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A mesoscopic stochastic mechanism of cytosolic calcium oscillations

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Abstract

Based on a model of intracellular calcium (Ca^{2+}) oscillation with self-modulation of inositol 1,4,5-trisphosphate signal, the mesoscopic stochastic differential equations for the intracellular Ca^{2+} oscillations are theoretically derived by using the chemical Langevin equation method. The effects of the finite biochemical reaction molecule number on both simple and complex cytosolic Ca^{2+} oscillations are numerically studied. In the case of simple intracellular Ca^{2+} oscillation, it is found that, with the increase of molecule number, the coherence resonance or autonomous resonance phenomena can occur for some external stimulation parameter values. In the cases of complex cytosolic Ca^{2+} oscillations, each extremum of concentration of cytosolic Ca^{2+} oscillations corresponds to a peak in the histogram of Ca^{2+} concentration, and the most probability appeared during the bursting plateau level for bursting, but at the largest minimum of Ca^{2+} concentration for chaos. For quasi-periodicity, however, there are only two peaks in the histogram of Ca^{2+} concentration, and the most probability is located at low concentration state. © 2006 Elsevier B.V. All rights reserved.

Keywords: Intracellular calcium oscillations; Chemical Langevin equation; Finite molecule number

1. Introduction

Calcium (Ca^{2^+}) is one of the most important messengers in the cytosol of living cells. Intracellular Ca^{2^+} oscillations play a significant role in signal transduction from receptors at the cell membrane to enzymes and genes controlling the complex biochemical network of cell [1–22]. The information in Ca^{2^+} oscillations can be encoded by their frequency [16–20] as well as by their amplitude [21–23].

Intracellular calcium dynamics has been intensively studied in different theoretical models (for comprehensive reviews see Schuster et al. [24] and Falcke [25]). In the deterministic approach, various models with the macroscopic differential equations have been developed for the widespread phenomenon in intra- and intercellular Ca²⁺ oscillating signalling, much insight has been gained into the processes involved in Ca²⁺ dynamics at the subcellular, cellular and intercellular levels, and the models

In realistic biological system, however, various internal and external fluctuations cannot be negligible. For the internal fluctuations, one source of fluctuations in intracellular Ca²⁺ concentration arises from random releasing of Ca²⁺ channels embedded in the membrane of intracellular stores like the endoplasmic reticulum (ER) [25,58,60]. Thus, some stochastic models have been developed for single Ca²⁺ channels [57], cluster of IP₃R channels [58,60–63], intracellular wave propagation [59,64–68] and intracellular oscillations [69,70]. On the other hand, the cellular biochemical reactions usually occurred in finite system, or the total numbers of biochemical reaction molecules in cell are often finite. At a given temperature, the occurrence of each biochemical reaction

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of calcium oscillations have become more elaborate and diversified [24–55]. In particular, various oscillating behaviors (e.g. bursting, chaos, and quasi-periodicity), different types of bifurcations, and the coupling between oscillating cells have been analyzed. There are many more models and ways to generate complex intracellular Ca²⁺ oscillations [24,25]. Complex Ca²⁺ oscillations may arise through the interplay between two oscillatory mechanisms, but this is not the only possibility, for instance, different agonists may induce different oscillating types in the same cell type [54].

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event in cellular systems is random due to thermal agitation, which is another source of intrinsic fluctuations in intracellular Ca²⁺ oscillations. Furthermore, this intrinsic fluctuation is self-originated in the system. An important fact about the internal fluctuation is that it scales with the system size, and vanishes in the thermodynamic limit. Therefore, for a typical living cell system, the macroscopic differential equations description to intracellular Ca²⁺ oscillations is no longer valid due to the finite size of system, one should consider the effects of finite biochemical reaction molecule number on the intracellular Ca²⁺ oscillations, and a mesoscopic stochastic kinetics should be offered to describe the mechanism of intracellular calcium oscillations. Recently, there has been an increasing interest in the finite size effects on some biological systems [57,71–74].

Now a question to be raised is how to describe the mesoscopic mechanism of intracellular Ca²⁺ oscillations in finite cellular volume, or what are the effects of finite biochemical reaction molecule number on the intracellular Ca²⁺ oscillations. In the present paper, as an example, the deterministic model of intracellular Ca²⁺ oscillation with self-modulation of IP₃ signal proposed by Houart et al. [46] to describe both simple and complex intracellular calcium oscillations has been chosen here. It should be stressed that, to obtain complex Ca²⁺ oscillations, the stochastic models do not require feedback of Ca²⁺ on IP₃ [25,58], for instance, the external noise-induced bursting in a two-variable cytosolic Ca²⁺ oscillation model [56].

The mesoscopic stochastic differential equations for this model have been theoretically derived by using the chemical Langevin equation (CLE) [75] in Section 2. In Section 3, the effects of internal fluctuation due to the finite biochemical reaction molecule number in cell on both simple and complex intracellular Ca²⁺ oscillations have been theoretically studied through numerical computation, respectively. In the case of simple intracellular Ca²⁺ oscillation, it is found that there might be a large signal-to-noise ratio (SNR) with the increase of total molecule number for some external stimulation parameter values, which corresponds to the coherence resonance or autonomous resonance phenomena. In the case of complex intracellular Ca²⁺ oscillations, the histograms of cytosolic Ca²⁺ concentration are different for each typical complex Ca2+ oscillatory behavior in finite biochemical reaction molecule number. Finally, we end with conclusions in Section 4.

2. Mesoscopic stochastic differential equations for intracellular Ca^{2^+} oscillations

The key species in the model of intracellular Ca^{2^+} oscillation with self-modulation of IP_3 signal [46] are the cytosolic Ca^{2^+} (its concentration is represented by Z), the Ca^{2^+} sequester in an internal store (its concentration is represented by Y), and the IP_3 (its concentration is represented by A) which is another important intracellular messenger. The time evolution of these species can be described by following macroscopic differential equations

$$\frac{dZ}{dt} = V_0 + V_1 \beta - V_2 + V_3 + k_f Y - kZ,\tag{1}$$

$$\frac{\mathrm{d}Y}{\mathrm{d}t} = V_2 - V_3 - k_\mathrm{f}Y,\tag{2}$$

$$\frac{\mathrm{d}A}{\mathrm{d}t} = \beta V_4 - V_5 - \varepsilon A,\tag{3}$$

where the external stimulation parameter β reflects the degree of stimulation of the cell by an agonist and thus only varies between 0 and 1, and has two Hopf bifurcation points. Detail description of other parameters in this model can be found in Table 1.

The three species S_i (i=1, 2, 3) considered here are cytosolic Ca^{2+} , Ca^{2+} sequester in internal store, and intracellular IP_3 , respectively. It is introduced that the number of calcium ions in cytosol as z, the number of calcium ions in internal store as y, and the number of IP_3 molecules in cytosol as a. Then, the relationship between the concentration and the total number of molecules is

$$Z = \frac{z}{Q}, \quad Y = \frac{y}{Q}, \quad A = \frac{a}{Q},$$
 (4)

where Ω is defined as the total molecule (including the cytosolic Ca^{2+} , intravesicular Ca^{2+} , and intracellular IP_3) number in cell. Nine elementary biochemical reaction processes are considered in the model of intracellular Ca^{2+} oscillation with self-modulation of IP_3 signal. Note that there are twelve reaction channels R_j ($j=1,\ldots,12$) for the three species S_i (i=1,2,3) as shown in Fig. 1, where the reaction channels R_3 and R_7 , R_4 and R_8 , and R_5 and R_9 represent the same biochemical reaction process, respectively. The transition rates of the twelve reaction channels are marked by $r_j=1,\ldots,12$. The corresponding transition rates for the twelve channels are described in Table 2, where the transition rates are proportional to the total number of intracellular molecules Ω . The transition rate r_j for R_j and the state-change vector $\overrightarrow{v_j}$, whose ith component $v_{i,i}$ is the change in

Table 1
Parameters in the model of intracellular Ca²⁺ oscillations [46]

Parameter	Description			
V_0	Constant input from extracellular medium			
V_1	Maximum rate of stimulus-induced influx from the extracellular medium			
V_2	Pump flux of cytosolic Ca ²⁺ into the internal stores with a maximum value V_{M2} , $V_2 = V_{M2} \frac{Z^2}{K_1^2 + Z^2}$			
V_3	Release of Ca ²⁺ from internal stores with a maximum value V_{M3} , $V_3 = V_{M3} \frac{Z^m}{K_+^m + Z^m} \frac{Y^2}{K^2 + Y^2} \frac{A^4}{K^4 + A^4}$			
V_4	Maximum rate of stimulus-induced synthesis of IP ₃			
V_5	Rate of phosphorylation of IP ₃ by the 3-kinase with a maximum value $V_{\rm M5}$ and a half-saturation constant K_5 , $V_5 = V_{\rm M5} \frac{A^P}{K^P + A^P} \frac{Z^n}{K^n + Z^n}$			
K_2	Threshold constant for pumping			
K_Y	Threshold constant for release by Ca ²⁺			
K_Z	Threshold constant for activation by Ca ²⁺			
K_A	Threshold constant for activation by IP ₃			
$k_{ m f}$	Rate constant measuring the passive, linear leak of Y into Z			
k	Rate constant of linear transport of cytosolic Ca ²⁺ into the extracellular medium			
K _d	Threshold constant for 3-kinase stimulated by Ca ²⁺			

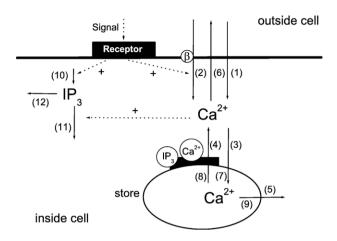


Fig. 1. The schematic description of the model [46]. There are twelve reaction channels marked with (1)–(12), respectively.

the number of S_i (i=1, 2, 3) molecules produced by one R_j (j=1, ..., 12) reaction channel, together completely specify the reaction channel R_j .

Table 2 Reaction channels and corresponding transition rates

Reaction channels	Description	Transition rates
$(1) z \rightarrow z + 1$	Constant input of cytosolic Ca ²⁺ from extracellular medium	$r_1 = \Omega V_0$
$(2) z \rightarrow z + 1$		$r_2 = \Omega V_1 \beta$
$(3) z \rightarrow z - 1$	Pumping of cytosolic Ca ²⁺ into internal store	$r_3 = \Omega \frac{V_{\text{M2}} Z^2}{K_Z^2 + Z^2}$
$(4) z \rightarrow z + 1$	Input of cytosolic Ca ²⁺ from internal store	$r_{4} = \Omega \frac{V_{\text{M3}} Z^{m}}{K_{Z}^{m} + Z^{m}} \frac{Y^{2}}{K_{Y}^{2} + Y^{2}} \frac{A^{4}}{K_{A}^{4} + A^{4}}$
$(5) z \rightarrow z+1$	Input of cytosolic Ca ²⁺ from internal store	$r_5 = \Omega k_{\mathrm{f}} Y$
$(6) z \rightarrow z - 1$	Leakage of cytosolic Ca ²⁺ into extracellular medium	$r_6 = \Omega kZ$
$(7) y \rightarrow y + 1$	Pumping of cytosolic Ca ²⁺ into internal store	$r_7 = \Omega \frac{V_{\rm M2} Z^2}{K_2^2 + Z^2}$
$(8) y \rightarrow y - 1$	Release of Ca ²⁺ from internal store	$r_8 = \Omega \frac{V_{\rm M3} Z^m}{K_Z^m + Z^m} \frac{Y^2}{K_Y^2 + Y^2} \frac{A^4}{K_A^4 + A^4}$
$(9) y \rightarrow y - 1$	Leakage of internal pool Ca ²⁺ from internal store	$r_9 = Qk_{\rm f}Y$
$(10) \ a \rightarrow a+1$	Stimulus-induced synthesis of IP ₃	$r_{10} = \Omega \beta V_4$
$(11) \ a \rightarrow a - 1$	Phosphorylated IP ₃ by 3-kinase	$r_{11} = \Omega \frac{V_{\text{M5}} A^P}{K_5^P + A^P} \frac{Z^n}{K_d^n + Z^n}$
$(12) \ a \rightarrow a - 1$	Metabolized IP ₃ by 5-phosphatase	$r_{12} = Q \epsilon A$

Following along the Gillespie's method [75], suppose the system's state at the current time t is known to be (z(t), y(t), a(t)). Let a random variable $K_j(z(t), y(t), a(t), \tau)$, for any $\tau > 0$, be the number of R_j reactions that occur in the subsequent time interval $[t, t+\tau]$. Since each of those reactions will increase the S_i population by $v_{j,i}$ (see Table 3), thus, the number of S_i molecules in the system at time $t+\tau$ will be

$$z(t+\tau) = z(t) + K_1(z(t), y(t), a(t), \tau) + K_2(z(t), y(t), a(t), \tau) - K_3(z(t), y(t), a(t), \tau) + K_4(z(t), y(t), a(t), \tau) + K_5(z(t), y(t), a(t), \tau) - K_6(z(t), y(t), a(t), \tau),$$
 (5)

$$y(t+\tau) = y(t) + K_7(z(t), y(t), a(t), \tau) -K_8(z(t), y(t), a(t), \tau) - K_9(z(t), y(t), a(t), \tau),$$
(6)

$$a(t+\tau) = a(t) + K_{10}(z(t), y(t), a(t), \tau) -K_{11}(z(t), y(t), a(t), \tau) - K_{12}(z(t), y(t), a(t), \tau).$$
(7)

An excellent approximation to $K_j(z(t), y(t), a(t), \tau)$ in Eqs. (5)–(7) can be obtained if the following two conditions are imposed [75]. (i) Require τ to be small enough that the change in the state during $[t, t+\tau]$ will be so slight that none of the propensity functions changes its value appreciably. Each $K_j(z(t), y(t), a(t), \tau)$ will be a statistically independent Poisson random variable, $\mathcal{P}_j(r_j(z(t), y(t), a(t)), \tau)$. Eqs. (5)–(7) are approximated by

$$\begin{split} z(t+\tau) &= z(t) + \mathcal{P}_1(r_1(z(t),y(t),a(t)),\tau) \\ &+ \mathcal{P}_2(r_2(z(t),y(t),a(t)),\tau) - \mathcal{P}_3(r_3(z(t),y(t),a(t)),\tau) \\ &+ \mathcal{P}_4(r_4(z(t),y(t),a(t)),\tau) + \mathcal{P}_5(r_5(z(t),y(t),a(t)),\tau) \\ &- \mathcal{P}_6(r_6(z(t),y(t),a(t)),\tau), \end{split}$$

(8)

$$y(t+\tau) = y(t) + \mathcal{P}_7(r_7(z(t), y(t), a(t)), \tau) -\mathcal{P}_8(r_8(z(t), y(t), a(t)), \tau) - \mathcal{P}_9(r_9(z(t), y(t), a(t)), \tau),$$
(9)

$$a(t+\tau) = a(t) + \mathcal{P}_{10}(r_{10}(z(t), y(t), a(t)), \tau) - \mathcal{P}_{11}(r_{11}(z(t), y(t), a(t)), \tau) - \mathcal{P}_{12}(r_{12}(z(t), y(t), a(t)), \tau).$$
(10)

(ii) Require τ to be large enough that the expected number of occurrences of each reaction channel R_j in $[t, t+\tau]$ be much larger than 1, which allows us to approximate each Poisson random variable $\mathcal{P}_j(r_j(z(t), y(t), a(t)), \tau)$ by a normal random variable with

Table 3 Population change $v_{i,i}$ of specie S_i (i = 1, 2, 3) in R_i (j = 1, ..., 12) reaction channel

Specie	Population change
S ₁ (cytosolic Ca ²⁺)	$v_{1,1} = v_{2,1} = 1, v_{3,1} = -1, v_{4,1} = v_{5,1} = 1, v_{6,1} = -1,$
S_2 (Ca ²⁺ in internal store)	$v_{7,1} = v_{8,1} = v_{9,1} = v_{10,1} = v_{11,1} = v_{12,1} = 0$ $v_{1,2} = v_{2,2} = v_{3,2} = v_{4,2} = v_{5,2} = v_{6,2} = 0, v_{7,2} = 1,$
S ₃ (intracellular IP ₃)	$v_{8,2} = v_{9,2} = -1$, $v_{10,2} = v_{11,2} = v_{12,2} = 0$ $v_{1,3} = v_{2,3} = v_{3,3} = v_{4,3} = v_{5,3} = v_{6,3} = v_{7,3} = v_{8,3} = v_{9,3} = 0$, $v_{10,3} = 1$, $v_{11,3} = v_{12,3} = -1$

the same mean and variance. That brings Eqs. (8)–(11) into the form

$$\begin{split} z(t+\tau) &= z(t) \\ &+ \mathcal{N}_1(r_1(z(t),y(t),a(t))\tau, r_1(z(t),y(t),a(t))\tau) \\ &+ \mathcal{N}_2(r_2(z(t),y(t),a(t))\tau, r_2(z(t),y(t),a(t))\tau) \\ &- \mathcal{N}_3(r_3(z(t),y(t),a(t))\tau, r_3(z(t),y(t),a(t))\tau) \\ &+ \mathcal{N}_4(r_4(z(t),y(t),a(t))\tau, r_4(z(t),y(t),a(t))\tau) \\ &+ \mathcal{N}_5(r_5(z(t),y(t),a(t))\tau, r_5(z(t),y(t),a(t))\tau) \\ &- \mathcal{N}_6(r_6(z(t),y(t),a(t))\tau, r_6(z(t),y(t),a(t))\tau), \end{split}$$

$$y(t+\tau) = y(t) + \mathcal{N}_{7}(r_{7}(z(t), y(t), a(t))\tau, r_{7}(z(t), y(t), a(t))\tau) - \mathcal{N}_{8}(r_{8}(z(t), y(t), a(t))\tau, r_{8}(z(t), y(t), a(t))\tau) - \mathcal{N}_{9}(r_{9}(z(t), y(t), a(t))\tau, r_{9}(z(t), y(t), a(t))\tau),$$
(12)

$$a(t+\tau) = a(t) + \mathcal{N}_{10}(r_{10}(z(t), y(t), a(t))\tau, r_{10}(z(t), y(t), a(t))\tau) - \mathcal{N}_{11}(r_{11}(z(t), y(t), a(t))\tau, r_{11}(z(t), y(t), a(t))\tau) - \mathcal{N}_{12}(r_{12}(z(t), y(t), a(t))\tau, r_{12}(z(t), y(t), a(t))\tau),$$
(13)

where $N_j(m_j, \sigma_j^2)$ denotes the normal random variable with mean m_j and variance σ_j^2 , and is statically independent with each other because of the statistically independent Poisson random variable. The molecular populations from discretely changing integer variables in Eqs. (8)–(10) are converted to continuously changing real variables in Eqs. (11)–(13) in effect. The linear combination theorem for normal random variables.

$$\mathcal{N}_j(m_j, \sigma_j^2) = m_j + \sigma_j \mathcal{N}_j(0, 1), \tag{14}$$

can now be invoked to bring Eqs. (11)-(13) into the form

$$z(t+\tau) = z(t) + r_1(z(t), y(t), a(t))\tau + r_2(z(t), y(t), a(t))\tau - r_3(z(t), y(t), a(t))\tau + r_4(z(t), y(t), a(t))\tau + r_5(z(t), y(t), a(t))\tau + [r_1(z(t), y(t), a(t))\tau]^{1/2}\mathcal{N}_1(0, 1) + [r_2(z(t), y(t), a(t))\tau]^{1/2}\mathcal{N}_2(0, 1) - [r_3(z(t), y(t), a(t))\tau]^{1/2}\mathcal{N}_3(0, 1) + [r_4(z(t), y(t), a(t))\tau]^{1/2}\mathcal{N}_4(0, 1) + [r_5(z(t), y(t), a(t))\tau]^{1/2}\mathcal{N}_5(0, 1) - [r_6(z(t), y(t), a(t))\tau]^{1/2}\mathcal{N}_6(0, 1)$$

$$(15)$$

$$y(t+\tau) = y(t) + r_7(z(t), y(t), a(t))\tau -r_8(z(t), y(t), a(t))\tau -r_9(z(t), yt, a(t))\tau +[r_7(z(t), y(t), a(t))\tau]^{1/2}\mathcal{N}_7(0, 1) -[r_8(z(t), y(t), a(t))\tau]^{1/2}\mathcal{N}_8(0, 1) -[r_9(z(t), y(t), a(t))\tau]^{1/2}\mathcal{N}_9(0, 1),$$
(16)

$$a(t+\tau) = a(t) + r_{10}(z(t), y(t), a(t))\tau -r_{11}(z(t), y(t), a(t))\tau -r_{12}(z(t), y(t), a(t))\tau + [r_{10}(z(t), y(t), a(t))\tau]^{1/2} \mathcal{N}_{10}(0, 1) - [r_{11}(z(t), y(t), a(t))\tau]^{1/2} \mathcal{N}_{11}(0, 1) + [r_{12}(z(t), y(t), a(t))\tau]^{1/2} \mathcal{N}_{12}(0, 1),$$

$$(17)$$

where $N_{j=1,...,12}(0, 1)$ are statically independent with each other. Let us regard any time interval τ that satisfies both conditions (i) and (ii) as a macroscopic infinitesimal, and denote it simply by dt, and write the unit normal random variable $N_j(0, 1)$ as $N_j(t)$. Eqs. (15)–(17) become

$$\begin{split} z(t+\mathrm{dt}) &= z(t) + r_1(z(t),y(t),a(t))\mathrm{dt} \\ &+ r_2(z(t),y(t),a(t))\mathrm{dt} - r_3(z(t),y(t),a(t))\mathrm{dt} \\ &+ r_4(z(t),y(t),a(t))\mathrm{dt} + r_5(z(t),y(t),a(t))\mathrm{dt} \\ &- r_6(z(t),y(t),a(t))\mathrm{dt} \\ &+ r_1^{1/2}(z(t),y(t),a(t))\mathcal{N}_1(t)(\mathrm{dt})^{1/2} \\ &+ r_2^{1/2}(z(t),y(t),a(t))\mathcal{N}_2(t)(\mathrm{dt})^{1/2} \\ &- r_3^{1/2}(z(t),y(t),a(t))\mathcal{N}_3(t)(\mathrm{dt})^{1/2} \\ &+ r_4^{1/2}(z(t),y(t),a(t))\mathcal{N}_4(t)(\mathrm{dt})^{1/2} \\ &+ r_5^{1/2}(z(t),y(t),a(t))\mathcal{N}_5(t)(\mathrm{dt})^{1/2} \\ &- r_5^{1/2}(z(t),y(t),a(t))\mathcal{N}_5(t)(\mathrm{dt})^{1/2} \\ &- r_6^{1/2}(z(t),y(t),a(t))\mathcal{N}_6(t)(\mathrm{dt})^{1/2}, \end{split}$$

$$y(t + dt) = y(t) + r_7(z(t), y(t), a(t)), dt$$

$$-r_8(z(t), y(t), a(t))dt - r_9(z(t), y(t), a(t))dt$$

$$+r_7^{1/2}(z(t), y(t), a(t)\mathcal{N}_7(t)(dt)^{1/2}$$

$$-r_8^{1/2}(z(t), y(t), a(t))\mathcal{N}_8(t)(dt)^{1/2}$$

$$-r_9^{1/2}(z(t), y(t), a(t))\mathcal{N}_9(t)(dt)^{1/2},$$
(19)

$$a(t + dt) = a(t) + r_{10}(z(t), y(t), a(t))dt - r_{11}(z(t), y(t), a(t))dt - r_{12}(z(t), y(t), a(t))dt + r_{10}^{1/2}(z(t), y(t), a(t))\mathcal{N}_{10}(t)(dt)^{1/2} - r_{11}^{1/2}(z(t), y(t), a(t))\mathcal{N}_{11}(t)(dt)^{1/2} - r_{12}^{1/2}(z(t), y(t), a(t))\mathcal{N}_{12}(t)(dt)^{1/2}.$$
(20)

Eqs. (18)–(20) have the canonical form of standard Langevin equations for multivariate continuous Markov processes. Thus, Eqs. (18)–(20) imply the equivalent white-noise form Langevin equation

$$\frac{\mathrm{d}z(t)}{\mathrm{d}t} = r_1(z(t), y(t), a(t)) + r_2(z(t), y(t), a(t)) \\
-r_3(z(t), y(t), a(t)) + r_4(z(t), y(t), a(t)) \\
+r_5(z(t), y(t), a(t)) - r_6(z(t), y(t), a(t)) \\
+r_1^{1/2}(z(t), y(t), a(t)) \xi_1(t) \\
+r_2^{1/2}(z(t), y(t), a(t)) \xi_2(t) \\
-r_3^{1/2}(z(t), y(t), a(t)) \xi_3(t) \\
+r_4^{1/2}(z(t), y(t), a(t)) \xi_4(t) \\
+r_5^{1/2}(z(t), y(t), a(t)) \xi_5(t) \\
-r_6^{1/2}(z(t), y(t), a(t)) \xi_6(t), \tag{21}$$

$$\frac{\mathrm{d}y(t)}{\mathrm{d}t} = r_7(z(t), y(t), a(t)) - r_8(z(t), y(t), a(t)) \\
-r_9(z(t), y(t), a(t)) \\
+r_7^{1/2}(z(t), y(t), a(t))\xi_7(t) \\
-r_8^{1/2}(z(t), y(t), a(t))\xi_8(t) \\
-r_9^{1/2}(z(t), y(t), a(t))\xi_9(t),$$
(22)

$$\begin{split} \frac{\mathrm{d}a(t)}{\mathrm{d}t} &= r_{10}(z(t), y(t), a(t)) - r_{11}(z(t), y(t), a(t)) \\ &- r_{12}(z(t), y(t), a(t)) \\ &+ r_{10}^{1/2}(z(t), y(t), a(t)) \xi_{10}(t) \\ &- r_{11}^{1/2}(z(t), y(t), a(t)) \xi_{11}(t) \\ &- r_{12}^{1/2}(z(t), y(t), a(t)) \xi_{12}(t), \end{split} \tag{23}$$

where $\xi_{i=1,...,12}(t)$ are temporally uncorrelated, statistically independent Gaussian white noises with $\langle \xi_i(t) \rangle = 0$ and $\langle \xi_i(t) \rangle = \delta_{ii}\delta(t-s)$.

By using the relationship between the concentration and the molecular number of each species, Eq. (4), the CLE corresponding to the macroscopic differential Eqs. (1)–(3) can be obtained from Eqs. (21)–(23) and read as

$$\frac{dZ}{dt} = (V_0 + V_1 \beta - V_2 + V_3 + k_f Y - kZ) + \frac{1}{\sqrt{\Omega}} \left[\sqrt{V_0} \xi_1(t) + \sqrt{V_1 \beta} \xi_2(t) - \sqrt{V_2} \xi_3(t) + \sqrt{V_3} \xi_4(t) + \sqrt{k_f Y} \xi_5(t) - \sqrt{kZ} \xi_6(t) \right],$$
(24)

$$\frac{dY}{dt} = (V_2 - V_3 - k_f Y) + \frac{1}{\sqrt{\Omega}} \left[\sqrt{V_2} \xi_7(t) - \sqrt{V_3} \xi_8(t) - \sqrt{k_f Y} \xi_9(t) \right], \tag{25}$$

$$\frac{\mathrm{d}A}{\mathrm{d}t} = (\beta V_4 - V_5 - \varepsilon A) + \frac{1}{\sqrt{\Omega}} \left[\sqrt{\beta V_4} \xi_{10}(t) - \sqrt{V_5} \xi_{11}(t) - \sqrt{\varepsilon A} \xi_{12}(t) \right].$$
(26)

It can be noted that the internal fluctuation item is proportional to $1/\sqrt{\Omega}$ in the mesoscopic stochastic differential Eqs. (24) (25) (26) when the other parameters are fixed, and vanishes in the thermodynamic limit (i.e. $\Omega \rightarrow \infty$).

The model Eqs. (1)–(3) for cytosolic Ca²⁺ behaviors shows not only for simple periodic Ca²⁺ oscillations, but also for some complex oscillatory phenomena. The dynamic behavior of the model in parameter space had been investigated in Ref. [46], and it was shown that the complex Ca²⁺ oscillatory behaviors include bursting, chaos and quasi-periodicity. Four sets of parameter values corresponding to the simple oscillations and the complex oscillatory behaviors are listed in Table 4. To study the

Table 4
Parameter values corresponding to the simple oscillations and the various types of complex oscillatory behavior in the Ca²⁺ oscillations model [46]

Parameters	Simple oscillation	Bursting	Chaos	Quasi-periodicity
n	4.0	2.0	4.0	4.0
m	2.0	4.0	2.0	2.0
p	2.0	1.0	1.0	2.0
β	0.50	0.46	0.65	0.51
$K_2 (\mu M)$	0.1	0.1	0.1	0.1
K_5 (μ M)	1.0	1.0	0.3194	0.3
K_A (μ M)	0.2	0.1	0.1	0.2
$K_{\rm d}$ (μ M)	0.4	0.6	1.0	0.5
$K_Y(\mu M)$	0.2	0.2	0.3	0.2
$K_Z(\mu M)$	0.5	0.3	0.6	0.5
$k (\min^{-1})$	10.0	10.0	10.0	10.0
$k_{\rm f}({\rm min}^{-1})$	1.0	1.0	1.0	1.0
$\epsilon (\text{min}^{-1})$	0.1	1.0	13.0	0.1
$V_0 (\mu \mathrm{M \ min}^{-1})$	2.0	2.0	2.0	2.0
$V_1 (\mu \text{Mmin}^{-1})$	2.0	2.0	2.0	2.0
$V_{\rm M2}~(\mu {\rm M~min}^{-1})$	6.0	6.0	6.0	6.0
$V_{\rm M3}~(\mu {\rm M~min}^{-1})$	20.0	20.0	30.0	20.0
$V_4 (\mu \mathrm{M \ min}^{-1})$	2.0	2.5	3.0	5.0
$V_{\rm M5}~(\mu {\rm M~min}^{-1})$	5.0	30.0	50.0	30.0

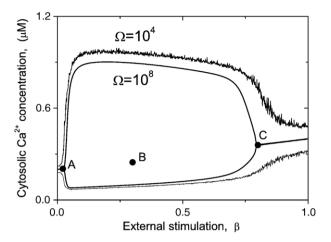


Fig. 2. Bifurcation diagram for simple intracellular calcium oscillation, where β =0.028 for point A, β =0.3 for point B, and β =0.8 for point C.

effects of the finite molecule number on both simple and complex Ca²⁺ oscillations, numerical simulations are needed, and the mesoscopic stochastic Eqs. (24)–(26) are simulated by using a simple forward Euler algorithm with a time step of 0.001 min. In each calculation the time evolution of the system lasted 1000 min after transient behavior was discarded.

3. Effects of finite total molecule number on intracellular calcium oscillations

It is well known that Ca²⁺ relays information within cells to regulate their activity, and the information can be encoded by both the frequency and the amplitude of Ca²⁺ oscillations. In this section, the effects of total molecule number on the various types of intracellular calcium oscillations have been investigated through the power spectrum and the distribution (the histogram) of Ca²⁺ concentration.

3.1. Simple intracellular calcium oscillation

The external control parameter β has two supercritical Hopf bifurcation points for the macroscopic kinetics. The intracellular Ca²⁺ concentration is oscillated for $\beta_1 < \beta < \beta_2$, and is stable steady state for $\beta < \beta_1$ or $\beta > \beta_2$. It is known that the smaller the molecule number is, the larger the internal fluctuation will be. For the mesoscopic kinetics (i.e. Eqs. (24)–(26)), the bifurcation diagrams for different molecule numbers are plotted in Fig. 2. Three values of the external control parameter β (=0.028, 0.3, 0.8, marked by A, B and C in Fig. 2) were chosen to discuss the effects of finite molecule number on the intracellular Ca²⁺ oscillation.

When β =0.3 (at which the intracellular Ca²⁺ concentration shows simple oscillations in the case of macroscopic kinetics), the cytosolic Ca²⁺ oscillation and the histogram of Ca²⁺ concentration are plotted in Fig. 3 for different molecule numbers Ω . It is shown that there is only one peak in the histogram of Ca²⁺ concentration, which is located at low cytosolic Ca²⁺ concentration for small Ω . With the increase of molecule number, a tiny peak appeared at high Ca²⁺ concentration states.

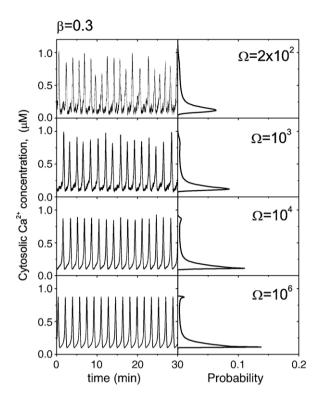


Fig. 3. Temporal evolution of cytosolic Ca^{2+} concentration (left) and corresponding histogram of Ca^{2+} concentration (right) for different molecule numbers Ω . The other parameter values corresponding to the simple cytosolic oscillation are listed in Table 4.

To characterize the effects of finite molecule number on the stochastic cytosolic Ca²⁺ oscillation, the coefficient of variation (CV) [76]

$$CV = \sqrt{\frac{\langle T^2 \rangle - \langle T \rangle^2}{\langle T \rangle}}$$
 (27)

has been calculated, where $\langle T \rangle := \lim_{N \to \infty} \sum_{i=1}^N (t_{i+1} - t_i)/N$ and $\langle T^2 \rangle := \lim_{N \to \infty} \sum_{i=1}^N (t_{i+1} - t_i)^2/N$ are the mean and mean-squared interspike intervals, respectively. A spike in the Ca²⁺ concentration $Z(t_i)$ occurred at time t_i . In our computations, a spike will occur when the Ca²⁺ concentration exceeds a threshold $(Z_c = 0.5 \ \mu\text{M})$. The CV presents a measure of spike coherence (i.e. the order degree of stochastic Ca²⁺ oscillations). With the increase of Ω , Fig. 4 shows that the mean interspike interval is increased, and then tends to a limit value 1.78218 min (the period of cytosolic Ca²⁺ oscillations in the case of macroscopic kinetics), while the CV is decreased and tends to zero if $\Omega \to \infty$.

When β =0.028 or β =0.8 (at which the intracellular Ca²⁺ concentration is stable steady state in the case of macroscopic kinetics), the power spectrum of cytosolic Ca²⁺ concentration for different Ω is plotted in Fig. 5. A Welch window function is used during the estimation of power spectrum. The time series of stochastic calcium concentration Z contains 2¹⁶ data points, and the smoothed curves of power spectrum are obtained by nearest averaging over 50 points. It can be found that there exists a peak of power spectrum. With the increase of molecule

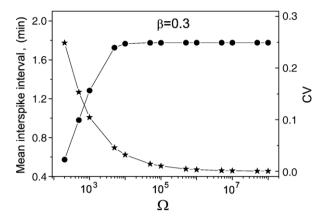


Fig. 4. Mean interspike interval $\langle T \rangle$ (dot line) and its coefficient of variation (CV) (star line) as a function of the molecule number. The other parameter values are the same as in Fig. 3.

number (in other words, with the decrease of internal fluctuation), (i) the height of power spectrum's peak is increased firstly, reaches a maximum, and then is decreased; (ii) the width of power spectrum's peak becomes narrower; (iii) our computation results also showed that the optimal frequency corresponding to the peak shifts from low to high when the molecule number is increased from 10^2 to 10^5 , however, the optimal frequency is nearly not varied when $\Omega > 10^5$.

The above results show that there exists the most pronounced peak of power spectrum for certain intermediate molecule number. To quantify the relative performance of the stochastic calcium concentration oscillations, one can define an effective SNR [60-62,71-74,76]:

$$SNR = R \frac{\omega_{\rm p}}{\Lambda \omega},\tag{28}$$

where ω_p is the frequency at the peak, $\Delta\omega$ is the width between ω_p and ω_1 satisfying $\omega_1 > \omega_p$ and $P(\omega_1) = P(\omega_p)/e$, $R = P(\omega_p)/P(\omega_2)$, and $P(\omega_2)$ is the smallest power spectrum value $(0 < \omega_2 - \omega_p)$. The SNR with the increase of molecule number is

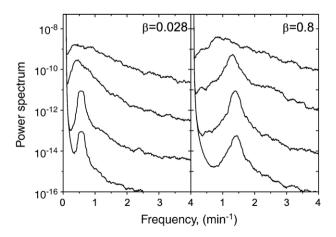


Fig. 5. Power spectrum of cytosolic Ca^{2+} concentration for β =0.028 (left) and β =0.8 (right). From top to bottom: Ω =10², 10⁴, 10⁶, and 10⁸. The other parameter values are the same as in Fig. 3.

respectively plotted for β =0.028 and β =0.8 in Fig. 6. It is clear that there is a maximum of SNR at certain optimal molecule numbers ($\Omega \sim 10^6$ under the given values of β here). This is a signature of resonance which is the typical coherence resonance or autonomous resonance phenomenon since there is no external periodic signal. It should be pointed out that (i) the maximum of SNR and the coherence resonance had been found by Meinhold and Schimansky-Geier [60] corresponding to the optimal IP₃R-I channel numbers of a single cluster there, but corresponding to the optimal biochemical reaction molecule numbers here. (ii) It was also found that [77,78] the optimal system size (i.e. the volume) of SNR depends on the external control parameter, the optimal volume matches well with the real cell volume when the control parameter is tuned near the left Hopf bifurcation point, and the size of real living cells in vivo is around $10^3 \, \mu m^3$ [68].

3.2. Complex intracellular calcium oscillation

Complex Ca^{2+} oscillatory behaviors of the model include bursting, chaos and quasi-periodicity. In the case of mesoscopic kinetics of intracellular Ca^{2+} oscillations, the time series of cytosolic Ca^{2+} concentration for the three types of complex Ca^{2+} oscillations is plotted in Fig. 7, respectively. When the molecule number Ω is small (e.g. Ω =200), various Ca^{2+} oscillatory types are hard to distinguish, and the irregularity of the oscillations shows up both in the amplitude and in the time interval between successive Ca^{2+} spikes due to the large internal fluctuation. However, with the increase of molecule number (or decrease of the intrinsic fluctuation), the differences between various cytosolic Ca^{2+} oscillation types of complex Ca^{2+} oscillations are much more remarkable.

To characterize the effects of finite molecule number on the complex cytosolic Ca^{2+} oscillations, the histograms of Ca^{2+} concentration for different molecule numbers Ω are shown in Fig. 8. For the bursting type, when the molecule number is very small (e.g. Ω =200), there is only one peak located at low Ca^{2+} concentration state in the histograms of Ca^{2+} concentration. With the increase of molecule number, more and more peaks

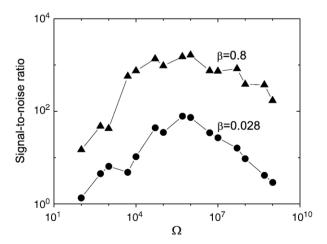
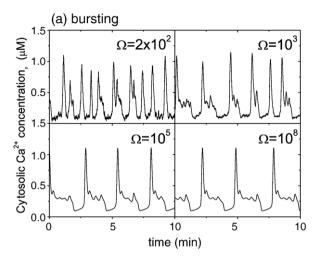
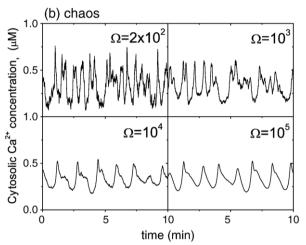


Fig. 6. Signal-to-noise ratio vs. the finite molecule number for β =0.028 and β =0.8. The other parameter values are the same as in Fig. 3.





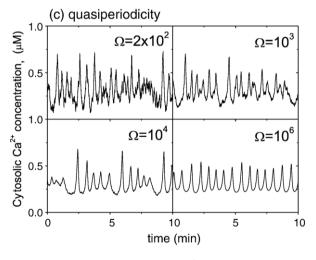


Fig. 7. Temporal evolution of three cytosolic Ca^{2+} oscillatory types for different molecule numbers Ω . The other parameter values corresponding to three oscillatory types are listed in Table 4.

appeared at moderate Ca^{2+} concentration states, and a tiny peak at highest concentration state also existed. For the chaos type, when $10^3 < \Omega < 5 \times 10^5$, there are visibly two peaks in the histograms of Ca^{2+} concentration; it can be seen that the height of peak at low concentration is higher than that at high concentration first (e.g. $\Omega = 10^3$); with the increase of molecule

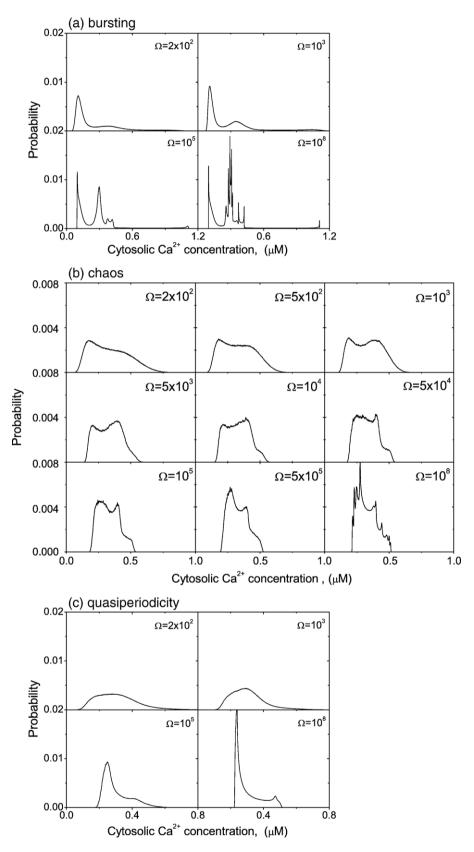


Fig. 8. Histograms of complex cytosolic Ca^{2^+} concentration for different molecule numbers Ω . The bin-size is 0.001 μ M. The other parameter values corresponding to various complex cytosolic Ca^{2^+} oscillatory types are listed in Table 4.

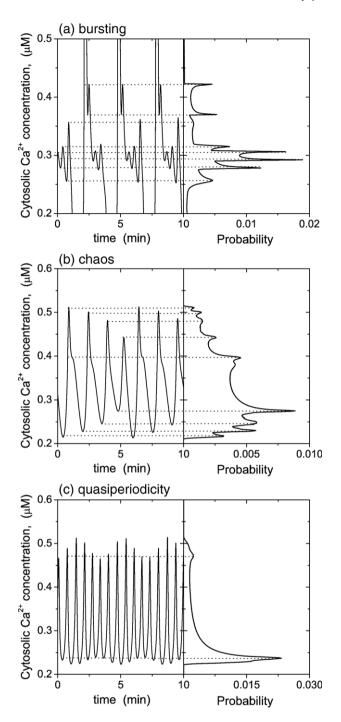


Fig. 9. Temporal evolution of cytosolic Ca^{2+} concentration (left) and corresponding histogram of Ca^{2+} concentration amplitude (right) when $\Omega=10^8$. The other parameter values corresponding to various oscillatory types are listed in Table 4.

number, the height of peak at low concentration becomes lower than that at high concentration (e.g. $\Omega = 5 \times 10^3$); when the molecule number is continuously increased, however, the height of peak at low concentration is higher than that at high concentration again (e.g. $\Omega = 5 \times 10^5$). For the quasi-periodicity type, there is only one peak located at low Ca²⁺ concentration in the histograms of amplitude of cytosolic Ca²⁺ concentration for different molecule numbers, and only a tiny peak appeared at high Ca²⁺ concentration when the molecule number is very large (e.g. $\Omega = 10^8$).

The above results also show that, when the molecule number Ω becomes large, the property of the histogram of cytosolic Ca^{2+} concentration for various complex Ca^{2+} oscillatory types is different. Fig. 9 shows the time series of the three Ca^{2+} oscillatory types and the corresponding histogram of Ca^{2+} concentration for $\Omega = 10^8$. For the bursting and chaos types, it

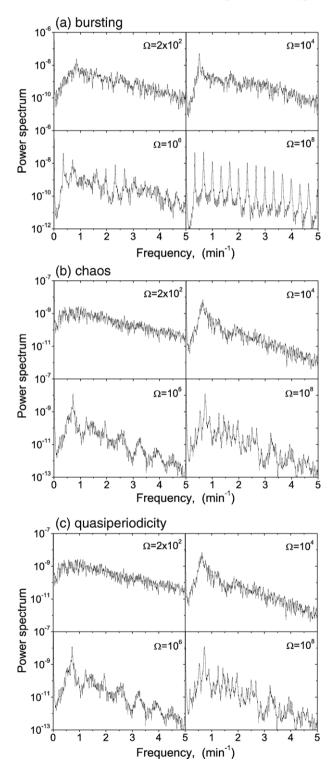


Fig. 10. Power spectrum of various complex cytosolic Ca^{2^+} oscillatory types for different molecule numbers Ω . The other parameter values corresponding to three complex Ca^{2^+} oscillations are listed in Table 4.

is found that each extremum (including maximum and minimum) of concentration of cytosolic Ca^{2+} oscillations corresponds to a peak in the histogram of Ca^{2+} concentration, and the most probability appeared during the bursting plateau level (i.e. $Z=0.292~\mu M$) for the bursting type, while it is at the largest minimum of Ca^{2+} concentration (i.e. $Z=0.275~\mu M$) for the chaos type. For the quasi-periodicity type, however, there are only two peaks (located at low Ca^{2+} concentration and high Ca^{2+} concentration, respectively) in the histogram of Ca^{2+} concentration, and the most probability is around the minimum of cytosolic Ca^{2+} concentration (i.e. $Z=0.237~\mu M$).

The power spectra of various complex cytosolic Ca2+ oscillatory types for different Ω are plotted in Fig. 10. A Welch window function is used during the estimation of power spectrum. The time series of stochastic cytosolic calcium concentration Z contains 2^{16} data points, and the smoothed curves of power spectrum are obtained by nearest averaging over 5 points. When the molecule number is very small (e.g. $\Omega = 2 \times 10^2$), or the intrinsic fluctuation of the system is very large, high irregularity of the intracellular Ca²⁺ oscillations shows up both in the amplitude and in the time interval between successive Ca²⁺ spikes for various Ca²⁺ oscillatory types (see Fig. 7), and the three oscillatory types are hard to distinguish. On the other hand, Fig. 10 shows that there is a peak of power spectrum for various types of complex oscillations for $\Omega = 2 \times 10^2$; the frequency corresponding to the maximum of power spectrum is called as the optimal frequency of cytosolic Ca²⁺ stochastic oscillations. With the increase of molecule number, the optimal frequency is shifted from high frequency to low one, meanwhile, the natural frequencies of system (all the frequencies discussed here are multiples of the smallest natural frequency, e.g. $\Omega = 10^8$) become distinguishable for bursting; the frequency corresponding to the unstable limit cycle in the case of macroscopic kinetics [46] becomes discernible for both chaos and quasi-periodicity.

4. Conclusions

Based on the model of intracellular Ca2+ oscillation with self-modulation of IP₃ signal [46] to describe both simple and complex intracellular calcium oscillations, the mesoscopic stochastic differential equations for the intracellular Ca²⁺ oscillation have been derived by using the Gillespie's method (or the chemical Langevin equation) [75] in this paper, which provide us with a convenient mathematical method to study the intrinsic fluctuation in cellular or subcellular level. It should be stressed that the mesoscopic stochastic kinetics for the intracellular Ca²⁺ oscillations is valid only under two conditions being satisfied, one is the macroscopically infinitesimal time increment must be of short enough duration that the propensity of each reaction term does not change significantly during the macroscopically infinitesimal time increment, the other is the macroscopically infinitesimal time increment must be of long enough duration that each reaction channel fires several times during the macroscopically infinitesimal time increment.

The effects of the finite biochemical reaction molecule number on both simple and complex cytosolic Ca²⁺ oscillations have been theoretically studied through numerical computation, respectively. In the case of simple intracellular ${\rm Ca^{2^+}}$ oscillation, with the increase of molecule number, it is found that the coherence resonance or autonomous resonance phenomena can occur for some external stimulation parameter values. In the cases of complex intracellular ${\rm Ca^{2^+}}$ oscillations, the histograms of ${\rm Ca^{2^+}}$ oscillations are different for each typical complex ${\rm Ca^{2^+}}$ oscillatory behavior in finite biochemical reaction molecule number.

The physiological information can be encoded by both the frequency and the amplitude of cytosolic Ca²⁺ oscillations; the above results might be helpful to understand how the intrinsic fluctuation due to the finite biochemical reaction molecule number in live cell affect the cytosolic Ca²⁺ relaying of the information within cells to regulate their activity. The above results about the effects of finite biochemical reaction molecule number on cytosolic Ca2+ oscillations are obtained at given external control parameter, that is, the extracellular stimulation is a constant. However, in a live cell system, the random extracellular agonists such as hormones and neurotransmitters are unavoidable. Therefore, we expect that more novel physical and richer physiological phenomena in the cellular and subcellular level could be observed when the random extracellular stimulations or the external fluctuations are considered, and this is our further work.

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