + \frac{Glu_{max}}{2} \left(1 + \tanh\left(\frac{GABA_N - g_{mid}}{g_{slope}}\right) \right)$$
(89)

$$J_{GABA,k} = g_{GABA}(v_k - E_{GABA}) \tag{90}$$

$$g_{GABA} = \frac{G_{GABA}}{2} \left(1 + \tanh\left(\frac{GABA_N - g_{mid}}{g_{slope}}\right) \right)$$
(91)

 G_{GABA} is the maximal conductance of the channel given by $0.3 \times G_{Cl,i}$ where $G_{Cl,i} = 1.34 \times 10^{-6} ~\mu\mathrm{M}~\mathrm{mV}^{-1}~\mathrm{ms}^{-1}$ is the conductance of the SMC Cl $^-$ leak channel taken from the excitatory NVU model (i.e. Wilson Cowan).

The flux of K^+ into the SC based on the extracellular K^+ (μMs^{-1}):

$$J_{KNEtoSC} = J_{NaNEtoSC} = c_{unit} k_{syn} \frac{dK_e}{dt}$$
(92)

 Cl^- concentration in the SC via electroneutrality (μM):

$$Cl_s = Na_s + K_s - HCO_{3_s} (93)$$

Cl⁻ flux through the Cl⁻ channel (μ M s⁻¹):

$$J_{Cl_k} = G_{Cl_k}(v_k - E_{Cl_k}) (94)$$

 K^+ flux through the K^+ channel ($\mu M s^{-1}$):

$$J_{K_k} = G_{K_k}(v_k - E_{K_k}) \tag{95}$$

Na⁺ flux through the Na⁺ channel (μ M s⁻¹):

$$J_{Na_{k}} = G_{Na_{k}}(v_{k} - E_{Na_{k}}) \tag{96}$$

Na⁺ and HCO₃⁻ flux through the NBC channel (μ M s⁻¹):

$$J_{NBC_k} = G_{NBC_k} \left(v_k - E_{NBC_k} \right) \tag{97}$$

Cl⁻ and K⁺ flux through the KCC1 channel (μ M s⁻¹):

$$J_{KCC1_k} = G_{KCC1_k} \phi \ln \left(\frac{K_s C l_s}{K_k C l_k} \right)$$
(98)

Na⁺, K⁺ and Cl⁻ flux through the NKCC1 channel (μ M s⁻¹):

$$J_{NKCC1_k} = G_{NKCC1_k} \phi \ln \left(\frac{Na_s K_s Cl_s^2}{Na_k K_k Cl_k^2} \right)$$

$$\tag{99}$$

Flux through the Na⁺/K⁺ ATP-ase pump (μ M s⁻¹):

$$J_{NaK_k} = J_{NaK_{max}} \frac{Na_k^{1.5}}{Na_k^{1.5} + K_{Na_s}^{1.5}} \frac{K_s}{K_s + K_{K_s}}$$
(100)

 K^+ flux through the BK channel (μ M s⁻¹):

$$J_{BK_{k}} = G_{BK_{k}} w_{k} (v_{k} - E_{BK_{k}}) \tag{101}$$

Nernst potential for the K^+ channel (mV):

$$E_{K_k} = \frac{\phi}{z_K} \ln \left(\frac{K_s}{K_k} \right) \tag{102}$$

Nernst potential for the Na⁺ channel (mV):

$$E_{Na_k} = \frac{\phi}{z_{Na}} \ln \left(\frac{Na_s}{Na_k} \right) \tag{103}$$

Nernst potential for the Cl⁻ channel (mV):

$$E_{Cl_k} = \frac{\phi}{z_{Cl}} \ln \left(\frac{Cl_s}{Cl_k} \right) \tag{104}$$

Nernst potential for the NBC channel (mV):

$$E_{NBC_k} = \frac{\phi}{z_{NBC}} \ln \left(\frac{Na_s HCO_{3_s}^2}{Na_k HCO_{3_k}^2} \right)$$
 (105)

Nernst potentical for TRPV4 channel (mV):

$$E_{TRPV_k} = \frac{\phi}{z_{Ca}} ln\left(\frac{Ca_p}{Ca_k}\right) \tag{106}$$

Nernst potential for the BK channel (mV):

$$E_{BK_k} = \frac{\phi}{z_K} \ln \left(\frac{K_p}{K_k} \right) \tag{107}$$

The time constant associated with the opening of the BK channel (s^{-1}):

$$\phi_n = \psi_n \cosh\left(\frac{v_k - v_3}{2v_4}\right) \tag{108}$$

The equilibrium state of the BK channel (-):

$$w_{\infty} = \frac{1}{2} \left(1 + \tanh\left(\frac{v_k + eet_{shift}eet_k - v_3}{v_4}\right) \right)$$
 (109)

The voltage associated with half open probability (mV):

$$v_3 = -\frac{v_5}{2} \tanh\left(\frac{Ca_k - Ca_3}{Ca_4}\right) + v_6 \tag{110}$$

The ratio ρ of bound to unbound metabotropic receptors on the astrocytic process adjacent to the SC (-):

$$\rho = \rho_{\min} + \frac{\rho_{\max} - \rho_{\min}}{Glu_{\max}}Glu \tag{111}$$

The ratio G of active to total G-protein due to metabotropic glutamate receptor (mGluR) binding on the astrocyte endfoot surround the SC (-):

$$G = \frac{\rho + \delta_G}{K_G + \rho + \delta_G} \tag{112}$$

Fast Ca²⁺ buffering in the astrocytic cytosol is described within the steady state approximation (-):

$$B_{cyt} = \left(1 + BK_{end} + \frac{K_{ex}B_{ex}}{(K_{ex} + Ca_k)^2}\right)^{-1}$$
(113)

The flux of Ca²⁺ through the IP₃R channel (μ M s⁻¹):

$$J_{IP3_k} = J_{max} \left[\left(\frac{IP3_k}{IP3_k + K_i} \right) \left(\frac{Ca_k}{Ca_k + K_{act_k}} \right) h_k \right]^3 \left(1 - \frac{Ca_k}{s_k} \right)$$
(114)

The flux of Ca^{2+} through the uptake pump ($\mu M s^{-1}$):

$$J_{pump_k} = V_{max} \frac{Ca_k^2}{Ca_k^2 + k_{pump}^2} \tag{115}$$

The flux of $\mathrm{Ca^{2+}}$ through the leak channel ($\mu\mathrm{M}~\mathrm{s^{-1}}$):

$$J_{ERleak_k} = P_L \left(1 - \frac{Ca_k}{s_k} \right) \tag{116}$$

flux of Ca²⁺ through CICR into astrocytic cytosol

$$J_{CICR_k} = C_k * \frac{s_k^4}{sc_k^4 + s_k^4} \frac{Ca_k^4}{cc_k^4 + Ca_k^4}$$
(117)

flux of Ca²⁺ through TRPV4 channel

$$J_{TRPV_k} = G_{TRPV_k} m_k (v_k - E_{TRPV_k}) \tag{118}$$

| Parameter | Description | Value |
|------------------|--|---|
| VR_{sk} | Volume ratio between the SC and astrocyte | 0.465 |
| Glu_{max} | Maximum glutamate concentration (one vesicle) | $1846~\mu\mathrm{M}$ |
| $K_{e_{switch}}$ | Threshold past which glutamate is released | 5.0 mM |
| Glu_{slope} | Slope of glutamate sigmoidal | 0.1 mM |
| c_{unit} | Constant to convert from mM to μ M | 10^{3} |
| k_{syn} | The number of active synapses per astrocytic process | 11.5 |
| G_{K_k} | Whole cell conductance of K ⁺ | $6907.77~\mu{ m M~mV^{-1}~s^{-1}}$ |
| G_{Na_k} | Whole cell conductance of Na ⁺ | 226.94 $\mu { m M~mV^{-1}~s^{-1}}$ |
| G_{NBC_k} | Whole cell conductance of the NBC cotransporter | $130.74~\mu{ m M~mV^{-1}~s^{-1}}$ |
| G_{KCC1_k} | Whole cell conductance of the KCC1 cotransporter | $1.728~\mu{ m M~mV^{-1}~s^{-1}}$ |
| G_{NKCC1_k} | Whole cell conductance of the NKCC1 cotransporter | $9.568~\mu{ m M}~{ m mV}^{-1}~{ m s}^{-1}$ |
| G_{BK_k} | Whole cell conductance of the BK channel | $10.25~\mu{ m M~mV^{-1}~s^{-1}}$ |
| G_{Cl_k} | Whole cell conductance of Cl ⁻ | $151.93~\mu{ m M~mV^{-1}~s^{-1}}$ |
| G_{TRPV_k} | Whole cell conductance of TRPV4 channel | $3.15 \mathrm{e}\text{-4}~\mu\mathrm{M}~\mathrm{mV}^{-1}~\mathrm{s}^{-1}$ |
| $J_{NaK_{max}}$ | Maximum flux through the Na ⁺ /K ⁺ ATP-ase pump | $2.37{	imes}10^4~\mu{ m M~s}^{-1}$ |
| K_{Na_k} | Na ⁺ /K ⁺ ATP-ase pump constant | $10 \times 10^3 \; \mu \mathrm{M}$ |
| K_{K_s} | Na ⁺ /K ⁺ ATP-ase pump constant | $1.5 \times 10^3 \; \mu \mathrm{M}$ |
| $ ho_{ m min}$ | Minimum ratio of bound to unbound IP ₃ receptors | 0.1 |
| $ ho_{ m max}$ | Maximum ratio of bound to unbound IP ₃ receptors | 0.7 |
| δ_G | Ratio of the activities of the unbound and bound receptors | 1.235×10^{-2} |
| K_G | G-protein disassociation constant | 8.82 |
| r_h | Maximum rate of IP ₃ production in astrocyte due to glutamate receptors | $4.8~\mu{ m M}~{ m s}^{-1}$ |
| k_{deg} | Rate constant for IP ₃ degradation in astrocyte | 1.25 s^{-1} |
| r_{buff} | Rate of Ca ²⁺ buffering at the endfoot compared to the astrocyte body | 0.05 |
| VR_{ERcyt} | Volume ratio between ER and astrocytic cytosol | 0.185 |
| BK_{end} | Ratio of endogenous buffer concentration to disassociation constant | 40 |
| K_{ex} | Disassociation constant of exogenous buffer | $0.26~\mu\mathrm{M}$ |
| B_{ex} | Concentration of exogenous buffer | $11.35~\mu\mathrm{M}$ |
| J_{max} | Maximum rate of Ca ²⁺ through the IP ₃ mediated channel | $2880~\mu{ m M}~{ m s}^{-1}$ |
| K_i | Disassociation constant for IP ₃ binding to an IP_3R | $0.03~\mu\mathrm{M}$ |

| K_{act_k} | Disassociation constant for Ca^{2+} binding to an activation site on an IP_3R | $0.17~\mu\mathrm{M}$ |
|---------------|---|--|
| k_{on} | Rate of Ca^{2+} binding to the inhibitory site on the IP_3R | $2~\mu\mathrm{M}^{-1}~\mathrm{s}^{-1}$ |
| K_{inh} | Disassociation constant of IP_3R | $0.1~\mu\mathrm{M}$ |
| V_{max} | Maximum rate of Ca ²⁺ uptake pump on the ER | $20~\mu\mathrm{M~s^{-1}}$ |
| k_{pump} | Ca ²⁺ uptake pump disassociation constant | $0.24~\mu\mathrm{M}$ |
| P_L | ER leak channel steady state balance constant | $0.0804~\mu{ m M~s^{-1}}$ |
| V_{eet} | EET production rate | 72 s^{-1} |
| $c_{k_{min}}$ | Minimum Ca ²⁺ concentration required for EET production | $0.1~\mu\mathrm{M}$ |
| k_{eet} | EET degradation rate | 7.2 s^{-1} |
| v_4 | Measure of the spread of w_{∞} | 8 mV |
| eet_{shift} | Describes the EET dependent voltage shift | 2 mV μ M $^{-1}$ |
| v_5 | Determines the range of the shift of w_{∞} as Ca^{2+} varies | 15 mV |
| v_6 | Shifts the range of w_{∞} | $-55~\mathrm{mV}$ |
| ψ_n | Characteristic time for the opening of the BK channel | 2.664 s^{-1} |
| Ca_3 | BK open probability Ca ²⁺ constant | $0.4~\mu\mathrm{M}$ |
| Ca_4 | BK open probability Ca ²⁺ constant | $0.35~\mu\mathrm{M}$ |
| AA_m | dissociation constant for AA_k | $0.161~\mu\mathrm{M}$ |
| AA_{max} | maximum rate of generation of AA_k | $29~\mu{ m M}~{ m s}^{-1}$ |
| Ca_0 | baseline Ca_k^{2+} | $0.1432~\mu\mathrm{M}$ |
| D_{AA} | diffusion coefficient for AA | $0.152~\mu m^2 m s^{-1}$ |
| E_{GABA} | Nernst potential for GABA | -75 mV |
| g_{mid} | midpoint of the GABA sigmoidal | 0.6 |
| g_{slope} | scaling for the GABA sigmoidal | 0.1 |
| C_k | V_{max} for astrocyte CICR | $30~\mu\mathrm{M}~\mathrm{ms}^{-1}$ |
| sc_k | first Disassociation constant for astrocytic CICR flux | $2~\mu\mathrm{M}$ |
| cc_k | second Disassociation constant for astrocytic CICR flux | $0.9~\mu\mathrm{M}$ |

Table 8: Parameters of the astrocyte and SC submodel, for references see Dormanns et al. [4], Kenny et al. [9].

6 PVS

6.1 ODEs

 K^+ concentration in the PVS (μM):

$$\frac{dK_p}{dt} = \frac{J_{BK_k}}{VR_{pk}} + \frac{J_{KIR_i}}{VR_{pi}} - K_{decay_p}(K_p - K_{min_p})$$
(119)

 Ca^{2+} concentration in the PVS (μ M):

$$\frac{dCa_p}{dt} = \frac{J_{TRPV_k}}{VR_{pk}} + \frac{J_{VOCC_i}}{VR_{pi}} - Ca_{decay_p}(Ca_p - Ca_{min_p})$$
(120)

The open probability of the transient receptor potential vanniloid-related 4 (TRPV4) channel (-):

$$\frac{dm_k}{dt} = \frac{m_{\infty_k} - m_k}{t_{TRPV_k}} \tag{121}$$

6.2 Algebraic Variables

The flux of Ca²⁺ through the VOCC which connects the SMC to the PVS (μ M s⁻¹):

$$J_{VOCC_i} = G_{Cai} \frac{v_i - v_{Ca1}}{1 + \exp\left[-(v_i - v_{Ca2})/R_{Cai}\right]}$$
(122)

This VOCC channel is altered when the interneuron model is used. See below equation (158)

The flux of Ca²⁺ through the TRPV4 channel (μ M s⁻¹):

$$J_{TRPV_k} = G_{TRPV_k} m_k (v_k - E_{TRPV_k})$$
(123)

The Nernst potential of the TRPV4 channel (mV):

$$E_{TRPV_k} = \frac{\phi}{z_{Ca}} \ln \left(\frac{Ca_p}{Ca_k} \right) \tag{124}$$

The equilibrium state of the TRPV4 channel (-):

$$m_{\infty_k} = \Gamma_m \left[\frac{1}{1 + H_{Ca_k}} \left(H_{Ca_k} + \tanh\left(\frac{v_k - v_{1,TRPV}}{v_{2,TRPV}}\right) \right) \right]$$
 (125)

The material strain gating term (-):

$$\Gamma_m = \frac{1}{1 + \exp\left(-\frac{\eta - \eta_0}{\kappa_k}\right)} \tag{126}$$

The strain on the perivascular endfoot of the astrocyte (-)

$$\eta = \frac{R - R_{init}}{R_{init}} \tag{127}$$

The Ca²⁺ inhibitory term (-)

$$H_{Ca_k} = \frac{Ca_k}{\gamma_{Cai}} + \frac{Ca_p}{\gamma_{Cae}} \tag{128}$$

| Parameter | Description | Value |
|----------------|--|--|
| VR_{pk} | Volume ratio between PVS and astrocyte | 0.001 |
| VR_{pi} | Volume ratio between PVS and SMC | 0.001 |
| K_{decay_p} | Rate of decay of K ⁺ in PVS | 0.15 s^{-1} |
| K_{min_p} | Steady state value of K ⁺ in PVS | $3 \times 10^3 \ \mu \text{M}$ |
| Ca_{decay_p} | Rate of decay of Ca ²⁺ in PVS | $0.5 \ s^{-1}$ |
| Ca_{min_p} | Steady state value of Ca ²⁺ in PVS | $2 	imes 10^3 \ \mu 	extsf{M}$ |
| G_{Cai} | VOCC whole cell conductance | $1.29\times 10^{-3}~\mu{\rm M~mV^{-1}~s^{-1}}$ |
| v_{Ca1} | VOCC reversal potential | 100 mV |
| v_{Ca2} | Half point of the VOCC activation sigmoidal | $-24~\mathrm{mV}$ |
| R_{Cai} | Maximum slope of the VOCC activation sigmoidal | 8.5 mV |
| G_{TRPV_k} | TRPV4 whole cell conductance | $3.15 \times 10^{-4} \; \mu \mathrm{M \; mV^{-1} \; s^{-1}}$ |
| t_{TRPV_k} | Characteristic time constant for m_k | 0.9 s |
| η_0 | Strain required for half activation of the TRPV4 channel | 0.1 |
| κ_k | TRPV4 channel strain constant | 0.1 |
| $v_{1,TRPV}$ | TRPV4 channel voltage gating constant | 120 mV |
| $v_{2,TRPV}$ | TRPV4 channel voltage gating constant | 13 mV |
| γ_{Cai} | Ca ²⁺ concentration constant | $0.01~\mu\mathrm{M}$ |
| γ_{Cae} | Ca ²⁺ concentration constant 33 | $200~\mu\mathrm{M}$ |

Table 9: Parameters of the PVS compartment, for references see Dormanns et al. [4], Kenny et al. [9].

SMC

7.1 **ODEs**

Cytosolic Ca²⁺ in the SMC (μ M):

$$\frac{dCa_i}{dt} = J_{IP_{3i}} - J_{SR_{uptake_i}} + J_{CICR_i} - J_{extrusion_i} + J_{SR_{leak_i}} \dots
-J_{VOCC_i} + J_{Na/Ca_i} - 0.1J_{stretch_i} + J_{Ca^{2+}-coupling_i}^{SMC-EC}$$
(129)

 Ca^{2+} in the sarcoplasmic reticulum (SR) of the SMC (μ M):

$$\frac{ds_i}{dt} = J_{SR_{uptake_i}} - J_{CICR_i} - J_{SR_{leak_i}} \tag{130}$$

Membrane potential of the SMC (mV):

$$\frac{dv_i}{dt} = -\gamma_v (J_{NaK_i} + J_{Cl_i} + 2J_{VOCC_i} + J_{Na/Ca_i} + J_{K_i} + J_{stretch_i} + J_{KIR_i}) + V_{coupling_i}^{SMC-EC}$$
(131)

Open state probability of Ca²⁺-activated K⁺ channels (-):

$$\frac{dw_i}{dt} = \lambda_i \left(K_{act_i} - w_i \right) \tag{132}$$

IP₃ concentration in the SMC (μ M):

$$\frac{dIP3_i}{dt} = -J_{degrad_i} + J_{IP_3-coupling_i}^{SMC-EC}$$
(133)

Arachidonic acid in the SMC, this is just diffused from the astrocyte.

$$\tau_{AA} = \frac{x_{ki}^2}{D_{AA}} \tag{134}$$

$$\tau_{AA} = \frac{x_{ki}^2}{D_{AA}}$$

$$\frac{dAA_i}{dt} = \frac{AA_k - AA_i}{\tau_{AA}}$$
(134)

20-HETE in the SMC

$$\frac{dH_i}{dt} = \frac{1}{1 + exp(\frac{(NO_i - NO_{rest})}{R_{NO}})} \frac{V_a A A_i}{K_a + A A_i} + \frac{V_f A A_i}{K_f + A A_i} - \lambda_h H_i$$
(136)

(137)

Here there are two CYP generators of 20-HETE. The CYP450 generator is inhibited by NO hence the function multiplying the first term of the o.d.e. for 20-HETE. This model is a phenomenological one and relies on the results from [12]

Algebraic Variables

Release of Ca²⁺ from IP₃ sensitive stores in the SMC (μ M s⁻¹):

$$J_{IP3_i} = F_i \frac{IP3_i^2}{K_{ri}^2 + IP3_i^2} \tag{138}$$

Uptake of Ca²⁺ into the SR (μ M s⁻¹):

$$J_{SR_{uptake_i}} = B_i \frac{Ca_i^2}{c_{bi}^2 + Ca_i^2}$$
 (139)

 Ca^{2+} induced Ca^{2+} release (CICR) (μ M s⁻¹):

$$J_{CICR_i} = C_i \frac{s_i^2}{s_{ci}^2 + s_i^2} \frac{Ca_i^4}{c_{ci}^4 + Ca_i^4}$$
(140)

 Ca^{2+} extrusion by Ca^{2+} -ATP-ase pumps ($\mu M s^{-1}$):

$$J_{extrusion_i} = D_i C a_i \left(1 + \frac{v_i - v_d}{R_{di}} \right) \tag{141}$$

Leak current from the SR (μ M s $^{-1}$):

$$J_{SR_{leak:}} = L_i s_i \tag{142}$$

Flux of ${\rm Ca^{2+}}$ exchanging with ${\rm Na^{+}}$ in the ${\rm Na^{+}/Ca^{2+}}$ exchange ($\mu{\rm M~s^{-1}}$):

$$J_{Na/Ca_{i}} = G_{Na/Ca_{i}} \frac{Ca_{i}}{Ca_{i} + c_{Na/Ca_{i}}} \left(v_{i} - v_{Na/Ca_{i}} \right)$$
(143)

 ${
m Ca^{2+}}$ flux through the stretch-activated channels in the SMC ($\mu{
m M~s^{-1}}$):

$$J_{stretch_i} = \frac{G_{stretch}}{1 + \exp\left(-\alpha_{stretch}\left(\frac{\Delta pR}{h} - \sigma_0\right)\right)} \left(v_i - E_{SAC}\right) \tag{144}$$

Flux through the Na⁺/K⁺ pump (μ M s⁻¹):

$$J_{NaK_i} = F_{NaK} \tag{145}$$

Cl⁻ flux through the Cl⁻ channel (μ M s⁻¹):

$$J_{Cl_i} = G_{Cli} \left(v_i - v_{Cli} \right) \tag{146}$$

 K^+ flux through K^+ channel ($\mu M s^{-1}$):

$$J_{K_i} = G_{K_i} w_i \left(v_i - v_{K_i} \right) \tag{147}$$

Flux through inward rectifying K⁺ (KIR) channels in the SMC (μ M s⁻¹):

$$J_{KIR_i} = G_{KIR_i}(v_i - v_{KIR_i}) \tag{148}$$

IP₃ degradation (μ M s⁻¹):

$$J_{dearad_i} = k_{di}IP3_i (149)$$

Nernst potential of the KIR channel in the SMC (mV):

$$v_{KIR_i} = z_1 K_p - z_2 (150)$$

Conductance of KIR channel (μ M mV⁻¹ s⁻¹):

$$G_{KIR_i} = F_{KIR_i} \exp(z_5 v_i + z_3 K_p - z_4)$$
(151)

Equilibrium distribution of open channel states for the SMC BK channels (-):

$$K_{act_i} = \frac{(Ca_i + c_{w,i})^2}{(Ca_i + c_{w,i})^2 + \alpha_{act_i} \exp(-([v_i - v_{Ca_{3i}} - h_{shift}(H_i - H_0)]/R_{Ki}))}$$
(152)

Translation factor, regulatory effect of cyclic guanosine monophosphate (cGMP) on the BK channel open probability (μ M):

$$c_{\mathbf{w},i} = \frac{\beta_{w,i}}{2} \left(1 + \tanh\left(\frac{\mathsf{cGMP}_i - \alpha_{w,i}}{\epsilon_{w,i}}\right) \right) \tag{153}$$

Heterocellular electrical coupling between SMCs and ECs (mV s⁻¹):

$$V_{coupling_i}^{SMC-EC} = -G_{coup}(v_i - v_j)$$
(154)

Heterocellular IP₃ coupling between SMCs and ECs (μ M s⁻¹):

$$J_{IP_3-coupling_i}^{SMC-EC} = -P_{IP_3}(IP_{3i} - IP_{3j})$$

$$\tag{155}$$

 Ca^{2+} coupling between SMCs and ECs (μ M s⁻¹):

$$J_{Ca^{2+}-coupling_{i}}^{SMC-EC} = -P_{Ca^{2+}}(Ca_{i} - Ca_{j})$$
(156)

Finally the flux of Ca²⁺ through the VOCC channel on the SMC is given by

$$J_{VOCC,i} = g_{VOCC} \frac{v_i - v_{Ca1}}{1 + \exp\left(\frac{-(v_i - v_{Ca2})}{R_{C}a}\right)}$$
(157)

which is the same as equation (122) where $v_{Ca1} = 100$ mV is the reversal potential, $v_{Ca2} = -24$ mV is the half-point of the VOCC activation sigmoidal, and $R_{Ca} = 8.5$ mV is the maximum slope of the sigmoidal (all taken from the NVU model). However for the interneuron model the NPY conductance is given by

$$g_{VOCC} = G_{Ca,i} \left(1 + \frac{N_{inc}}{2} \tanh \left(\frac{NPY_N - N_{mid}}{N_{slope}} \right) \right)$$
 (158)

where $G_{Ca,i}=1.29\times 10^{-6}~\mu\mathrm{M}~\mathrm{mV}^{-1}~\mathrm{ms}^{-1}$ is the base conductance of the VOCC (taken from the NVU model), $N_{inc}=0.05$ is the proportional increase of the conductance from baseline due to NPY (i.e. 0.05 means a 5% increase from $G_{Ca,i}$ when NPY is released, model estimate), $N_{mid}=0.6$ is the midpoint of the sigmoidal, and N_{slope} is the slope of the sigmoidal (both model estimates). Hence when $NPY_N=0$, $g_{VOCC}=G_{Ca,i}$ and when $NPY_N=1$, $g_{VOCC}=1.05\times G_{Ca,i}$.

| Parameter | Description | Value |
|---------------|---|--|
| λ_i | Rate constant for opening | 45 s^{-1} |
| F_i | Maximal rate of activation-dependent Ca ²⁺ influx | $0.23~\mu{ m M}~{ m s}^{-1}$ |
| K_{ri} | Half-saturation constant for agonist-dependent Ca ²⁺ entry | $1~\mu\mathrm{M}$ |
| B_i | SR uptake rate constant | $2.025~\mu{ m M~s^{-1}}$ |
| c_{bi} | Half-point of the SR ATP-ase activation sigmoidal | $1 \mu M$ |
| C_i | CICR rate constant | $55~\mu{ m M}~{ m s}^{-1}$ |
| s_{ci} | Half-point of the CICR Ca ²⁺ efflux sigmoidal | $2~\mu\mathrm{M}$ |
| c_{ci} | Half-point of the CICR activation sigmoidal | $0.9~\mu\mathrm{M}$ |
| D_i | Rate constant for Ca ²⁺ extrusion by the ATP-ase pump | 0.24 s^{-1} |
| v_d | Intercept of voltage dependence of extrusion ATP-ase | $-100~\mathrm{mV}$ |
| R_{di} | Slope of voltage dependence of extrusion ATP-ase | 250 mV |
| L_i | Leak from SR rate constant | $0.025 \ \mathrm{s^{-1}}$ |
| G_{Cai} | Whole-cell conductance for VOCCs | $1.29{	imes}10^{-3}~\mu{ m M~mV^{-1}~s^{-1}}$ |
| $v_{Ca_{1i}}$ | Reversal potential for VOCCs | 100 mV |
| $v_{Ca_{2i}}$ | Half-point of the VOCC activation sigmoidal | $-24~\mathrm{mV}$ |
| R_{Cai} | Maximum slope of the VOCC activation sigmoidal | 8.5 mV |
| $G_{Na/Cai}$ | Whole-cell conductance for Na ⁺ /Ca ²⁺ exchange | $3.16{	imes}10^{-3}~\mu{ m M~mV^{-1}~s^{-1}}$ |
| $c_{Na/Cai}$ | Half-point for activation of Na ⁺ /Ca ²⁺ exchange by Ca ²⁺ | $0.5~\mu\mathrm{M}$ |
| $v_{Na/Cai}$ | Reversal potential for the Na ⁺ /Ca ²⁺ exchanger | -30 mV |
| $G_{stretch}$ | Whole cell conductance for stretch activated channels (SACs) | $6.1 \times 10^{-3} \ \mu \mathrm{M \ mV^{-1} \ s^{-1}}$ |

| $\alpha_{stretch}$ | Slope of stress dependence of the SAC activation sigmoidal | $7.4{	imes}10^{-3}~{ m mmHg}^{-1}$ |
|--------------------|---|---|
| Δp | Pressure difference over vessel | 30 mmHg |
| σ_0 | Half-point of the SAC activation sigmoidal | 500 mmHg |
| E_{SAC} | Reversal potential for SACs | -18 mV |
| F_{NaK} | Rate of the K ⁺ influx by the Na ⁺ /K ⁺ pump | $4.32 \times 10^{-2} \ \mu \text{M s}^{-1}$ |
| G_{Cli} | Whole-cell conductance for Cl ⁻ current | $1.34{	imes}10^{-3}~\mu{ m M~mV^{-1}~s^{-1}}$ |
| v_{Cli} | Reversal potential for Cl ⁻ channels | $-25~\mathrm{mV}$ |
| G_{Ki} | Whole-cell conductance for K ⁺ efflux | $4.46 \times 10^{-3} \ \mu \mathrm{M \ mV^{-1} \ s^{-1}}$ |
| v_{K_i} | Nernst potential | -94 mV |
| k_{di} | Rate constant of IP ₃ degradation | $0.1 \ s^{-1}$ |
| F_{KIR_i} | Scaling factor of K ⁺ efflux through the KIR channel | $1.285{	imes}10^{-6}\mu{ m M~mV^{-1}~s^{-1}}$ |
| z_1 | Model estimation for membrane voltage KIR channel | $4.5{	imes}10^3$ mV μ M $^{-1}$ |
| z_2 | Model estimation for membrane voltage KIR channel | 112 mV |
| z_3 | Model estimation for the KIR channel conductance | $4.2 	imes 10^{-4}~\mu$ M $^{-1}$ |
| z_4 | Model estimation for the KIR channel conductance | 12.6 |
| z_5 | Model estimation for the KIR channel conductance | $-7.4 	imes 10^{-2} \; \mathrm{mV^{-1}}$ |
| α_{act_i} | Translation factor for v_i dependence of K_{act_i} sigmoidal | $0.13~\mu\mathrm{M}^{2}$ |
| $v_{Ca_{3i}}$ | Half-point for the K_{act_i} activation sigmoidal | $-27~\mathrm{mV}$ |
| R_{Ki} | Maximum slope of the K_{act_i} activation sigmoidal | 12 mV |
| h_{shift} | 20-HETE scaling parameter in K_{act_i} activation sigmoidal | 10 |
| H_0 | 20-HETE shift parameter in K_{act_i} activation sigmoidal | $0.126~\mu\mathrm{M}$ |
| λ_h | 20-HETE degradation parameter | $2.0e-3 s^{-1}$ |
| $\beta_{w,i}$ | Constant to fit data | $1~\mu\mathrm{M}$ |
| $\alpha_{w,i}$ | Constant to fit data | $10.75~\mu\mathrm{M}$ |
| $\epsilon_{w,i}$ | Constant to fit data | $0.668~\mu\mathrm{M}$ |
| G_{coup} | Heterocellular electrical coupling coefficient | $0.5 \ \mathrm{s}^{-1}$ |
| P_{IP_3} | Heterocellular IP ₃ coupling coefficient | $0.05 \ \mathrm{s^{-1}}$ |
| $P_{Ca^{2+}}$ | Heterocellular $P_{Ca^{2+}}$ coupling coefficient | 0.05 s^{-1} |

Table 10: Parameters of the SMC compartment, for references see Dormanns et al. [4].

8 EC

8.1 ODEs

Cytosolic Ca $^{2+}$ concentration in the EC (μ M):

$$\frac{dCa_j}{dt} = J_{IP_{3j}} - J_{ER_{uptake_j}} + J_{CICR_j} - J_{extrusion_j} \dots
+ J_{ER_{leak_j}} + J_{cation_j} + J_{0_j} - J_{stretch_j} - J_{Ca^{2+}-coupling_j}^{SMC-EC}$$
(159)

 $\mathrm{Ca^{2+}}$ concentration in the ER in the EC ($\mu\mathrm{M}$):

$$\frac{ds_j}{dt} = J_{ER_{uptake_j}} - J_{CICR_j} - J_{ER_{leak_j}} \tag{160}$$

Membrane potential of the EC (mV):

$$\frac{dv_j}{dt} = -\frac{1}{C_{m_j}} (I_{K_j} + I_{R_j}) - V_{coupling_j}^{SMC-EC}$$

$$\tag{161}$$

 IP_3 concentration of the EC (μ M):

$$\frac{dIP3_j}{dt} = J_{PLC} - J_{degrad_j} - J_{IP_3-coupling_j}^{SMC-EC}$$
(162)

8.2 Algebraic Variables

Release of Ca^{2+} from IP₃-sensitive stores in the EC (μ M s⁻¹):

$$J_{IP_{3j}} = F_j \frac{IP3_j^2}{K_{rj}^2 + IP3_j^2} \tag{163}$$

Uptake of Ca²⁺ into the endoplasmic reticulum (μ M s⁻¹):

$$J_{ER_{uptake_{j}}} = B_{j} \frac{Ca_{j}^{2}}{c_{b_{j}}^{2} + Ca_{j}^{2}}$$
(164)

CICR (μ M s⁻¹):

$$J_{CICR_j} = C_j \frac{s_j^2}{s_{cj}^2 + s_j^2} \frac{Ca_j^4}{c_{cj}^4 + Ca_j^4}$$
(165)

 ${
m Ca^{2+}}$ extrusion by ${
m Ca^{2+}}$ -ATP-ase pumps ($\mu{
m M~s^{-1}}$):

$$J_{extrusion_j} = D_j C a_j (166)$$

 ${
m Ca^{2+}}$ flux through the stretch-activated channels in the EC ($\mu{
m M~s^{-1}}$):

$$J_{stretch_j} = \frac{G_{stretch}}{1 + \exp\left(-\alpha_{stretch}\left(\frac{\Delta pR}{h} - \sigma_0\right)\right)} \left(v_j - E_{SAC}\right)$$
(167)

Leak current from the ER (μ M s⁻¹):

$$J_{ER_{leak_j}} = L_j s_j \tag{168}$$

 Ca^{2+} influx through nonselective cation channels ($\mu M s^{-1}$):

$$J_{cation_j} = G_{cat_j} (E_{Ca_j} - v_j) \frac{1}{2} \left(1 + \tanh \left(\frac{\log_{10} (Ca_j / c_{log}) - m_{3_{cat_j}}}{m_{4_{cat_j}}} \right) \right)$$
 (169)

 K^+ current through the BK_{Caj} channel and the SK_{Caj} channel (fA):

$$I_{K_j} = G_{totj}(v_j - v_{K_j}) \left(I_{BK_{Caj}} + I_{SK_{Caj}} \right)$$

$$\tag{170}$$

 K^+ efflux through the BK_{Caj} channel (-):

$$I_{BK_{Caj}} = 0.2 \left(1 + \tanh \left(\frac{(\log_{10}(Ca_j/c_{log}) - c)(v_j - b_j) - a_{1j}}{m_{3bj}(v_j + a_{2j}(\log_{10}(Ca_j/c_{log}) - c) - b_j)^2 + m_{4bj}} \right) \right)$$
(171)

 K^+ efflux through the SK_{Caj} channel (-):

$$I_{SK_{Caj}} = 0.3 \left(1 + \tanh \left(\frac{\log_{10}(Ca_j/c_{log}) - m_{3sj}}{m_{4sj}} \right) \right)$$
 (172)

Residual current regrouping Cl⁻ and Na⁺ current flux (fA):

$$I_{R_i} = G_{R_i}(v_j - v_{restj}) \tag{173}$$

IP₃ degradation (μ M s⁻¹):

$$J_{degrad_j} = k_{dj} IP3_j (174)$$

coupling between SMC and EC

membrane coupling

$$V_{coupling_j}^{SMC-EC} = G_{coup}(v_i - v_j)$$
(175)

Ca²⁺ coupling

$$J_{Ca^{2+}-coupling_j}^{SMC-EC} = Ca_{coup}(Ca_i - Ca_j)$$

$$(176)$$

IP₃ coupling

$$J_{IP_3-coupling_j}^{SMC-EC} = IP_{3,coup}(I_i - I_j)$$
(177)

| Parameter | Description | Value |
|----------------|---|--|
| J_{0_j} | Constant Ca ²⁺ influx | $0.029~\mu{ m M~s^{-1}}$ |
| C_{m_j} | Membrane capacitance | 25.8 pF |
| J_{PLC} | IP ₃ production rate | $0.11~\mu{ m M}~{ m s}^{-1}$ |
| F_j | Maximal rate of activation-dependent Ca ²⁺ influx | $0.23~\mu{ m M~s^{-1}}$ |
| K_{rj} | Half-saturation constant for agonist-dependent Ca ²⁺ entry | $1~\mu\mathrm{M}$ |
| B_j | ER uptake rate constant | $0.5~\mu{ m M}~{ m s}^{-1}$ |
| c_{bj} | Half-point of the SR ATP-ase activation sigmoidal | $1~\mu\mathrm{M}$ |
| C_j | CICR rate constant | $5~\mu\mathrm{M}~\mathrm{s}^{-1}$ |
| s_{cj} | Half-point of the CICR Ca ²⁺ efflux sigmoidal | $2~\mu\mathrm{M}$ |
| c_{cj} | Half-point of the CICR activation sigmoidal | $0.9~\mu\mathrm{M}$ |
| D_j | Rate constant for Ca ²⁺ extrusion by the ATP-ase pump | $0.24 \ { m s}^{-1}$ |
| L_j | Rate constant for Ca ²⁺ leak from the ER | $0.025 \ \mathrm{s}^{-1}$ |
| G_{catj} | Whole-cell cation channel conductivity | $6.6{	imes}10^{-4}~\mu{ m M~mV^{-1}~s^{-1}}$ |
| E_{Caj} | Ca ²⁺ equilibrium potential | 50 mV |
| c_{log} | Log constant | $1~\mu\mathrm{M}$ |
| G_{totj} | Total K ⁺ channel conductivity | 6927 pS |
| v_{Kj} | K ⁺ equilibrium potential | -80 mV |
| $m_{3_{catj}}$ | Model constant, further explanation see [10] | -0.18 |
| $m_{4_{catj}}$ | Model constant, further explanation see [10] | 0.37 |
| 2 | Model constant, further explanation see [10] | -0.4 |
| b_j | Model constant, further explanation see [10] | $-80.8~\mathrm{mV}$ |
| a_{1j} | Model constant, further explanation see [10] | 53.3 mV |
| a_{2j} | Model constant, further explanation see [10] | 53.3 mV |
| m_{3bj} | Model constant, further explanation see [10] | $1.32{	imes}10^{-3}~{ m mV}^{-1}$ |
| m_{4bj} | Model constant, further explanation see [10] | 0.30 mV |
| m_{3sj} | Model constant, further explanation see [10] | -0.28 |
| m_{4sj} | Model constant, further explanation see [10] | 0.389 |
| G_{R_i} | Residual current conductivity | 955 pS |
| v_{restj} | Membrane resting potential | $-31.1~\mathrm{mV}$ |
| k_{dj} | Rate constant of IP ₃ degradation | $0.1 \; \mathrm{s}^{-1}$ |
| G_{coup} | coupling coefficient for membrane potential | $0.5 \; \mathrm{s}^{-1}$ |
| Ca_{coup} | coupling coefficient for Ca ²⁺ | $0.05~{ m s}^{-1}$ |
| $IP_{3,coup}$ | coupling coefficient for IP ₃ | $0.05~{ m s}^{-1}$ |

Table 11: Parameters of the EC compartment, for references see Dormanns et al. [4].

9 NO pathway

9.1 ODEs

 Ca^{2+} concentration in the neuron (μ M):

$$\frac{dCa_n}{dt} = \frac{1}{1 + \lambda_{\text{buf}}} \left(\frac{I_{\text{Ca,tot}}}{2FV_{\text{spine}}} - \kappa_{\text{ex}} (Ca_n - [Ca]_{\text{rest}}) \right)$$
(178)

Activated nNOS (μ M):

$$\frac{d[\text{nNOS}]_n}{dt} = \frac{V_{\text{max,nNOS}}[\text{CaM}]_n}{K_{\text{m,nNOS}} + [\text{CaM}]_n} - \mu_{\text{deact},n}[\text{nNOS}]_n$$
(179)

we have changed some of the parameters for this nNOS equation to get a more reasonable profile see Table 12. These values give a maximum $nNOS_{act}$ of 0.2408 μM . Because of this change initial values for all state variables have been changed (only slightly) to reflect the new vaues of $nNOS_{act}$ and NO.

NO concentration in the neuron (μ M):

$$\frac{dNO_n}{dt} = p_{NO,n} - c_{NO,n} + d_{NO,n} \tag{180}$$

NO concentration in the astrocyte (μ M):

$$\frac{dNO_k}{dt} = p_{NO,k} - c_{NO,k} + d_{NO,k}$$
 (181)

NO concentration in the SMC (μ M):

$$\frac{dNO_i}{dt} = p_{NO,i} - c_{NO,i} + d_{NO,i}$$
 (182)

Activated endothelial NO synthase (eNOS) (μ M):

$$\frac{d[\text{eNOS}]_j}{dt} = \gamma_{\text{eNOS}} \frac{K_{\text{dis}} C a_j}{K_{\text{m,eNOS}} + C a_j} + (1 - \gamma_{\text{eNOS}}) g_{\text{max}} F_{\text{wss}} - \mu_{\text{deact},j} [\text{eNOS}]_j$$
(183)

NO concentration in the EC (μ M):

$$\frac{dNO_j}{dt} = p_{NO,j} - c_{NO,j} + d_{NO,j}$$
 (184)

Fraction of soluble guanylyl cyclase (sGC) in the basal state (-):

$$\frac{dE_b}{dt} = -k_1 E_b \text{NO}_i + k_{-1} E_{6c} + k_4 E_{5c}$$
(185)

Fraction of sGC in the intermediate form (-):

$$\frac{dE_{6c}}{dt} = k_1 E_b NO_i - (k_{-1} + k_2) E_{6c} - k_3 E_{6c} NO_i$$
(186)

Concentration of cGMP in the SMC (μ M):

$$\frac{dcGMP_i}{dt} = V_{\text{max,sGC}} E_{5c} - V_{\text{max,pde}} \frac{cGMP_i}{K_{\text{m,nde}} + cGMP_i}$$
(187)

9.2 Algebraic Variables

Fraction of open NR2A N-methyl-D-aspartate (NMDA) receptors (-):

$$w_{\text{NR2},A} = \frac{Glu}{K_{m,A} + Glu} \tag{188}$$

Fraction of open NR2B NMDA receptors (-):

$$w_{\text{NR2},B} = \frac{Glu}{K_{\text{m},B} + Glu} \tag{189}$$

Inward Ca²⁺ current per open NMDA receptor (fA):

$$I_{\text{Ca}} = \frac{4v_n G_{\text{M}}(P_{\text{Ca}}/P_{\text{M}})([Ca]_{\text{ex}}/[\text{M}])}{1 + \exp(\alpha_v(v_n + \beta_v))} \frac{\exp(2v_n/\phi))}{1 - \exp(2v_n/\phi))}$$
(190)

Total inward Ca²⁺ current for all open NMDA receptors per synapse (fA):

$$I_{\text{Ca,tot}} = (n_{\text{NR2},A} w_{\text{NR2},A} + n_{\text{NR2},B} w_{\text{NR2},B}) I_{\text{Ca}}$$
(191)

 Ca^{2+} -calmodulin complex concentration (μ M):

do we use Ca²⁺-calmodulin complex at all?

$$[\operatorname{CaM}]_n = \frac{Ca_n}{m_c} \tag{192}$$

Neuronal NO production flux (μ M s⁻¹):

$$p_{NO,n} = V_{\text{max,NO},n}[\text{nNOS}]_n \frac{[O_2]_n}{K_{\text{m,O2},n} + [O_2]_n} \frac{[LArg]_n}{K_{\text{m,LArg},n} + [LArg]_n}$$
(193)

Neuronal NO consumption flux (μ M s⁻¹):

$$c_{NO,n} = k_{O2,n}[NO]_n^2 [O_2]_n$$
(194)

Neuronal NO diffusive flux (μ M s⁻¹):

$$d_{NO,n} = \frac{[\text{NO}]_k - [\text{NO}]_n}{\tau_{nk}} \tag{195}$$

Time for NO to diffuse between the centres of the neuron and the astrocyte (s):

$$\tau_{nk} = \frac{x_{nk}^2}{2D_{\text{c,NO}}} \tag{196}$$

Astrocytic NO production flux (μ M s⁻¹):

$$p_{NO,k} = 0 ag{197}$$

Astrocytic NO consumption flux (μ M s⁻¹):

$$c_{NO,k} = k_{O2,k} [NO]_k^2 [O_2]_k$$
(198)

Astrocytic NO diffusive flux (μ M s⁻¹):

$$d_{NO,k} = \frac{[NO]_n - [NO]_k}{\tau_{nk}} + \frac{[NO]_i - [NO]_k}{\tau_{ki}}$$
(199)

Time for NO to diffuse between the centres of the astrocyte and the SMC (s):

$$\tau_{ki} = \frac{x_{ki}^2}{2D_{\text{c,NO}}} \tag{200}$$

SMC NO production flux (μ M s⁻¹):

$$p_{NO,i} = 0 (201)$$

SMC NO consumption flux (μ M s⁻¹):

$$c_{NO,i} = k_{\text{dno}}[\text{NO}]_i \tag{202}$$

SMC NO diffusive flux (μ M s⁻¹):

$$d_{NO,i} = \frac{[NO]_k - [NO]_i}{\tau_{ki}} + \frac{[NO]_j - [NO]_i}{\tau_{ij}}$$
(203)

sGC kinetics rate constant (s^{-1}):

$$k_4 = C_4 [\text{cGMP}]_i^2 \tag{204}$$

Fraction of sGC in the fully activated form (-):

$$E_{5c} = 1 - E_b - E_{6c} (205)$$

Regulatory effect of cGMP on myosin dephosphorylation (-):

$$R_{\text{cGMP}} = \frac{[\text{cGMP}]_i^2}{K_{\text{m,mlcp}}^2 + [\text{cGMP}]_i^2}$$
(206)

Maximum cGMP production rate (μ M s⁻¹):

$$V_{\text{max,pde}} = k_{\text{pde}}[\text{cGMP}]_i \tag{207}$$

Time for NO to diffuse between the centres of the SMC and the EC (s):

$$\tau_{ij} = \frac{x_{ij}^2}{2D_{\text{c,NO}}} \tag{208}$$

Fraction of the elastic strain energy stored within the membrane (-):

$$F_{\text{wss}} = \frac{1}{1 + \alpha_{\text{wss}} \exp(-W_{\text{wss}})} - \frac{1}{1 + \alpha_{\text{wss}}}$$
(209)

Strain energy density (-):

$$W_{\text{wss}} = W_0 \frac{(\tau_{\text{wss}} + \sqrt{16\delta_{\text{wss}}^2 + \tau_{\text{wss}}^2 - 4\delta_{\text{wss}})^2}}{\tau_{\text{wss}} + \sqrt{16\delta_{\text{wss}}^2 + \tau_{\text{wss}}^2}}$$
(210)

Wall shear stress (Pa):

$$\tau_{\text{wss}} = \frac{R\Delta P}{2L} \tag{211}$$

 O_2 concentration in the EC (μ M):

$$[O_2]_j = c_{unit} O_2 \tag{212}$$

Oxygen in EC taken as O_2 from lumen (diffusion very fast so plausible!) instead of constant, in uM, hence $c_{unit} = 1$. EC NO production flux (μ M s⁻¹):

$$p_{NO,j} = V_{\text{max,NO},j}[\text{eNOS}]_j \frac{[O_2]_j}{K_{\text{m,O2},j} + [O_2]_j} \frac{[LArg]_j}{K_{\text{m,L-Arg},j} + [LArg]_j}$$
(213)

EC NO consumption flux (μ M s⁻¹):

$$c_{NO,j} = k_{O2,j}[NO]_j^2[O_2]_j$$
 (214)

EC NO diffusive flux (μ M s⁻¹):

$$d_{NO,j} = \frac{[\text{NO}]_i - [\text{NO}]_j}{\tau_{ij}} - \frac{4D_{\text{c,NO}}[\text{NO}]_j}{r_l^2}$$
(215)

| Parameter | Description | Value |
|--------------------------|---|--|
| [Glu] _{max} | Maximum glutamate concentration | $1846~\mu\mathrm{M}$ |
| $\lambda_{ m buf}$ | Buffer capacity | 20 |
| $V_{ m spine}$ | Dendritic spine volume | $8 	imes 10^{-5} \ \mathrm{pL}$ |
| $\kappa_{ m ex}$ | Decay rate constant of internal Ca ²⁺ concentration | $1.6 \times 10^3 \; \mathrm{s}^{-1}$ |
| $[Ca]_{rest}$ | Resting internal Ca ²⁺ concentration | $0.1~\mu\mathrm{M}$ |
| $V_{ m max,nNOS}$ | Maximum nNOS activation rate | $0.7 \times 10^{-3} \ \mu \text{M s}^{-1} \text{ new val}$ |
| • | | ues |
| $K_{\mathrm{m.nNOS}}$ | Michaelis constant | $0.8 \mu M$ new values |
| $\mu_{\mathrm{deact},n}$ | Rate constant at which nNOS is deactivated | 0.02 s^{-1} new values |
| $K_{m,A}$ | Michaelis constant | $650~\mu\mathrm{M}$ |
| $K_{m,B}$ | Michaelis constant | $2800~\mu\mathrm{M}$ |
| v_n | Neuronal membrane potential | $-40~\mathrm{mV}$ |
| $G_{ m M}$ | Conductance of NMDA receptor | 0.46 pS new value |
| $P_{\rm Ca}/P_{\rm M}$ | Ratio of Ca ²⁺ permeability to monovalent ion permeability | 3.6 |
| $[Ca]_{\text{ex}}$ | External Ca ²⁺ concentration | $2 \times 10^3 \ \mu M$ |
| [M] | Concentration of monovalent ions | $1.3 \times 10^5 \mu\text{M}$ |
| α_v | Voltage-dependent Mg ²⁺ block parameter | $-0.08 \mathrm{mV}^{-1}$ |
| eta_v | Voltage-dependent Mg ²⁺ block parameter | 20 mV |
| | Average number of NR2A NMDA receptors | 0.63 |
| $n_{\text{NR2},A}$ | Average number of NR2B NMDA receptors Average number of NR2B NMDA receptors | 11 |
| $n_{\mathrm{NR2},B}$ | Number of Ca ²⁺ ions bound per calmodulin | 4 |
| m_c | Maximum catalytic rate of neuronal NO production | 4.22 s^{-1} |
| $V_{\max,NO,n}$ | O ₂ concentration in the neuron | $200 \mu M$ |
| $[O_2]_n$ | | , |
| $K_{\mathrm{m,O2},n}$ | Michaelis constant for nNOS for O ₂ | 243 μM |
| $[LArg]_n$ | L-Arg concentration in the neuron | $100 \mu \text{M}$ |
| $K_{m,LArg,n}$ | Michaelis constant for nNOS for LArg | 1.5 μ M |
| $k_{\mathrm{O2},n}$ | O ₂ reaction rate constant | $9.6 \times 10^{-6} \ \mu M^{-2} \ s^{-1}$ |
| x_{nk} | Distance between centres of neuron and astrocyte | $25 \mu \text{m}$ |
| $k_{\text{O2},k}$ | O ₂ reaction rate constant | $9.6 \times 10^{-6} \mu \text{M}^{-2} \text{s}^{-1}$ |
| $[O_2]_k$ | O ₂ concentration in the astrocyte | $200~\mu\mathrm{M}$ |
| x_{ki} | Distance between centres of astrocyte and SMC compartments | $25 \mu\mathrm{m}$ |
| k_{-1} | sGC kinetics rate constant | 100 s^{-1} |
| k_1 | sGC kinetics rate constant | $2 \times 10^{3} \ \mu \text{M}^{-1} \ \text{s}^{-1}$ |
| k_2 | sGC kinetics rate constant | $0.1 \mathrm{s}^{-1}$ |
| k_3 | sGC kinetics rate constant | $3~\mu\mathrm{M}^{-1}~\mathrm{s}^{-1}$ |
| $V_{ m max,sGC}$ | Maximal cGMP production rate | $0.8520~\mu{ m M~s^{-1}}$ |
| $K_{ m m,pde}$ | Michaelis constant | $2~\mu\mathrm{M}$ |
| $k_{ m dno}$ | Constant reflecting the activity of various NO scavengers | $0.01 \ s^{-1}$ |
| C_4 | sGC rate scaling constant | $0.011~\mu\mathrm{M}^{-2}~\mathrm{s}^{-1}$ |
| $K_{ m m,mlcp}$ | Hill coefficient | $5.5~\mu\mathrm{M}$ |
| $k_{\rm pde}$ | Phosphodiesterase rate constant | 0.0195 s^{-1} |
| x_{ij} | Distance between centres of SMC and EC compartments | $3.75~\mu\mathrm{m}$ |
| $\gamma_{ m eNOS}$ | Relative strength of the Ca ²⁺ dependent pathway for eNOS activation | 0.1 |
| $\mu_{\mathrm{deact},j}$ | eNOS-caveolin association rate | 0.0167 s^{-1} |
| $K_{ m dis}$ | eNOS-caveolin disassociation rate | $0.09~\mu{ m M}~{ m s}^{-1}$ |

| $K_{ m m,eNOS}$ | Michaelis constant | $0.45~\mu\mathrm{M}$ |
|----------------------------|---|--|
| $g_{ m max}$ | Maximum wall-shear-stress-induced eNOS activation | $0.06~\mu{ m M}~{ m s}^{-1}$ |
| $lpha_{ m wss}$ | Zero shear open channel constant | 2 |
| W_0 | Shear gating constant | $1.4 \ Pa^{-1}$ |
| $\delta_{ m wss}$ | Membrane shear modulus | 2.86 Pa |
| $V_{\max, \mathrm{NO}, j}$ | Maximum catalytic rate of NO production | $1.22 \; \mathrm{s}^{-1}$ |
| $K_{m,O2,j}$ | Michaelis constant for eNOS for O ₂ | $7.7~\mu\mathrm{M}$ |
| $[LArg]_j$ | L-Arg concentration in the neuron | $100 \ \mu M$ |
| $K_{m,L-Arg,j}$ | Michaelis constant for L-Arg | $1.5~\mu\mathrm{M}$ |
| $\Delta P/L$ | Pressure drop over length of arteriole | $9.1 	imes 10^{-2}$ Pa μ m $^{-1}$ |
| $k_{\mathrm{O2},j}$ | O ₂ reaction rate constant | $9.6 	imes 10^{-6}~\mu\mathrm{M}^{-2}~\mathrm{s}^{-1}$ |
| $D_{ m c,NO}$ | NO diffusion coefficient | $3300~\mu\mathrm{m}^{~2}\mathrm{s}^{-1}$ |
| r_l | Constant of lumen radius | $25~\mu\mathrm{m}$ |

Table 12: Parameters of the NO submodel, for references see Dormanns et al. [3].

10 AA 20-HETE NO pathway

In our previous model we noted that the rate of increase of $nNOS_{act}$ and NO was linear and reached a peak at 3 seconds after stimulation. we found a mistake in the NMDA receptor conductance value g_{Ca} (in the code). This allowed for Ca_{neuron} to be about $6\mu M$ which is similar to that found by [21] for 10 receptors per post-synapse. The new values (in red in Table 12) provide a maximum of $nNOS_{act} = 0.25 \mu M$ and a maximum of $NO = 0.13 \mu M$.

Our results have so far indicated that the NO-cGMP pathway is very slow (even when using experimental data from [28]). This has led to a reconsideration of the NO model and an investigation into the relationship between NO, as a vasodilator and 20-HETE as a vasoconstrictor. Liu et al [12] pprovides an hypothesis on the relationship between NO and 20-HETE and EETs in neurovascular coupling. This is illustrated in Figure 17. The key for vasoconstriction is the modulation (inhibition) of the K_{Ca} channel by 20-HETE. Whilst in contrast NO in addition to the cGMP pathway inhibits the production of 20-HETE from arachidonic acid (AA) via the CYP4A enzyme. In this hypothesis AA diffuses to the SMC from the astrocyte. Using the model of Haffield et al [6] we can write conservation equations for AA and 20-HETE as shown below.

$$[AA] = [AA]_b + \frac{AA_{max}\Delta[Ca^{2+}]}{AA_M}$$
(216)

$$[AA] = [AA]_b + \frac{AA_{max}\Delta[Ca^{2+}]}{AA_M}$$

$$\frac{d}{dt}[20 - HETE] = f_{NO}\frac{V_{A11}[AA]}{K_{A11} + [AA]} + \frac{V_{F2}[AA]}{K_{F2} + [AA]} - \mu[20 - HETE]$$
(216)

$$f_{NO} = \frac{1}{1 + exp(\frac{NO_i - NO_{rest}}{R_{NO}})}$$
(218)

The first RHS two terms are based on the fact that both CYPA2 and CYPF2 enzymes contribute to the 20-HETE production. Nitric Oxide degrades the CYPA2 enzyme and hence the function f_{NO} provides a phenomenological way of expressing this degradation. The parameter values are given in the Table below

| Parameter | Description | Value |
|------------|--------------------------|---------------|
| $[AA]_b$ | archidonic acid baseline | $9.3~\mu m$ |
| AA_M | Michaelis constant | $0.161~\mu M$ |
| AA_{max} | AA max reaction rate | $29~\mu M$ |

| V_{A11} | CYPA 20-HETE max reaction rate | $0.212 \ s^{-1}$ |
|-------------|--|--------------------|
| K_{A11} | CYPA Michaelis constant | $228~\mu M$ |
| V_{F2} | CYPF 20-HETE max reaction rate | $0.0319 \ s^{-1}$ |
| K_{F2} | CYPF Michaelis constant | $23.5~\mu M$ |
| μ | 20-HETE decay constant | $0.139 \ s^{-1}$ |
| D_{AA} | Diffusion coefficient of AA ¹ | $10^{-8}m^2s^{-1}$ |
| NO_{rest} | NO resting concentration for f_{NO} | $0.02047~\mu M$ |
| R_{NO} | scaling parameter for f_{NO} | $0.02 \ s^{-1}$ |

Table 13: parameter values for the production of archidonic acid and 20-HETE.

11 Wall mechanics

11.1 **ODEs**

Fraction of free phosphorylated cross-bridges (-):

$$\frac{d[Mp]}{dt} = \chi_w \left(K_4[AMp] + K_1[M] - (K_2 + K_3)[Mp] \right)$$
 (219)

Fraction of attached phosphorylated cross-bridges (-):

$$\frac{d[AMp]}{dt} = \chi_w \left(K_3[Mp] + K_6[AM] - (K_4 + K_5)[AMp] \right)$$
 (220)

Fraction of attached dephosphorylated cross-bridges (-):

$$\frac{d[AM]}{dt} = \chi_w \left(K_5 [AMp] - (K_7 + K_6) [AM] \right) \tag{221}$$

Vessel radius (μ m):

$$\frac{dR}{dt} = \frac{R_{init}}{\eta} \left(\frac{RP_T}{h} - E \frac{R - R_0}{R_0} \right) \tag{222}$$

11.2 Algebraic Variables

Fraction of free non-phosphorylated cross-bridges (-):

$$[M] = 1 - [AMp] - [Mp]$$
(223)

Rate constants for phosphorylation of M to Mp and of AM to AMp (s^{-1}) :

$$K_1 = K_6 = \gamma_{cross} C a_i^{n_{cross}}$$
(224)

Rate constants for dephosphorylation of Mp to M and of AMp to AM (s^{-1}) :

$$K_2 = K_5 = \delta_K \left(k_{\text{mlpc,b}} + k_{\text{mlpc,c}} R_{\text{cGMP}} \right) \tag{225}$$

Wall thickness of the vessel (μ m):

$$h = 0.1R \tag{226}$$

Fraction of attached myosin cross-bridges (-):

$$F_r = [AM_p] + [AM] \tag{227}$$

¹Kong et al Fluid Phase Equilibria, **297**(3)pp 162-167

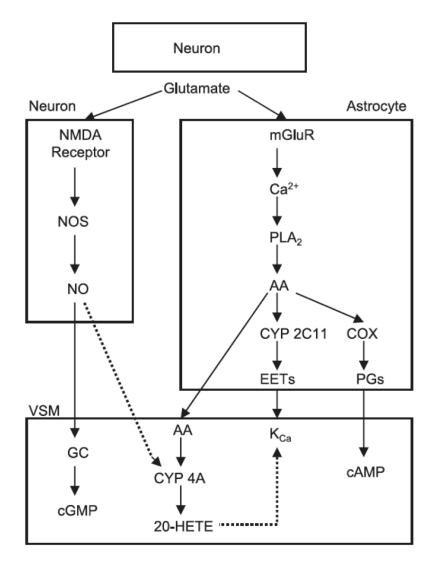


Figure 17: proposed pathway sketch by Liu et al [12] of 20-HETE and NO. Dashed lines are inhibition

Young's modulus (Pa):

$$E = E_{pas} + F_r \left(E_{act} - E_{pas} \right) \tag{228}$$

Initial radius (μ m):

$$R_0 = R_{init} + F_r(\alpha_R - 1)R_{init} \tag{229}$$

| Parameter | Description | Value |
|-----------------------|--|--|
| χ_w | Scaling constant for wall mechanics | 1.7 |
| K_3 | Rate constant for attachment of phosphorylated crossbridges | $0.4 \ { m s}^{-1}$ |
| K_4 | Rate constant for detachment of phosphorylated crossbridges | 0.1 s^{-1} |
| K_7 | Rate constant for detachment of dephosphorylated crossbridges | 0.1 s^{-1} |
| γ_{cross} | Sensitivity of the contractile apparatus to Ca ²⁺ | $17~\mu\mathrm{M}^{-3}\mathrm{s}^{-1}$ |
| n_{cross} | Fraction constant of the phosphorylation crossbridge | 3 |
| δ_K | Constant to fit data | 58.14 |
| $k_{\rm mlpc,b}$ | Basal MLC dephosphorylation rate constant | $8.6 \times 10^{-3} \text{ s}^{-1}$ |
| $k_{\mathrm{mlpc,c}}$ | First-order rate constant for cGMP regulated MLC dephosphorylation | $32.7 \times 10^{-3} \text{ s}^{-1}$ |

| η | Viscosity | 10^4 Pa s |
|------------|---------------------------------------|--------------------------------|
| P_T | Transmural pressure | 4×10^3 Pa |
| E_{pas} | Young's moduli for the passive vessel | 66×10^3 Pa |
| E_{act} | Young's moduli for the active vessel | $233 \times 10^{3} \text{ Pa}$ |
| α_R | Scaling factor for initial radius | 0.6 |

Table 14: Parameters of the wall mechanics submodel, for references see Dormanns et al. [4].

12 Tissue Slice Model

12.1 ODEs

 K^+ concentration in the astrocyte of NVU block i with four neighbours j (μ M):

$$\frac{dK_k^i}{dt} = -\frac{1}{\Delta x} \sum_{i} J_{K,i\to j} - J_{K_k}^i + 2J_{NaK_k}^i + J_{NKCC1_k}^i + J_{KCC1_k}^i$$
 (230)

Membrane potential in the astrocyte of NVU block i with four neighbours j (mV):

$$\frac{dv_k^i}{dt} = \gamma_v \left[-\frac{1}{\Delta x} \sum_j z_K J_{K,i\to j} - J_{BK_k}^i - J_{K_k}^i - J_{Cl_k}^i - J_{NBC_k}^i - J_{Na_k}^i - J_{NaK_k}^i - 2J_{TRPV_k}^i \right]$$
(231)

 K^+ concentration in the ECS of NVU block i with four neighbours j (mM):

$$\frac{dK_e^i}{dt} = -\frac{1}{\Delta x} \sum_j J_{K,i \to j}^e + \frac{1}{Ff_e} \left(\frac{A_s I_{K,tot_{sa}}^i}{V_s} + \frac{A_d I_{K,tot_d}^i}{V_d} \right) - \frac{d\mathbf{Buff}_e^i}{dt}$$
(232)

 Na^+ concentration in the ECS of NVU block i with four neighbours j (mM):

$$\frac{dNa_e^i}{dt} = -\frac{1}{\Delta x} \sum_j J_{Na,i \to j}^e + \frac{1}{Ff_e} \left(\frac{A_s I_{Na,tot_{sa}}^i}{V_s} + \frac{A_d I_{Na,tot_d}^i}{V_d} \right)$$
(233)

12.2 Algebraic Variables

Gap junction flux of K_k from NVU block i to neighbour j (μ Mm s⁻¹):

$$J_{K,i\to j} = -\frac{D_{gap}}{\Delta x^2} \left((K_k^j - K_K^i) + \frac{z_K F}{RT} \frac{K_k^i + K_k^j}{2} (v_k^j - v_k^i) \right)$$
(234)

Extracellular electrodiffusive flux of K_e and Na_e from NVU block i to neighbour j (mMm s⁻¹):

$$J_{K,i\to j}^{e} = -\frac{D_{K,e}}{\Delta x} \left[(K_e^j - K_e^i) - z_K \left(\frac{K_e^i + K_e^j}{2} \right) \left(\frac{z_K D_{K,e} (K_e^j - K_e^i) + z_{Na} D_{Na,e} (N a_e^j - N a_e^i)}{z_K^2 D_{K,e} \frac{K_e^i + K_e^j}{2} + z_{Na}^2 D_{Na,e} \frac{N a_e^i + N a_e^j}{2}} \right) \right]$$
(235)

$$J_{Na,i\to j}^{e} = -\frac{D_{Na,e}}{\Delta x} \left[(Na_{e}^{j} - Na_{e}^{i}) - z_{Na} \left(\frac{Na_{e}^{i} + Na_{e}^{j}}{2} \right) \left(\frac{z_{K}D_{K,e}(K_{e}^{j} - K_{e}^{i}) + z_{Na}D_{Na,e}(Na_{e}^{j} - Na_{e}^{i})}{z_{K}^{2}D_{K,e} \frac{K_{e}^{i} + K_{e}^{j}}{2} + z_{Na}^{2}D_{Na,e} \frac{Na_{e}^{i} + Na_{e}^{j}}{2} \right) \right]$$
(236)

| Parameter | Description | Value |
|--------------|---|---|
| D_{gap} | Astrocytic gap junction diffusion coefficient | $3.1 \times 10^{-9} \text{ m}^2 \text{ s}^{-1}$ |
| Δx^2 | Width of one NVU block | $1.24 \times 10^{-4} \mathrm{m}$ |
| $D_{K,e}$ | Extracellular K ⁺ diffusion coefficient | $3.8 \times 10^{-9} \; \mathrm{m^2 \; s^{-1}}$ |
| $D_{Na,e}$ | Extracellular Na ⁺ diffusion coefficient | $2.5 \times 10^{-9} \ \mathrm{m^2 \ s^{-1}}$ |

Table 15: Parameters of the large scale tissue slice model.

13 Initial values

These initial values take into account the new $nNOS_{act_j}$ and NO equations. All concentrations are in micromolar and membrane potential in millivolts.

13.1 Neuron

 $E_t = 0$; $I_t = 0$; $K_e = 3.5$; $Na_{sa} = 9.37$; $Na_d = 9.42$; O2 = 0.02566; CBV = 1.204; HbR = 0.7641; $Ca_n = 0.1$; $nNOS_{act_n} = 0.01056$; $NO_n = 0.02425$;

13.2 Astrocyte

 $Na_k = 18740; K_k = 92660; HCO_{3_k} = 9085; Cl_k = 8212; Na_s = 149200; K_s = 2932; HCO_{3_s} = 16980; K_p = 3039; w_k = 8.26e-5; Ca_k = 0.1435; s_k = 480.8; h_k = 0.4107; I_k = 0.048299; eet_k = 0.4350; m_k = 0.513; Ca_p = 1853; NO_k = 0.02234; v_k = -88.79;$

13.3 **SMC/EC**

 $Ca_i = 0.2641$; $s_i = 1.1686$; $v_i = -34.7$; $w_i = 0.2206$; $I_i = 0.275$; $K_i = 99994.8$; $NO_i = 0.02047$; $E_b = 0.6372$; $E_6c = 0.2606$; $cGMP_i = 6.1$; $Ca_j = 0.8339$; $s_j = 0.6262$; $v_j = -68.39$; $I_j = 0.825$; $eNOS_{act_j} = 0.4451$; $NO_j = 0.02051$;

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