

POSTER PRESENTATION

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A novel method for approximating equilibrium single-channel Ca^{2+} domains

Victor Matveev

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Localized calcium (Ca^{2+}) signals control some of the most fundamental physiological processes, including synaptic transmission as well as its activity-dependent plasticity. Computational and mathematical modeling played a crucial role in the understanding of spatio-temporal Ca^{2+} dynamics that drives these processes, and showed that Ca^{2+} concentration around a single Ca^{2+} channel reaches a quasi-stationary distribution (known as the Ca^{2+} “nanodomain”) within tens of microseconds after the opening of the channel, and collapses as rapidly after the closing of the channel. Such localization of Ca^{2+} in time and space is achieved by its rapid diffusion as well as its binding to its multiple interaction partners collectively called Ca^{2+} buffers and Ca^{2+} sensors. One of the successes of mathematical modeling was the development of several analytic approximations describing the equilibrium concentration of Ca^{2+} as a function of distance from the open Ca^{2+} channel, such as the Rapid Buffering Approximation (RBA), the Linear Approximation (LA) and the Excess Buffering Approximation (EBA) [1-4]. Each of these approximations has a particular applicability parameter regime created by the interplay between the properties of Ca^{2+} buffers, in particular their mobility and Ca^{2+} binding rates, and the strength of the Ca^{2+} current. Here we present a novel approximation method which does not rely on a specific range of the relevant Ca^{2+} and buffer parameters, and is based on matching the low-distance and large-distance asymptotic behavior of the concentration function. Even at low orders, the resulting approximation is as accurate as the second-order RBA and EBA approximations [4], but its validity extends far beyond the parameter range of applicability of RBA and EBA. The usefulness of the resulting approximation is two-fold: first, together with the previously developed approximations, the novel method could provide a deeper intuition into the

dependence of Ca^{2+} nanodomain properties on the relevant buffering parameters, and second, it constitutes an efficient numerical approximation tool in the modeling of the Ca^{2+} signals underlying presynaptic and postsynaptic phenomena.

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Correspondence: matveev@njit.edu
New Jersey Institute of Technology, NJ 07030, USA



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