Examining the Electrical Excitation, Calcium Signaling, and Mechanical Contraction Cycle of a Heart Cell

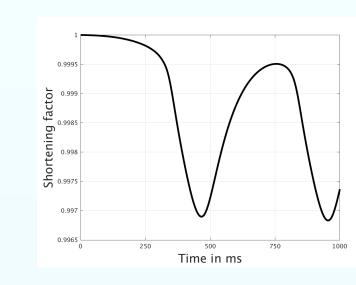
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Goal

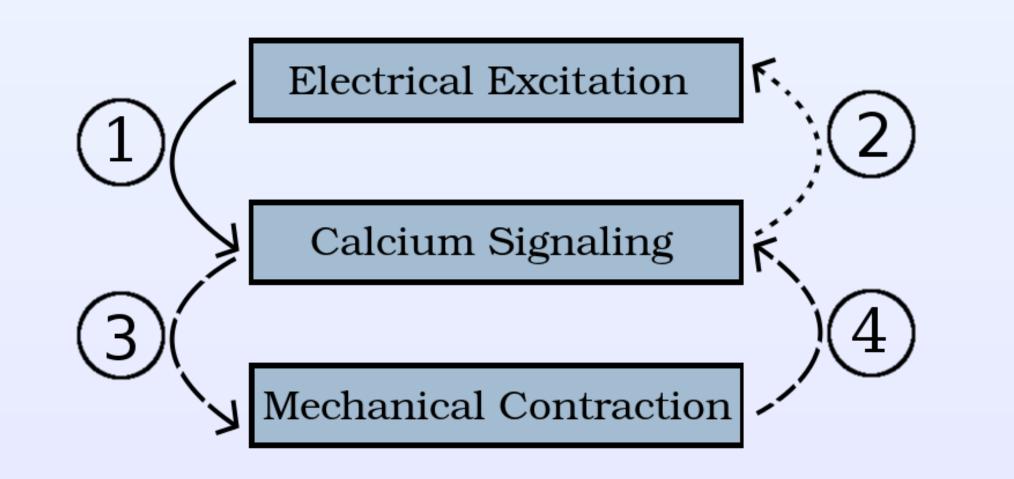
In order to better understand the heart, this work studies the connection between the electrical excitation and mechanical contraction components of the calcium induced calcium release (CICR) cycle in a single heart cell. The heart beats when a collection of heart cells contract through this CICR process, driven by the voltage in the electrical system and ultimately depending on the calcium concentration in the cytosol.

Seven-Variable Simulations

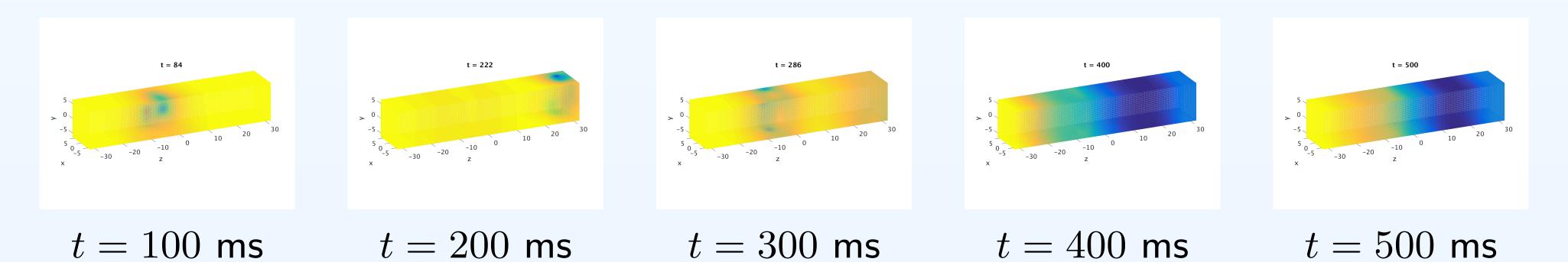
Voltage stimulates release of calcium from the store $s(\mathbf{x}, t)$, limited by its depletion, via CRUs to the cytosol $c(\mathbf{x}, t)$. This calcium binds to actin-myosin cross-bridges $b_3^{(c)}(\mathbf{x}, t)$, which contract the cell. \Rightarrow



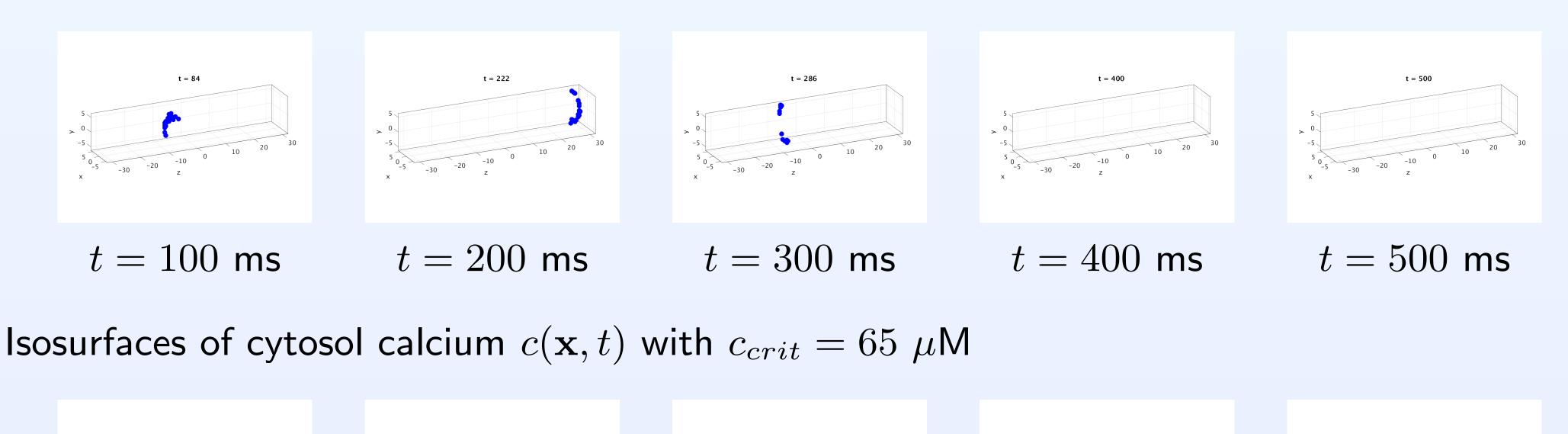
Shortening factor



This work creates a model and executes simulations linking all components of the CICR cycle, including the mechanical component with Links (3) and (4). Isosurfaces of the store SR calcium $s(\mathbf{x},t)$ with $s_{crit} = 5,000 \ \mu M$

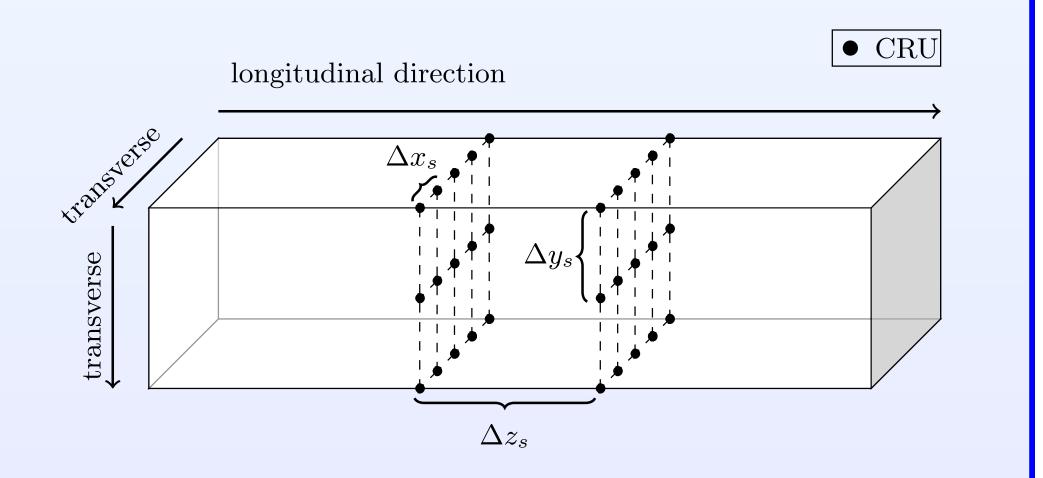


Plots of open calcium release units (CRUs)



Model

This work models a heart cell through the use of seven coupled time-dependent non-linear reaction-diffusion equations. Calcium releases from the store in the sacroplasmic reticulum (SR) into the cytosol through calcium release units (CRUs).



 $t = 100 \text{ ms} \qquad t = 200 \text{ ms} \qquad t = 300 \text{ ms} \qquad t = 400 \text{ ms} \qquad t = 500 \text{ ms}$ Isosurfaces of inactive actin-myosin cross-bridges $b_3^{(c)}(\mathbf{x}, t)$ with $b_{3_{crit}}^{(c)} = 120 \ \mu\text{M}$ $\int_{a}^{b} \frac{1}{a} \frac{1}{$

t = 100 ms t = 200 ms t = 300 ms t = 400 ms t = 500 ms

- Created first fully coupled simulations including mechanical contraction.
- Identified set of parameter values showing realistic effects.

Simulations used FVM with $32 \times 32 \times 128$ mesh for 1 million DOF in each time step.

Acknowledgments

The CRUs form a regular net modeled by a lattice of $15 \times 15 \times 31 = 6,975$ point sources throughout the interior of the 3-D cell.

[1] Kallista Angeloff, Carlos Barajas, et al., *Spora*, vol. 2, 2016.

References

[2] Full technical report: HPCF-2017-15

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• **REU Site**: hpcreu.umbc.edu

• NSF, NSA, DOD, UMBC, HPCF, CIRC