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Sensitivity Analysis in a Rescaled 0D Pacemaker-Bimembrane Model

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Abstract

We present a new 0D model of a pacemaker device coupled to a surrogate model of cardiac tissue, based on the work by Pannetier et al. [1]. This model utilizes a simplified internal circuit of an implantable cardiac pacemaker device and includes equivalent models for electrode interfaces and a rescaled minimal myocyte model. Our model integrates the polarization effects commonly observed at electrodes-tissue interface, and allows for the simulation of pacemaker interactions with cardiac tissue, providing insights into energy consumption and device functionality.

1. Introduction

Pacemakers are critical devices that regulate heart rhythm by delivering electrical pulses through electrodes implanted into the cardiac tissue. These pulses are characterized by a duration and an amplitude which, when appropriately selected, provoke a cardiac beat. This is called *capture* by the manufacturers.

As the lifespan of pacemaker batteries is limited, understanding capture and its impact on energy consumption is essential to maximize the durability of the device. We propose an efficient computational model of a pacemaker coupled to a surrogate model of cardiac tissue, based on our previous work [1]. The model is built by simplifying the electronics of a real device into a circuit which allows us to study energy usage during a stimulation cycle. We use the model to compute the so-called *Lapicque* curve in the stimulation (duration, amplitude) space. The curve splits this space into two regions of functional and non-functional stimulations, respectively.

By exploring the parameters of our mathematical model, we aim to provide valuable insights that can help calibrate this model to match experimental data. This calibration is crucial for assessing the credibility of the model and ensuring its accuracy.

2. Model of a pacemaker coupled to a 2-membranes myocyte model

Our new 0D model is built upon an internal simplified circuit of a pacemaker and includes two equivalent models describing the electrode interfaces and a rescaled minimal myocyte model used as a surrogate for cardiac tissue. The circuit for this model is shown in Figure 1.

As in [1], we consider a stimulation signal divided into four parts, and based on a commercial device (Borea™, MicroPort CRM). First, an initial pulse is delivered that is aiming to trigger cellular excitation by discharging a large capacitor. Then, a reverse signal is applied to counteract the polarization which occurs on the surface of the electrodes. This is made possible by two switches, called S_p and S_0 , which swap the sub-circuits inside the device. To account for polarization [2], the electrodes are represented by R//C equivalent circuits (a resistor in parallel with a capacitor). We chose this model because it is the most descriptive model for which the parameters are still identifiable by calibration against experimental data [1].

The tissue is considered through a myocyte model incorporating two rescaled equivalent membranes based on the Beeler-Reuter ionic model [3], and connected by two conductances at cellular level, for the intracellular and extracellular medium.

2.1. Mathematical model

Let I be the current flowing through the pacemaker circuit and V_{+} the two electrode interfaces voltages. The usual Kirchhoff's laws on current yields the following equations [1] at electrodes,

$$
C_{\pm} \frac{dV_{\pm}}{dt} + \frac{V_{\pm}}{R_{\pm}} = I,
$$
 (1)

with R_{\pm} and C_{\pm} the resistances and capacitances of equivalent circuits. We define the time constants $\tau_+ := R_+C_+$.

Additionally, the surrogate tissue part of the circuit verifies the following equations on transmembrane voltages

Figure 1. Electrical circuit describing the model. Left side (black) the device internal circuitry, center (green) the equivalent R//C model of contacts, and right side (red) the two membrane rescaled myocyte equivalent model containing extracellular and intracellular equivalent conductances.

(TMV) $V_{\text{m},1}$, $V_{\text{m},2}$ and ionic states h_1, h_2 ,

$$
C_{\rm m} \frac{\mathrm{d}V_{\rm m,1}}{\mathrm{d}t} + I_{\rm ion,1} = \frac{G_{\rm m}}{S_{\rm m}} \left(V_{\rm m,1} - V_{\rm m,2} - \frac{I}{G_{\rm e}} \right), \tag{2}
$$

$$
\frac{dh_1}{dt} + f_1(h_1, V_{m,1}) = 0,
$$
\n(3)

$$
C_{\rm m} \frac{\mathrm{d}V_{\rm m,2}}{\mathrm{d}t} + I_{\rm ion,2} = \frac{G_{\rm m}}{S_{\rm m}} \left(V_{\rm m,1} - V_{\rm m,2} + \frac{I}{G_{\rm e}} \right), \tag{4}
$$

$$
\frac{dh_2}{dt} + f_2(h_2, V_{m,2}) = 0,
$$
\n(5)

where $G_{\rm m} \coloneqq G_{\rm i} G_{\rm e} (G_{\rm e} + G_{\rm i})^{-1}$ denotes the equivalent conductance of extracellular and intracellular media in mS, C_m is the membrane capacitance in μ F cm⁻², S_m is a scaling coefficient in cm^2 , and with Beeler-Reuter ionic models $(I_{\text{ion},1}, f_1)$ and $(I_{\text{ion},2}, f_2)$.

Finally, the system is closed by writing Kirchhoff's law on voltage on the device part of the circuit,

$$
R_k \frac{dI}{dt} + \frac{I}{C_k} = \frac{V_-}{\tau_-} + \frac{V_+}{\tau_+} + \frac{G_m}{G_e} \left(\frac{dV_{m,1}}{dt} - \frac{dV_{m,2}}{dt} \right),\tag{6}
$$

where R_k and C_k are the resistance and capacitances of the device circuit depending on states of switches S_p and $S_{\rm o}$.

The complete system is solved in our $CircE¹$ software.

2.2. Experimental data

Mathematical model design and parametrisation was guided by experiments using *ex vivo* sheep hearts (N=7, aged 1-2 years old and body weights of 43-52 kg).

Dual coronary-perfused ventricular wedge preparations from hearts opened along the posterior left and right ventricles and posterior septum were stretched on support frames for *ex vivo* assessment into a bath of saline solution, as described in [4] (Fig. 2).

Figure 2. Tissue wedge preparation with the pacemaker electrode in place into the second experiment protocol.

Controlled electrical pulses were applied initially with fixed pulse durations but variable amplitudes that decreased every 8 pulses, starting above the capture threshold. The amplitude decreased until absence of capture is detected by the device. Repeating this process for several durations yields the experimental Lapicque curve.

However, we observed that the pacemaker device is unsuitable due to its limited temporal and voltage resolutions, which are insufficient for an accurate localization of the Lapicque curve. Instead, using a PSA (Pulse System Analyzer) combined with electrical monitoring of the tissue and the lead (PowerLab systems), allowed to test of a larger band of amplitudes and durations with more steps. This setup also allows reaching 100 kHz temporal data acquisition for precise observation of events, such as the rapid pacemaker pulse events.

¹https://carmen.gitlabpages.inria.fr/circe/

3. Global sensitivity analysis

To effectively compare experimental and numerical Lapicque curves, it is essential to establish a good criterion to characterize the capture phenomenon. Whereas experimentally we only have the voltage measured at electrodes, the model gives direct access to the TMVs to define more criteria. Consequently, we introduced several specific quantities derived from both TMVs and voltage V_{meas} , as measured during the experiment at the pacemaker entries (MV). These quantities should be able to provide insights into the membrane's behavior, and local/global response to stimulations.

After studying several quantities on voltages (TMVs and MV), we kept the following observables:

• baseline-adjusted integrals; the baseline is defined as a first value of predefined window starting 5 ms before each stimulation,

• values at end of measurement window; these values typically correspond to the state of the model just before the next stimulation,

• duration between the peak of TMVs and the average distance between it and TMVs baseline (D50), corresponding to the APD50 for capturing stimulations,

• extrema of voltages and of their time-derivatives,

• relative squared differences on pulse, switch and discharge parts of the cycle compared with experimental data obtained on sheep hearts.

We conducted a Sobol sensitivity analysis [5] using the SALib² Python library, to measure the effects of the most uncertain parameters on the observables. The sampling technique and bounds of tested parameters are chosen carefully, as we have no significant a priori estimates for them. The bounds (see table 1) were extended compared to the previous ones [1] since the model was partially modified. The model was evaluated for 786, 432 samples of parameters, but the simulation failed for 2% of them due to physically incompatible values. The evaluation of these failed samples is replaced by the average evaluation of those remaining during the analysis of the Sobol indices. The results are shown on figure 3.

Table 1. Bounds of parameters used in the Sobol sensitivity analysis of the 0D model.

Figure 3. Results of total order indices of Sobol analysis on several outputs features on 786, 432 samples. The three top blocks report the indices for observables derived from $V_{\text{m},1}$, $V_{\text{m},2}$ and V_{meas} respectively. The lower block is dedicated to L^2 relative difference between the model and experimental data.

The analysis shows a strong influence of the tissue parameters (S, G_i, G_e) on the various characteristics of the TMVs, particularly for the first membrane. They also have a significant influence on the integral and on the last value of V_{meas} . Surprisingly, the capacitance value C_{ets} has little influence on these outputs. Additionally, we can also note the parameters that are not identifiable from the experimental data, such as the capacitance C_{ets} and the two resistances of the pacemaker circuit R_p and R_{int} . These parameters should be excluded from any forthcoming calibration procedure.

Finally, this study also provides information on which observables can be used to detect capture. For both TMVs, all quantities are ranked at the same level, but only the integral and the last value of V_{meas} seem to be impacted by parameter changes.

4. Hysteresis of stimulation threshold detection

To further investigate the issue of capture characterization, we used the model to reproduce the hysteresis that is observed when searching the stimulation threshold using either ascending or descending amplitudes. This is commonly observed in [6,7]. It became apparent that although all the criteria can be used to identify the transition from non-capturing to capturing stimulation, the criteria based on the TMVs derivatives or the V_{meas} integral do not have good mathematical properties and do not define a clear threshold. In figure 4, we plot the remaining quantities defined in the previous section, depending on the stimulation amplitude during a threshold search, as performed by the pacemaker.

Different thresholds are observed depending on whether the stimulation amplitude is increasing or decreasing. This confirms that our model is able to reproduce the observed hysteresis. Except for the D50 on the first membrane, all quantities show a large variation around the same threshold amplitude. This should tell that any of these quantities is a good indicator of capture. On the contrary, the values at the end of the stimulation cycle show a very small difference compared to the amplitude of signal, less than 1% for the TMVs (not plotted), and even lower thant 10^{-6} for the MV. This suggests that other observables, such as the maximum or the D50, are safer to characterize capture.

Figure 4. Illustration of hysteresis in the 0D model of 4 quantities. The amplitude varies on x-axis.

5. Conclusion

The hysteresis effect observed numerically underlines the model's ability to reproduce the capture phenomenon on different observables, even for quantities absent from the experimental data (TMVs). However, high Sobol indices for these markers indicate that calibration of the parameters with respect to data must be performed with care to ensure the validity of the model.

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