

Analysing High-Dimensional Neuroscience Models: Neurovascular Coupling.

Tim David¹, Pierre Gremaud², Joey Hart²

1.Department of Mechanical Engineering,

University of Canterbury, New Zealand

2.Department of Mathematics,

North Carolina State University, Raleigh, NC

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Abstract

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1 INTRODUCTION

During the last two decades functional magnetic resonance imaging (fMRI) has proven to be an established tool in studying the human brain. This is especially true in the case of the blood-oxygen-level dependent (BOLD) signal, where changes in blood oxygen levels can

be detected via the magnetic signal [17]. However due to the constraint on the resolution of BOLD, fMRI methodology has not been used extensively to study the underlying cellular neural architecture and their associated cerebral functions. Complex models that address this important relationship and constructing a detailed compartmental model with the relevant cell types involved will allow simulations relating certain brain functions performed in a region to its fMRI BOLD response. The neurovascular coupling (NVC) mechanism, the cerebral metabolic rate of oxygen consumption, and the cerebral blood volume (CBV) are known to contribute to the fMRI BOLD response [3], however a thorough understanding of these factors has yet to be fully established.

The NVC response, the ability to locally adjust vascular resistance as a function of neuronal activity, is believed to be mediated by a number of different signalling mechanisms. Roy and Sherrington [19] first proposed a mechanism based on a metabolic negative feedback theory. According to this theory, neural activity leads to a drop in oxygen or glucose levels and increases in CO_2 , adenosine, and lactate levels. All of these signals could dilate arterioles and hence were believed to be part of the neurovascular response. However, recent experiments illustrated that the NVC response is partially independent of these metabolic signals [10, 11, 16, 18, 13]. An alternative to this theory was proposed where the neuron releases signalling molecules to directly or indirectly affect the blood flow. Many mechanisms such as the potassium (K^+) signalling mechanism [8], the nitric oxide (NO) signalling mechanism or the arachidonic acid to epoxyeicosatrienoic acid (EET) pathway are found to contribute to the neurovascular response [2].

The K^+ signalling mechanism of NVC seems to be supported by significant evidence, although new evidence shows that the endfoot astrocytic calcium (Ca^{2+}) could play a significant role. The K^+ signalling hypothesis mainly utilises the astrocyte, positioned to enable the communication between the neurons and the local perfusing blood vessels. The astrocyte and the endothelial cells (ECs) surrounding the perfusing vessel lumen exhibit a striking similarity in ion channel expression and thus can enable control of the smooth muscle cell (SMC) from both the neuronal and blood vessel components [12]. Whenever there is neuronal activation K^+ ions are released into the extracellular space (ECS) and synaptic cleft (SC). The astrocyte is depolarised by taking up K^+ released by the neuron and releases it into the perivascular space (PVS) via the endfeet through the BK channels [7]. This increase in ECS K^+ concentration ($3 - 10$ mM) near the arteriole hyperpolarises the SMC through the inward rectifying K^+ (KIR) channel, effectively closing the voltage-gated Ca^{2+} channel, reducing smooth muscle cytosolic Ca^{2+} and thereby causing dilation. Higher K^+ concentrations in the PVS cause contraction due to the reverse flux of the KIR channel [6].

Amidst the difficulty in monitoring and measuring the rapid changes in metabolic demands in the highly heterogeneous brain, speculative estimates of the relative demands of the cerebral processes that require energy were given based on different experimental data by Ames [1]. As per the estimate, the vegetative processes that maintain the homeostasis including protein synthesis accounted for $10 - 15\%$ of the total energy consumption. The costliest function seems to be in restoring the ionic gradients during neural activation. The sodium potassium (Na^+/K^+) exchange pump is estimated to consume $40 - 50\%$, while the

Ca^{2+} influx from organelles and extracellular fluid consumes 3 – 7%. The processing of neurotransmitters such as uptake or synthesis consumes 10 – 20%, while the intracellular signalling systems which includes activation and inactivation of proteins consumes 20 – 30%. The rest of the energy is estimated to be consumed by the axonal and dendritic transport in both directions.

Previous work [?] has provided the construction of an experimentally validated numerical (*in silico*) model based on experimental data to simulate the fMRI BOLD signal associated with NVC along with the associated metabolic and blood volume responses. An existing neuron model [14, 15] has been extended to include an additional transient sodium (Na^+) ion channel (NaT) expressed in the neuron, and integrated into a complex NVC model [5, 4, 9]. This present model is based on the hypothesis that the K^+ signalling mechanism of NVC is the primary contributor to the vascular response and the Na^+/K^+ exchange pump in the neuron is the primary consumer of oxygen during neural activation. The model contains 160 parameters, most of which come from non-human experiments.

Such a complex model constructed with a high-dimensional parameter space is not easily amenable to sensitivity analyses considering the significant computing resource required. Indeed no formal theory exists which allows direct mathematical investigation of the variability of the large dimensional parameter vector and the resulting output. From a purely physiological perspective an understanding of the dominant cellular mechanisms resulting in cerebral tissue perfusion after neuronal stimulation would be of particular interest.

We have used the cerebral blood flow (CBF) change from the experimental data [20]

taken from the rat barrel cortex.

INTRODUCE NOTATION AND TERMINOLOGY WE CAN REFER TOO THROUGH-
OUT THE ARTICLE

2 Methodology

The ODE system is solved using ode15s in Matlab with relative and absolute tolerances of 10^{-4} .

2.1 QoIs

We analyse 3 quantities of interest (QoIs) with respect to two different stimulations, a 10 second square stimulation pulse and a 16 second stimulus used in lab experiments. They are displayed in Figure 1. Assuming the stimulation occurs for $t_1 \leq t \leq t_2$, the QoIs are defined

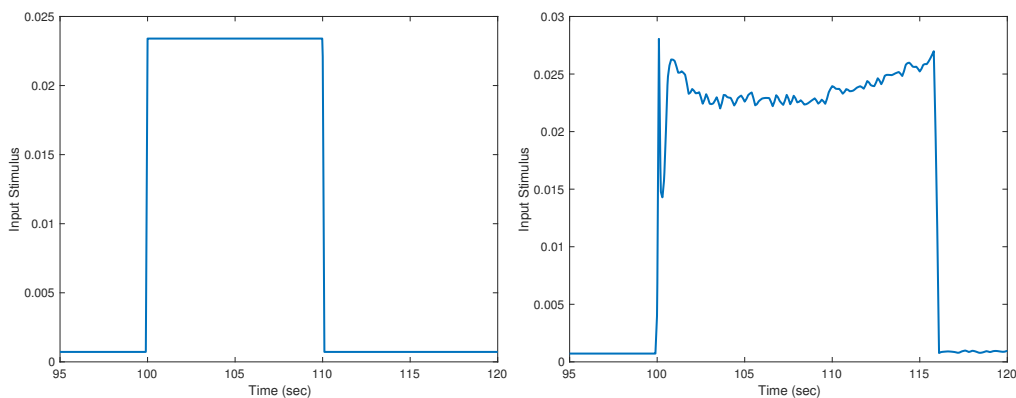


Figure 1: Left: rectangular pulse input stimulus; right: stimulus used in lab experiments.

as follows.

1. ECS potassium has a distinct effect on the flux into the Neuron. Hence we look at the average.

$$\frac{1}{t_2 - t_1} \int_{t_1}^{t_2} [K^+]_{ECS}(s) ds \quad (2.1)$$

2. As a representation of the volumetric flow rate in the cerebral tissue

$$\frac{1}{t_2 - t_1} \int_{t_1}^{t_2} \left(\frac{R(s)}{R_0} \right)^4 ds \quad (2.2)$$

3. The combined concentration of the actin myosin complex, both phosphorylated and unphosphorylated, determines the effect stress due to the contraction for the smooth muscle cell.

$$[AM + AM_p]_{min} \quad (2.3)$$

In addition, we consider the average concentration of AC potassium. It exhibits very small variation with respect to the parameters so we do not report any additional results for this QoI.

2.2 Surrogate models and Sobol' indices

DESCRIBE THE SURROGATES AND SOBOL' INDICES

3 Results

This section details our approach to varying the parameters and contains numerical results for the 3 QoI's, with both the square pulse stimulus and the experimental data stimulus for

each. The section is organized in four subsections, one on the parameter sampling and one for each of the three QoI's.

3.1 Parameter sampling

Distributions for the parameters are not known a priori, nor are they able to be estimated from experimental data. The subsection below details our computational approach to determine a distribution for the parameters. The computation time required for model evaluations with a stimulus applied is too great for the computational approach below. Rather, we focus on properties of the solution when no stimulus is applied, i.e. the steady state solution(s). This affords the needed computational acceleration.

Parameter distributions are determined by first assuming independence of the parameters and assigning each parameter a uniform distribution with $\pm 10\%$ uncertainty around its nominal value (larger amounts of uncertainty may be desirable but are not computationally feasible). Parameters are likely to be correlated but we do not know the correlation structure a priori. The model is run, with no stimulus applied, for 919 different parameter samples. Of these 919 runs, 670 of them fail due to ode15s terminating prematurely because its time steps became too small. The remaining 249 yields 139 solutions which the radius reaches a stable steady state (at least for the duration of the time integration) and 110 solutions for which the radius reaches a steady state but subsequently because transient. The left panel of Figure 2 displays four representative solutions for the radius; the red curves remain in steady state for the duration of our time integration whereas the black curves revert

into a transit regime after some time in or near steady state. The right panel of Figure 2 displays the samples for two parameters (in the buffer equation) which are highly influential in determining the behavior of the solution. A blue * indicates a sample where the solver terminated prematurely, a yellow + indicates a sample where an unstable steady state was observed, a red o indicates a sample where the solution remained in steady state. A very clear (and strong) correlation is observed between these parameters which has significant influence on the behavior of the solution.

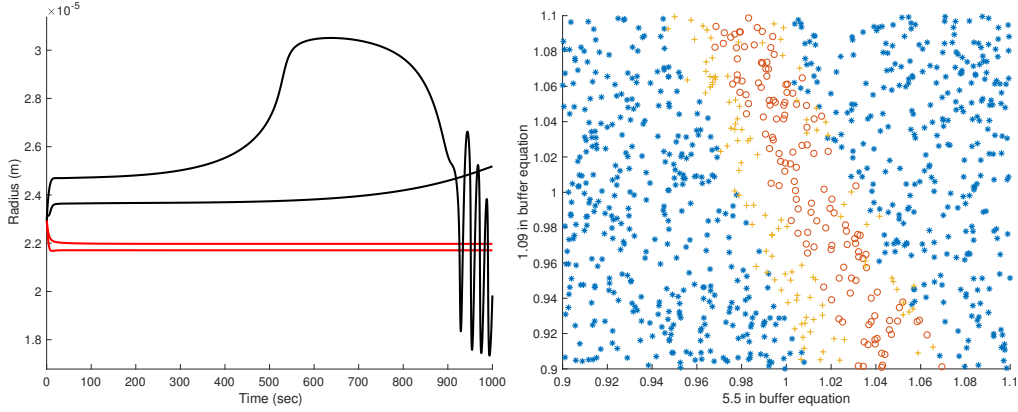


Figure 2: Left: examples of stable (red) and unstable (black) steady state solutions. Right: samples of the buffer parameters using uniform independent sampling. A blue * indicates the sample yielded a premature termination of the solver, a yellow + indicates the sample yielded an unstable steady state, a red o indicates the sample yielded a stable steady state.

Observing this correlation, we use the samples for which the radius remained in steady state to fit the two buffer parameters with a bivariate distribution with beta marginals and a Frank copula. The experiment was repeated by sampling the two correlated parameters with the bivariate distribution and all other parameters from their original uniform distributions.

The samples yielding steady state solutions were collected, a bivariate distribution fit to them, and the experiment was repeated. After three additional iterations we were able to generate 902 out of 960 samples which yielded solutions with stable steady states (51 solutions had unstable steady states and 7 had premature solver terminations). This distribution is used for all subsequent analysis.

Samples are drawn and the model, with a stimulus applied (in two separate cases, the 10 second rectangular pulse and the 16 second stimulus from experimental data), is run for each sample. This results in solutions exhibiting three different physiological regimes; they are displayed in Figure 3 where the radius is plotted as a function of time. The leftmost panel corresponds to the typical case when the radius increases in response to the stimulus and then decreases when it is removed; the center panel corresponds to an atypical case where the radius has an initial decrease in response to the stimulus; the right panel corresponds to another atypical case where the radius reaches another steady state and does not decrease after the stimulus is removed.

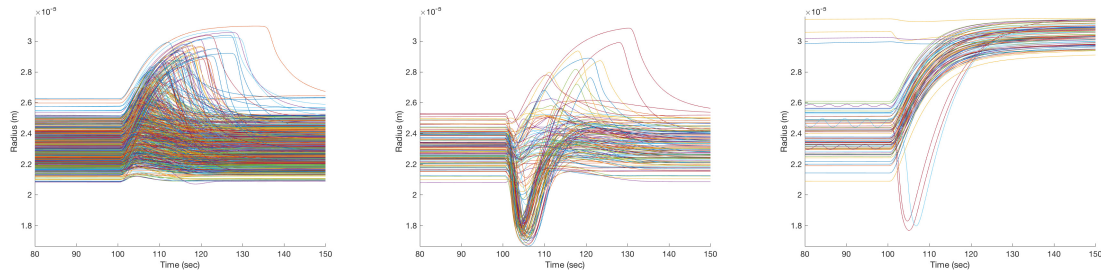


Figure 3: Radii corresponding to samples (using the rectangular pulse stimulus). Left: curves showing an increase in response to the stimulus; center: curves with a decrease in response to the stimulus; right: curves which settle in a different steady state.

This study focus on the typical case so we remove samples where the radius does not increase in response to the stimulus and decrease when it is removed. This processing yields 660 samples for analysis when the rectangular pulse stimulus is applied and 438 samples when the stimulus from experimental data is applied. The results presented below use these samples.

ADD A DISCUSSION OF WHICH PARAMETERS ARE MOST INFLUENTIAL IN DETERMINING THE SOLUTION REGIME

3.2 QoI (1) (K_{ECS} Mean)

ADD AN INTERPRETATION OF THE FIGURE AND TABLE

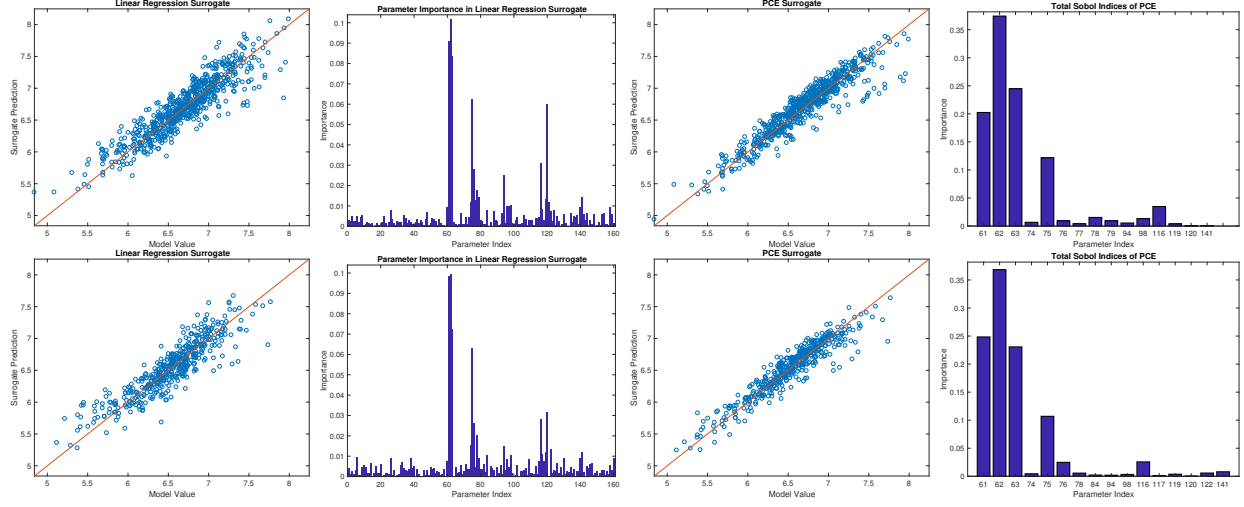


Figure 4: Top row: results with a rectangular pulse stimulus; bottom row: results with experimental data stimulus. From left to right, linear regression predictions, linear regression variable importance, PCE predictions, total Sobol' indices for PCE.

Index	Identification	Total Sobol' Index (RP)	Total Sobol' Index (ES)
62	0.143 in m4alpha and m4 beta	0.3744	0.3683
63	5.67 in m4alpha and m4 beta	0.2446	0.2307
61	gKleak_d in Neuron	0.2025	0.2485
75	34.9 in m6alpha	0.1217	0.1070
116	dhod in Neuron	0.0348	0.0257

Table 1: K_{ECS} Mean

3.3 QoI (2.2) (volumetric flow rate)

ADD AN INTERPRETATION OF THE FIGURE AND TABLE

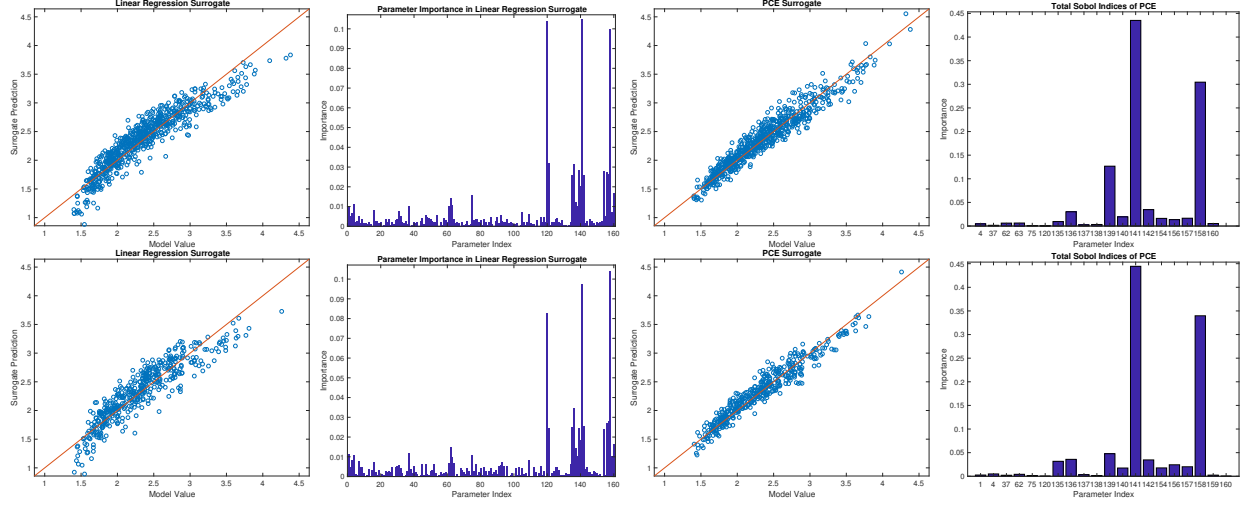


Figure 5: Top row: results with a rectangular pulse stimulus; bottom row: results with experimental data stimulus. From left to right, linear regression predictions, linear regression variable importance, PCE predictions, total Sobol' indices for PCE.

Index	Identification	Total Sobol' Index (RP)	Total Sobol' Index (ES)
141	z_4 in SMCEC	0.4353	0.4445
158	n_cross in WallMechanics	0.3045	0.3398
139	z_2 in SMCEC	0.1267	0.0478
142	z_5 in SMCEC	0.0347	0.0347
136	G_K_i in SMCEC	0.0302	0.0359

Table 2: volumetric flow rate

3.4 QoI (2.3) ($AM + AM_p$ Min)

ADD AN INTERPRETATION OF THE FIGURE AND TABLE

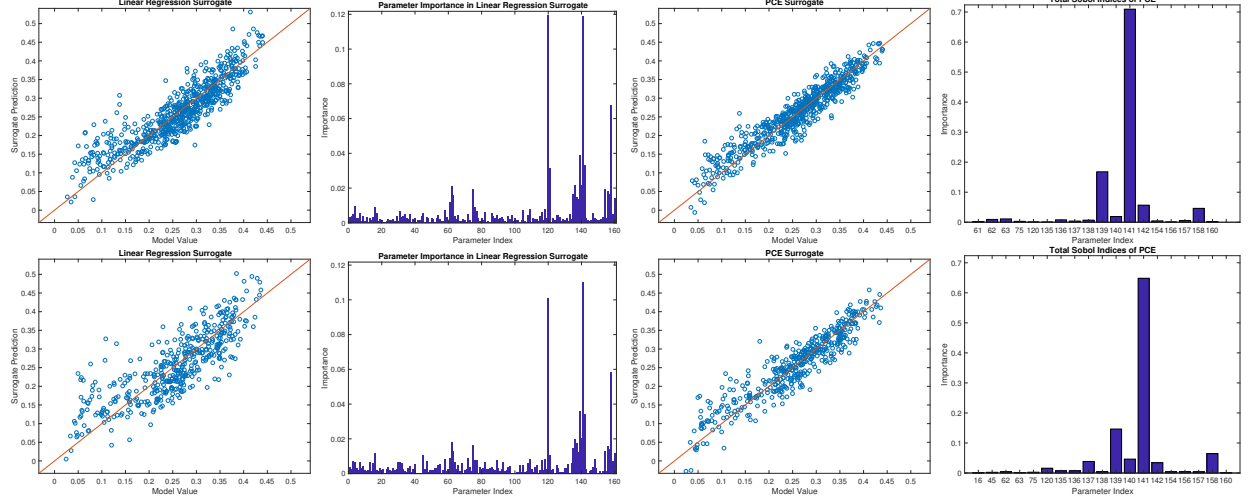


Figure 6: Top row: results with a rectangular pulse stimulus; bottom row: results with experimental data stimulus. From left to right, linear regression predictions, linear regression variable importance, PCE predictions, total Sobol' indices for PCE.

Index	Identification	Total Sobol' Index (RP)	Total Sobol' Index (ES)
141	z_4 in SMCEC	0.7095	0.6485
139	z_2 in SMCEC	0.1681	0.1465
142	z_5 in SMCEC	0.0569	0.0341
158	n_cross in WallMechanics	0.0463	0.0645
140	z_3 in SMCEC	0.0189	0.0463

Table 3: $AM + AM_p$ Min

4 Extra figures we probably won't include

4.1 AM_p Time Lag

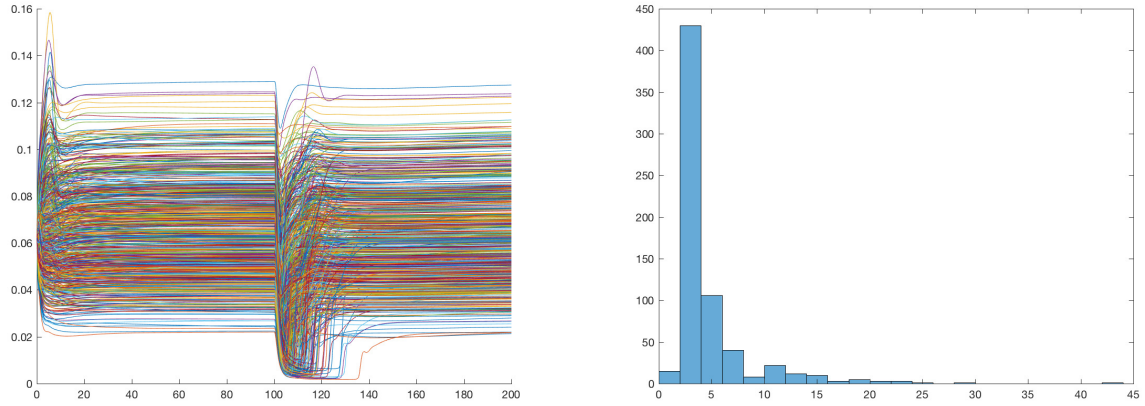


Figure 7: AM_p curves on the left and a histogram of the time lags on the right.

4.2 Radius Time Lag

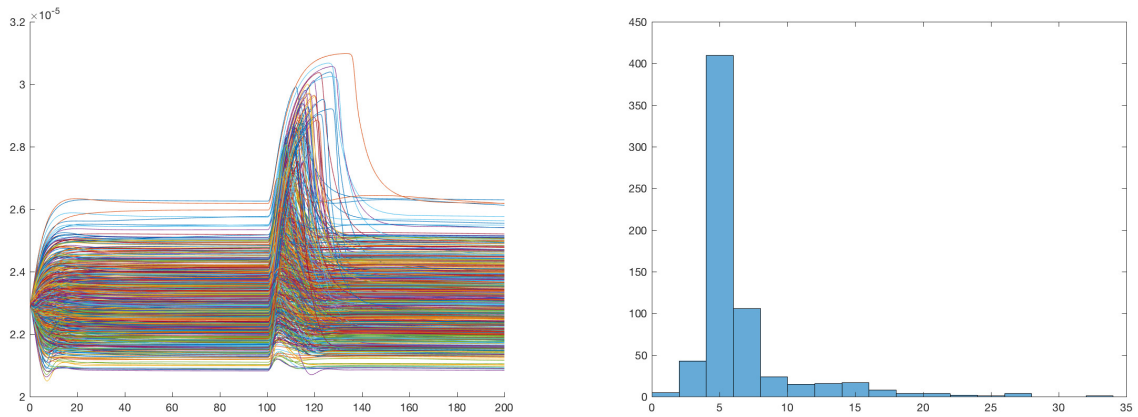


Figure 8: Radius curves on the left and a histogram of the time lags on the right.

5 Discussion

ADD A DISCUSSION OF THE RESULTS IN TERMS OF THEIR PHYSIOLOGICAL INTERPRETATION

6 Conclusion

conclusion goes here

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