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# **Fast Kinetics of Calcium Signaling and Sensor Design**

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## **Abstract**

Fast calcium signaling is regulated by numerous calcium channels exhibiting high spatiotemporal profiles which are currently measured by fluorescent calcium sensors. There is still a strong need to improve the kinetics of genetically encoded calcium indicators (sensors) to capture calcium dynamics in the millisecond time frame. In this review, we summarize several major fast calcium signaling pathways and discuss the recent developments and application of genetically encoded calcium indicators to detect these pathways. A new class of genetically encoded calcium indicators designed with site-directed mutagenesis on the surface of beta-barrel fluorescent proteins to form a pentagonal bipyramidal-like calcium binding domain dramatically accelerates calcium binding kinetics. Furthermore, novel genetically encoded calcium indicators with significantly increased fluorescent lifetime change are advantageous in deep-field imaging with high light-scattering and notable morphology change.

# **Introduction**

Calcium  $(Ca^{2+})$ , a second messenger and the most ubiquitous signaling molecule, plays an important role in regulating various biological functions in living organisms (Figure 1A). The time scale of calcium ion flow varies from milliseconds in muscle contractions to days for fertilization and development (Figure 1B) [1]. Rapid calcium signaling regulates calcium channels, excitation-contraction coupling, action potential, calcium sparks, and release of neurotransmitters (Figure 1A). Voltage gated calcium channels (VGCCs) exhibit a high open and close frequency and deliver fast calcium movement through a hydrophilic path in response to plasma membrane voltage changes, allowing precise calcium signaling within milliseconds [2, 3]. During channel activation, calcium concentration is estimated to be hundreds of micromolar within several nanometers from the mouth of the channels, generating  $Ca^{2+}$  microdomains. A high  $Ca^{2+}$  gradient is generated between the microdomain and bulk cytosol [4, 5].

In muscle cells, electrical stimuli applied to the plasma membrane can be converted to muscle contraction by a process known as excitation-contraction coupling (EC coupling). In

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skeletal muscle, an action potential activates the dihydropyridine receptor (DHPR) anchored in the T tubule of the sarcolemma. DHPR then physically interacts with ryanodine receptors (RyR) expressed in the sarcoplasmic reticulum (SR) membrane to induce SR calcium release; this interaction occurs within milliseconds. After stimulation, a transient asymmetric calcium spike lasting several to tens of milliseconds occurs in the cytosol, with a fast calcium recovery phase due to SERCA pump refilling of SR calcium and buffering effects of calcium binding proteins in the cytosol [6].

The VGCC is transiently activated after the initial  $Na^+$  influx and  $K^+$  efflux in cardiac muscles, forming a plateau and a sequential slow decayed phase of membrane potential lasting for about 200 ms, much longer than that of skeletal muscle or neurons lasting for only 2–4 ms. This limits the firing rate up to several Hz, preventing the tetanus contraction of cardiac muscles. The fast calcium influx through the calcium channel triggers SR calcium release through calcium-induced calcium release (CICR) to further elevate cytosolic calcium before decreasing. The  $Ca^{2+}$  influx is terminated by closing of the VGCC with cytosolic calcium pumped back into the SR by the SERCA pump or extruded to the extracellular space by the sodium-calcium exchanger (NCX) [7]. A normal contracting cardiac muscle cell exhibits a train of cytosolic calcium spikes with the time to peak around tens of milliseconds, and a decay phase within hundreds of milliseconds.

Calcium sparks, elementary events of the CICR through the RyR in cardiac EC coupling, were discovered by fast fluorescence imaging [8]. The opening of the RyRs in cardiac or skeletal muscle cells produces calcium transients with 10 ms to peak and 20 ms half-decay, restricted around 2 μm. Activation of numerous RyRs produces multiple simultaneous calcium sparks, ranging from 50 to 5000 in a cell [9], which is regulated by the SR calcium content. The summation of the sparks generates the cytosolic calcium change. The counterpart of the calcium sparks are  $Ca^{2+}$  blinks, the transient decrement of  $Ca^{2+}$  in SR exhibiting similar fast kinetics and a much smaller region.

The neurotransmitter released from the presynaptic vesicles [10] triggered by presynaptic calcium channel activation will induce the postsynaptic receptors, for the synaptic transmission. Calcium microdomains and high calcium gradients are formed around the presynaptic calcium channels during the sub-millisecond calcium influx. In varying cell types, the local calcium concentration ranges from 10 μM to 200 μM for membrane fusion. Additionally, numerous calcium sensor proteins on the vesicle membrane surface are activated throughout the neurotransmitter release process [11]. Calcium dependent inactivation regulates the rapid termination of the calcium influx within  $1-2$  ms [12].

#### **Need for calcium sensors with fast kinetics**

The link between abnormal calcium channels and pathology requires precise tools to study them. Fluorescence microscopy with calcium sensors is indispensable in understanding live cell processes with high spatiotemporal resolution. Development of cell permeable calcium dyes has contributed greatly to our understanding of intracellular calcium signaling [1, 13]. Dyes such as Fluo-5N with fast off rates are commonly used to monitor calcium dynamics (Table 1). However, calcium dyes without targeting are not suitable for microdomain

imaging. The current genetically encoded calcium indicators (GECIs) possessing the EFhand motif show slow kinetics of signal decay (Table 1) that hamper their applications in probing fast calcium transients, especially in neuronal and skeletal muscle cells. Most calcium dyes with high affinity can only perfuse up to 80 nm from the channel, and are unable to measure tens of micromolar calcium in calcium microdomains. Imaging of calcium sparks should be deconvoluted with equations for intrinsic signals, and significant modification of the decay curve [14]. A similar convolution was applied for skeletal muscle myoplasmic calcium measurement, for which an initial peak is sometimes missing during the long-last pulse stimulation [15]. Furthermore, the complete decay of calcium in each spike was not measured in high frequency action potential stimulation, which is different from the membrane voltage changes. Rather a single transient peak was recorded for both single and multiple action potentials [16]; summation of calcium peaks is possible but not clear. This separation or summation of calcium spikes can be influenced by the choice of sensor [17, 18]. Therefore, there is a strong demand for tools to image fast transient or highfrequency calcium signaling generated by the activity of numerous calcium channels and receptors. GECIs with fast on and off rates and simple quantitation are required to monitor rapid calcium signaling in microdomains.

## **What has been done in the field?**

Calcium imaging has evolved from aequorin and calcium dyes to GECIs [19]. Figure 1C summarizes the structures of the major classes of calcium indicators. Calcium dyes based on BAPTA are mainly applied in bulk cytosol due to their strong binding affinity. Relatively weak affinity dyes are used to measure calcium concentration in internal calcium stores containing high calcium [20], but require plasma membrane permeabilization, an obstacle for intact live cell imaging. Novel strategies have been applied to improve the targeting capability of calcium dyes [21], but they are not as practical as GECIs. The kinetics of calcium sensors are highly dependent on the calcium binding protein used as the sensor and the target peptides used. Both calmodulin and troponin C based sensors bind calcium cooperatively between four EF-hands with a fluorescence change resulting from the calcium induced global protein conformational change [22–26]. These types of calcium sensors have been successfully applied to measure cytosolic calcium due to fast on rates [27–29]. Applying these sensors to measure rapid calcium release from the ER is limited by their slow off rate [30]. A fast GCaMP sensor has recently been reported to have a 20 fold improvement in the  $k_{off}$  rate [31], and progress has also been made for red fluorescent sensors [32]. However, the slow off rates of these sensors may originate from the coupled EF hand calcium binding motifs and the subsequent large global conformational change [33–36]. Multiple binding processes are introduced by modification of multiple calcium binding sites in calmodulin and troponin C raising additional challenges for quantitative measurements of rapid calcium dynamics by the developed sensors.

# **CatchER (a GFP-based Calcium sensor for detecting high concentrations in the high calcium concentration environment such as ER)**

A previous calcium sensor—G1 reported by our lab was created by grafting an EF-hand motif into EGFP and exhibited ratiometric change in response to calcium but its kinetics

were limited by an intermediate state [37]. The design of CatchER with a fast off rate was achieved based on key determinants for fine-tuning  $Ca^{2+}$  binding affinity and  $Ca^{2+}$ -induced conformational changes and the established chromophore properties of fluorescent proteins. We reasoned that  $Ca^{2+}$  sensors with fast fluorescence response can be better designed by coupling  $Ca^{2+}$  binding sites directly to the chromophore rather than relying on stretched protein-protein interaction to modulate chromophore conformation. First, four or five oxygen ligand atoms from protein residues (typically, carboxyl groups of D, E, N, Q) are situated in the pentagonal bipyramidal geometry [38] to form a calcium binding site similar to natural calcium binding proteins [39–42]. A single calcium binding site without cooperativity from protein coupling is important for fast off rate. Second, appropriate choice of residue charge and type can fine-tune  $Ca^{2+}$  binding affinity and metal selectivity [43, 44]. Third, diffusion-limited access of  $Ca^{2+}$  to the site requires good solvent accessibility [45]. Fourth, propagating  $Ca^{2+}$ -induced, local conformational and electrostatic changes to the chromophore can be achieved by proper location of the charged ligand residues with respect it [46, 47]. Fifth, these changes must occur rapidly—more rapidly than the rate of conversion from neutral to anionic state ascribed to these chromophores [37, 48]. Sixth, the created binding site must not interfere with the chromophore's synthesis and formation. We have designed CatchER by introducing five acidic residues to form a hand-like metal binding pocket in EGFP (Figure 2). The calcium binding site exhibits a pentagonal bipyramidal geometry and spreads across three antiparallel beta-sheets on the surface of EGFP. This designed calcium binding pocket is in close proximity to the hydrophobic chromophore, with hydrogen bond interaction, but facing outwards with high solvent accessibility. Calcium binding affects the hydrogen bond network between the binding residues and chromophore, triggering fluorescence change. Calcium binding to CatchER induces the ratiometric changes in the absorption spectra, as well as an increase in fluorescence emission at 510 nm upon excitation at both 395 and 488 nm. This novel designed calcium pocket lacks two binding ligands from its ideal coordination, thereby exhibits lower binding affinity and fast kinetics [35]. The X-ray crystal structures of CatchER have been resolved at 1.78–1.20 Å in  $Ca^{2+}$  loaded and  $Ca^{2+}$  free form, respectively (Figure 3A) [35]. Calcium binding altered the conformations of Thr203 and Glu222 associated with two forms of Tyr66 of the chromophore to induce the optical change [49]. From the X-ray crystal structure,  $Ca^{2+}$  ions were partially occupied in two locations within the extended designed binding site, suggesting the ability of  $Ca^{2+}$  ions to move around these possible locations may be responsible at least in part for the fast kinetics of metal-ion binding to CatchER [49]. A more dynamic observation was achieved by NMR [35].

Our novel GECI—CatchER has been applied to measure the ER calcium dynamics in nonexcitable mammalian cells [50]. Furthermore, having fast kinetics, CatchER was able to record the SR luminal  $Ca^{2+}$  in flexor digitorum brevis (FDB) muscle fibers during voltage stimulation (Figure 2), and successfully determined the decreased SR calcium release in aging mice [51]. Thermostablity and brightness impaired the application of the early version of GCaMP, which was dim and experienced difficulty folding at 37°C [52]. Later it was improved using site directed mutagenesis. We have further improved the optical properties of CatchER. The new version of CatchER variant has been recently developed for optimal expression at 37°C (Reddish et al., unpublished results). Recently, we have designed red

fluorescent protein based CatchER with improved pKa value (Zhuo et al., unpublished results).

For multi-color imaging with different fluorescent sensors, a red fluorescent protein-based calcium sensor (R-CatchER) has been designed to expand the imaging wavelength (Zhuo et al., unpublished results). The red fluorescent protein based ER sensor R-LAR-GECO1 was reported recently with a weaker binding affinity of 24 μM by modifying the interaction surface between CaM and M13 of the cytosolic calcium sensor R-GECO1 [53, 54]. It is interesting progress since the binding affinity decreased around 50 folds, whereas earlier it was reported as impossible to tune the binding affinity of GCaMP significantly lower by the same mutations found in Cameleon [22, 52].

#### **From Intensity to Lifetime**

Two photon fluorescence lifetime imaging is an indispensable method for understanding molecular signaling in environments with highly dynamic morphological changes and light scattering, such as the brain [55–58]. Until now, synthesized calcium indicators, such as OGB1, have single exponential lifetime decay, allowing for quantification of calcium change with high accuracy. However, there are limited reports of calcium fluorescence lifetime imaging with GECIs. A study reported that there was no significant lifetime change before and after adding  $Ca^{2+}$  to Cameleon YC3.1 that used ECFP and Venus [59], which are commonly used by other FRET-based calcium sensors. The multiple component lifetime decay of CFP makes it difficult to accurately determine the lifetime change. A fluorescent protein with a single exponential lifetime decay and high photostablity would be a good alternative for the FRET pair donor. Recently, an mTFP-Cit pair based TnC sensor was reported to exhibit significant lifetime change [60]. However, the kinetic property of this sensor was not recorded.

We have recently reported that CatchER is the first protein-based calcium indicator with the single fluorescent moiety to show the direct correlation between the lifetime and calcium binding (Figure 3C). Calcium binding increases the mean fluorescence lifetime of the deprotonated form of CatchER by 44%. Compared to wild type GFP, the site-directed mutagenesis to create calcium binding ligands disrupted the proton-transfer path and altered the optical properties. Calcium binding rescues this path and affects the fluorescence intensity (Figure 3B) [61]. Our finding provides important insights for a strategy to design calcium sensors and suggests that CatchER could be a useful probe for FLIM imaging of calcium *in situ*.

#### **Perspectives**

Fluorescent sensor based calcium imaging has dramatically increased our knowledge of cellular functions. However, there are still pressing needs for the accurate determination of fast calcium signaling besides simulation, especially in excitable cells expressing calcium channels. For further development, accelerating the kinetics and broadening the binding affinities of GECIs will compensate current imaging limitations. Looking for different calcium binding motifs other than the EF-hand for GECIs may provide a beneficial approach in overcoming these issues. For neuronal and deep field imaging with strong light-scattering,

it could be advisable to design calcium sensors with high sensitivity for fluorescence lifetime imaging. Furthermore, EGFP-based calcium sensors are generally pH sensitive to the physiological conditions, as CatchER exhibited a pKa around 6.9 in the presence of  $Ca<sup>2+</sup>$ , with a 0.7 unit decrease than apo-form. Recently reported GCaMPs displayed a more significant pKa shift from  $8~10$  in the apo-form to  $6~7.2$  in the holo-form [16]. The pH insensitive sensors such as blue fluorescent protein [53] or red fluorescent protein-based sensor (Zhuo et al, unpublished results) are superior for monitoring calcium signaling in the Golgi and lysosome. The new, advanced GECIs will significantly improve our understanding of the molecular basis of  $Ca^{2+}$  homeostasis in health and pathology.

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# **Highlights**

**•** Fast calcium signaling is mainly regulated by calcium channels.

- **•** Calcium indicators facilitate our understanding of fast calcium signaling.
- **•** It is necessary to improve the kinetics of genetically encoded calcium indicators.
- **•** CatchER with a pentagonal bipyramidal-like binding domain exhibits fast kinetics.
- **•** Fluorescent lifetime calcium imaging is advantageous for deep field imaging.



#### **Figure 1.**

Calcium signaling and fluorescent calcium sensors. (A) Fast calcium signaling regulated by the voltage-gated calcium channels (VGCC) includes EC coupling, cardiac action potential, calcium sparks and neurotransmitter release. (B) Different time scales of calcium signaling. (C) Representatives of calcium dye and genetically encoded calcium indicators. Fura-2 is synthesized calcium dye. Aequorin is used as genetically encoded calcium indicator. Cameleon and TnC sensor are composed by calmodulin and troponin C containing four EFhands, respectively, flanked with FRET pair. GCaMP, GECO and R-CaMP are single fluorescent protein based sensor with calmodulin. CatchER is the single fluorescent protein with a designed calcium binding site on the surface of EGFP.



#### **Figure 2.**

CatchER tracks  $Ca^{2+}$  release and uptake kinetics in response to single or repetitive muscle fiber depolarization. (A) SR  $Ca^{2+}$  transients were recorded in FDB fibers expressing either CatchER or D1ER elicited by 100-ms command pulses to 20 mV under whole-cell patchclamp. CatchER's and D1ER citrine's fluorescence is illustrated to compare their amplitude and kinetics. D1ER citrine emission (535 nm) was chosen over the cyan fluorescence protein (485 nm) because of its larger amplitude. The D1ER excitation wavelength was set at 436 nm. D1ER's and CatchER's fluorescence was recorded using a spectrofluorometer and confocal microscope in the line-scan mode (see above), respectively. The dashed line indicates the baseline. (B) Time to peak, half recovery time, and response amplitude normalized to basal fluorescence were analyzed for CatchER and D1ER citrine fluorescence ( $F/F$ ). Asterisks indicate a statistically significant difference ( $P < 0.01$ ). Values are mean ± SEM for 19 and 13 fibers expressing CatchER or D1ER FDB, respectively. Transient changes in CatchER's fluorescence response to various 20-mV command pulses: (C) a 10 ms pulse, (D) five 10-ms pulses at 10 Hz, (E) a 10-ms pulse at 3.3 Hz, (F) a 10-ms pulse at 1.6 Hz, (G) a 100-ms pulse, (H) 100-ms pulse at 5 Hz, and (I) 50-ms pulse at 3.3 Hz. Images provided with kind permission from Springer Science and Business Media and Tang et al., *PNAS* Copyright © 2011.



#### **Figure 3.**

Structure and fluorescence lifetime of CatchER (A) The X-ray crystal structure of  $Ca^{2+}$ -CatchER (PDB ID: 4L12). The residues shown in stick are the designed calcium binding site. Residues 164-168 were shown with 50% transparency. The residue shown in green is the chromophore. The atom shown in grey is the calcium atom. The oxygen atoms and nitrogen atoms are indicated in red and blue, respectively. (B) The proton wire observed in the crystal structure of  $Ca^{2+}$ -CatchER (PDB ID: 4L12). The H-bonds are shown in the black dash line with the cut off of 3.5. Only side chains were shown in S72, T203, S205 and E222. (C) Fluorescence decay traces of apo-CatchER (dash line) and CatchER supplemented with 10 mM Ca2+ (Holo, solid line). CatchER was excited at 372 nm and emitted at 510 nm in the time range of 15 ns. Ex and Em are short for excitation and emission, respectively. Reprinted (adapted) with permission from Zou, et al., *J. Phys. Chem. B,* 2014. Copyright © 2015 American Chemical Society.

#### **Table 1**

#### Kinetics of recent major calcium indicators.



<sup>\*</sup>The unit is  $M^{-n}s^{-1}$ . n(CH-GECO2.1)=0.58 and n(R-GECO1)=1.6.