



Langevin, population density and moment-based modeling of local and global aspects of intercellular calcium signaling

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Abstract

Markov chain models of the coupled gating of intracellular calcium (Ca^{2+}) channels are often used to study the stochastic dynamic of local Ca^{2+} release events and whole cell Ca^{2+} homeostasis. However, the runtime of the Markov chain description of Ca^{2+} channel gating is exponential in the number of Ca^{2+} channel states and may thus result in a combinatorial state space explosion when the number of channel states is large. This dissertation presents several novel stochastic modeling approaches that capture important aspects of Ca^{2+} signaling while improving computational efficiency. This dissertation presents several novel stochastic modeling approaches that capture important aspects of calcium Ca^{2+} signaling. First, we present a Ca^{2+} release site modeling approach based on a Langevin description of stochastic Ca^{2+} release. This Langevin model facilitates our investigation of correlations between successive puff/spark amplitudes, durations and inter-spark intervals, and how such puff/spark statistics depend on the number of channels per release site and the kinetics of Ca^{2+} -mediated inactivation of open channels. Second, we show that when the Ca^{2+} channel model is minimal, Langevin equations in a whole cell model involving a large number of release sites may be replaced by a single Fokker-Planck equation. This yields an extremely compact and efficient local/global whole cell model that reproduces and helps interpret recent experiments investigating Ca^{2+} homeostasis in permeabilized ventricular myocytes. Last but not least, we present a population density and moment-based approach to modeling L-type Ca^{2+} channels. Our approaches account for the effect of heterogeneity of local Ca^{2+} signals on whole cell Ca currents. Moreover, they facilitate the study of domain Ca-mediated inactivation of L-type Ca channels.