Calcium spikes accompany the cleavage furrow ingression and cell separation during

fission yeast cytokinesis

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Running title: Calcium spikes during yeast cytokinesis

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Abstract

The role of calcium signaling during cytokinesis has long remained ambiguous. The studies of

embryonic cell division discovered that calcium concentration increases transiently at the

division plane just before the cleavage furrow ingression, leading to the hypothesis that these

calcium transients trigger the contractile ring constriction. However, such calcium transients

have only been found in animal embryos and their function remains controversial. Here we

explored cytokinetic calcium transients in the model organism fission yeast. We adopted

GCaMP, a genetically encoded calcium indicator, to determine the intracellular calcium level.

We validated GCaMP as a highly sensitive calcium indicator which allowed us to capture the

calcium transients stimulated by osmotic shocks. To identify calcium transients during

cytokinesis, we first identified a correlation between the intracellular calcium level and cell

division. Next, we discovered calcium spikes at the start of the cleavage furrow ingression and

the end of the cell separation using time-lapse microscopy to. Inhibition of these calcium spikes

slowed down the furrow ingression and led to frequent lysis of the daughter cells. We conclude

that like the larger animal embryos fission yeast triggers cytokinetic calcium transients which

promote the ring constriction and daughter cell integrity (194).

Highlight summary for TOC:

Calcium rises transiently at the division plane during embryonic cell cytokinesis, but the

conservation and function of such calcium transients remain unclear. We identified similar

calcium spikes during fission yeast cytokinesis and demonstrated that these spikes promote the

contractile ring constriction and the daughter cell integrity (257).

Introduction

Calcium is an essential secondary messenger in many cellular processes but its role during cytokinesis, the last stage of cell division, remains ambiguous. Eukaryotic cells maintain their intracellular calcium at much lower concentration than that of their extracellular environment. A transient increase of the free calcium level, through either release from the intracellular storage or influx through the plasma membrane, triggers a number of essential calcium signaling pathways (for a review see (Clapham, 2007)).

Although the importance of calcium to cytokinesis had long been known (Arnold, 1975), the calcium transients accompanying cytokinesis were discovered much later and remain poorly understood. Fluck et al. first observed two localized calcium waves at the cell division plane of medaka fish embryos during cytokinesis (Fluck et al., 1991). The first wave initiates just before the cleavage furrow ingression while the second wave appears after the cell separation. Followed-up studies discovered similar localized increase of calcium in other animal embryos, including those of zebra fish *Danio rerio* and African frog *Xenopus laevis* (Chang and Meng, 1995; Miller et al., 1993; Noguchi and Mabuchi, 2002). However, it remains unexplored whether such calcium transients are universal among all eukaryotic cells. Neither is there a consensus on the exact molecular function of these transients. The prevailing hypothesis (Chang and Meng, 1995; Fluck et al., 1991) holds that these transients can activate myosin light chain kinase (MLCK) to promote the activity of myosin II motor in the contractile ring and trigger the ring constriction (Craig et al., 1983; Scholey et al., 1980). This theory draws a parallel between the role of calcium in cytokinesis to that in smooth muscle constriction (Nishimura et al., 1990). This proposal, although attractive, has remained unproven even in the animal embryos. Adding to the ambiguity, inhibition of the cytokinetic calcium transients in *X. laevis* embryos has

produced conflicting results (Miller et al., 1993; Noguchi and Mabuchi, 2002; Snow and

Nuccitelli, 1993). To date, both the nature and molecular function of the cytokinetic calcium

transients remain undetermined.

Fission yeast *Schizosacchromyces pombe* has emerged as an excellent model organism

for the study of cytokinesis in the last twenty years. The molecular mechanism of cytokinesis in

this unicellular organism has been largely conserved in higher eukaryotes (for review see

(Pollard and Wu, 2010)). Like animal cells, fission yeast assembles an actomyosin contractile

ring at the cell division plane (Wu et al., 2003), a process mediated by essential cytokinetic

proteins such as Myo2p, Cdc12p and Adf1p (Balasubramanian et al., 1998; Chang et al., 1997;

Chen and Pollard, 2011). The assembly of contractile ring is further promoted by a continuous

polymerization of actin filaments in the ring (Courtemanche et al., 2016). As in higher

eukaryotes, the actomyosin ring constricts to drive the cleavage furrow ingression, assisted by a

growing septa (Laplante et al., 2015; Mishra et al., 2013). The terminal step of cell separation

(for review see (Garcia Cortes et al., 2016; Sipiczki, 2007) requires degradation of the septa

(Cortes et al., 2002; Cortes et al., 2012; Liu et al., 1999; Munoz et al., 2013) as well as proper

regulation of the turgor pressure (Atilgan et al., 2015; Proctor et al., 2012).

Although it still remains a challenge to quantify the free calcium concentration in a live

cell, the rapid development of genetically encoded calcium indicators (GECI) over the last

twenty years has made it far easier than before. Compared to the more widely used synthetic

probes, GECIs can be expressed in vivo without intrusive probe-loading procedure. This is

particularly advantageous in such genetically modifiable systems as fission yeast. Although most

GECIs still trail the chemical probes in their sensitivity, some of them such as GCaMP (Nagai et

al., 2001), which we chose for this study, have come close. This single fluorophore sensor

consists of three domains: a circularly permuted enhanced green-fluorescent protein (cp-EGFP),

a calcium-binding calmodulin motif and a M13 peptide of human myosin-light chain kinase

(MLCK). Binding of calcium to the calmodulin motif dissociates the M13 peptide from cp-

EGFP, increases the fluorescence of GCaMP significantly (Nakai et al., 2001). Further

modifications of GCaMP have substantially reduced its response time and enhanced its signal-to-

noise ratio (Chen et al., 2013; Tian et al., 2009). We picked one of the most sensitive variants,

GCaMP6s (Chen et al., 2013), to explore the potential calcium transients during fission yeast

cytokinesis.

Few studies have examined the calcium level of fission yeast at the single cell level, even

though its calcium signaling pathways have been characterized extensively. Two central

calcium-signaling molecules, calmodulin and calcineurin, have both been identified (Moser et

al., 1997; Takeda and Yamamoto, 1987; Yoshida et al., 1994). As in animal cells, they play a

critical role in a plethora of cellular processes including cytokinesis in fission yeast. In particular,

the essential calmodulin Cam1p is required for cell division, underlying the importance of

calcium-signaling in this model organism (Moser et al., 1997). Past studies have mostly

employed bulk assays to determine the average calcium level of many cells, using either

chemical or indirect probes (Deng et al., 2006; Ma et al., 2011). As a result, the regulation of

calcium at the sub-cellular level during many processes including cytokinesis has been left

undetermined.

In this study we first explored the feasibility of GCaMP-based calcium imaging in fission

yeast. We then deployed this probe to capture the calcium transients triggered by various stimuli

to validate its effectiveness. Finally, we identified the cytokinetic calcium spikes and examined

their importance to cytokinesis.

Results

Expression and localization of the genetically calcium indicator GCaMP in fission yeast

To probe the intracellular calcium level of fission yeast cells with fluorescence microscopy, we expressed a single copy of the GCaMP coding sequence at an endogenous locus. A strong constitutive ADH promoter drove the expression of GCaMP6s (referred to as GCaMP) (Fig. 1A). To measure the expression and sub-cellular localization of this calcium indicator, we tagged its C-terminus with another fluorescence protein mCherry. GCaMP-mCherry localized uniformly throughout the cytoplasm and the nucleus but it was excluded from the vacuoles (Fig. 1B). The reporter nevertheless was slightly enriched in the nucleus, compared to the cytoplasm (ratio = 1.5 ± 0.1, average ± S.D., n = 66). The ratio of GCaMP to mCherry intracellular fluorescence varied very little among the cells (Fig. 1C), indicating that the expression of this reporter is homogenous. We concluded that GCaMP can be expressed homogenously in the fission yeast cells and this calcium indicator localizes throughout the intracellular space.

We next determined whether the expression of GCaMP perturbs any cellular functions, a potential concern for endogenously expressed reporters. Overall the GCaMP expressing cells exhibited no apparent morphological defects. Their length and width were similar to the wild-type cells (Fig. 1D). Their mitotic and septation indexes were also normal, similar to the wild type (Fig. 1E-F). Moreover, these GCaMP expressing cells did not exhibit any hyper-sensitivity to either calcium, or sorbitol, or EGTA, all of which likely perturb the intracellular calcium homeostasis (Fig. S1). Lastly, cytokinesis in these GCaMP-expressing cells were unperturbed, as determined by time-lapse fluorescence microscopy. In the GCaMP-expressing cells, the contractile ring assembly and maturation took 44 ± 5 mins (average ± S.D., n = 31), starting from the appearance of precursor nodes, similar to the wild-type cells (43 ± 4 mins, n = 40). This was

followed by the ring constriction that took 30 ± 3 mins (n = 43), same as the wild type (30 ± 3

mins, n = 70). The last stage of cytokinesis the cell separation required 26 ± 4 mins (n = 29) in

the GCaMP-expressing cells, not significantly different from the wild type (25 ± 3 mins, n = 67).

These quantitative measurements indicated that expression of GCaMP dose not interfere with

cytokinesis either. Since GCaMP does not produce any discernable artifacts, we combined our

GCaMP-expressing strain with quantitative microscopy to examine the intracellular calcium

level of fission yeast cells.

Calcium transients in fission yeast cells

To validate GCaMP as an effective calcium indicator, we first determined whether an increase of

extracellular calcium alters the fluorescence of GCaMP-expressing cells. In the rich YE5s media,

addition of 30mM $CaCl_2$ increased average fluorescence of the GCaMP expressing cells by

about three folds, compared to those in YE5s (Fig. 2A-B). The increased extracellular calcium

concentration also resulted in more frequent fluctuations of the intracellular GCaMP

fluorescence (Fig. 2C-D and Supplemental Movie S1). Thus, GCaMP-based imaging is fully

capable of detecting alterations of calcium concentration.

Next, we employed the GCaMP-based time-lapse imaging to capture rapid changes of

intracellular calcium level, the calcium transients. The resting fission yeast cells produced only

sporadic calcium transients (Fig. 2C). We applied various stimuli through microfluidics to

produce more transients. First, we used hypo-osmotic shock, one of the best-known stimuli of

calcium transients in yeast (Batiza et al., 1996; Denis and Cyert, 2002). We found highly

synchronized calcium transients, captured by an increase of GCaMP fluorescence, in the cells

under such shock (Fig. 3A and Supplemental Movie S2). The intracellular calcium rose quickly

following the shock and peaked 2 mins after the osmotic shock on average. These calcium

transients accompanied the expansion of cell volume closely (Fig. 3B-C). Average amplitudes of

the calcium increase were proportionally to the strength of hypo-osmotic shocks (Fig. 3D).

Secondly, we applied hyper-osmotic shocks to the cells. This surprisingly did not elicit

detectable calcium transients (Fig. 3E and Fig. S2A-B). Lastly, we abrupted increased the

extracellular calcium concentration to stimulate the cells. This also provoked strong calcium

transients, but they (QC) appeared only ~5 mins after the infusion of calcium (Fig. 3F). As a

result of the stimulation, the average intracellular calcium concentration remained elevated after

more than 20 mins. This response to the external calcium shock confirmed a similar observation

made through the bulk assays (Deng et al., 2006). Our results demonstrated that the GCaMP-

based time-lapse imaging can capture the calcium transients in fission yeast cells, making it

entirely feasible to identify the cytokinetic calcium transients.

Identification of calcium spikes during cytokinesis

As the first step to determine the link between calcium and cytokinesis, we analyzed the

intracellular calcium level in a large number of asynchronized cells. Dividing fission yeast cells

can be identified through their increased length. These rod-shaped cells grow by tip extension,

starting to grow from a length of ~7 μm after birth. They expand throughout interphase until

stopping at a length of ~14 μm at the mitotic entry (Mitchison and Nurse, 1985). We found that

the calcium level was constant among most cells regardless of their length (n = 407) (Fig. 4A).

However, the calcium centration of a few cells was significantly higher than the average

(>110%). The length distribution of these "outliers" was interestingly bimodal. Two peak

fractions are ~14 μm and ~8 μm respectively (Fig. 4B), roughly equaling to the length of the

dividing and new-born cells respectively. The existence of the outlier cells with high calcium

level provided the first hint of calcium transients during cell division.

Next, we examined the temporal regulation of intracellular calcium in the dividing cells

through time-lapse microscopy. To facilitate our analysis, these cells expressed both GCaMP and

well-established fluorescence protein markers of cytokinesis. First, we looked closely at the

calcium level during mitosis, by imaging the cells expressing both GCaMP and Sad1p-GFP, a

marker of spindle pole bodies (SPB). In fission yeast, the separation of SPBs is concomitant with

the start of prometaphase and anaphase B ends ~30 mins after (Fig. S3A) (Wu et al., 2003).

During this period, we captured very few calcium transients. About 34 mins after the separation

of SPBs, the calcium level started to rise quickly throughout the intracellular space, which we

termed calcium spikes. These spikes appeared to be well-synchronized, coinciding with

telophase as well as early cytokinesis (Fig. S3B-C). Next, we determined whether these calcium

spikes initiated exactly at the beginning of furrow ingression as those calcium transients did in

the fish embryos (Chang and Meng, 1995; Fluck et al., 1991). We imaged the cells expressed

GCaMP and a contractile ring marker Rlc1p-tdTomato, the regulatory myosin light chain (Fig.

5A). Our movies recorded very few calcium spikes during the assembly and maturation of the

contractile ring (Fig. 5A-B). In comparison, our time-lapse microscopy consistently captured

calcium spikes accompanying the start of ring constriction in most dividing cells. The spikes also

appeared frequently during the constriction. We called them "constricting spikes" (Fig. 5B-C and

Supplemental Movie S3). In these cytokinetic cells, the average intracellular calcium level

increased ~1.7 fold within 1 ± 3 min (average $\pm$ S.D., n = 64) of the start of the cleavage furrow

ingression (Fig. 5D).

Besides the dividing cells, the intracellular calcium of new-born cells appeared to be high

as well, prompting us to examine the calcium spikes during cell separation. Although few

calcium spikes were captured following the ring closure and before the cell separation, spikes

were found in most cells following the cell separation (Fig. 6A). These separating spikes were

concomitant with the initial appearance of new ends. They distributed asymmetrically between

the two daughter cells (Fig. 6B-C). Among the cells (n = 73) in which at least one separating

spike was captured, 56% triggered the spike in only one daughter cells. The other 44% triggered

the calcium spikes that oscillated between the two daughter cells (Fig. 6B-C). On average, the

separating spikes increased the intracellular GCaMP fluorescence by ~1.7 folds (Fig. 6D-E). The

calcium level peaked ~2 mins after the end of cell separation (Supplemental Movie S4). We

observed similar constricting and separating spikes in the dividing cells inoculated in the

synthetic EMM5s media (Fig. S3D-E), further confirming the existence of these cytokinetic

spikes.

To better understand the spatiotemporal regulation of these cytokinetic calcium spikes,

we characterized them by fast imaging. We increased the frequency of time-lapse acquisition by

40 folds, from one frame every two minutes to one frame every three seconds. We only captured

a limited number of constricting and separating spikes in this way (Fig. 7A and D). Close

examination of the spikes found that they increased intracellular calcium level most significantly

at the tip, the peri-nuclear space, and the division plane (Fig. 7B and E). All of them exhibited a

lifespan of 50-200 seconds (Fig. 7C and F). Their amplitudes were highly variable (Fig. 7C and

F). Some constricting spikes exhibited a more than 10-fold increase of GCaMP fluorescence

above the baseline level (Fig. 7C). We concluded that the cytokinetic calcium spikes rise

throughout the cytoplasm and they are highly heterogeneous.

Functions of the fission yeast cytokinetic calcium spikes

The close temporal correlation between the calcium spikes and cytokinesis prompted us to

investigate whether these spikes are required for cytokinesis. To this end, we depleted the

extracellular calcium from the media, hypothesizing that the calcium influx contributes to the

spikes. The calcium depletion was only temporary to minimize the effects on the other calcium-

dependent processes. The cells were observed in either EMM5s media without any supplemented

calcium or YE5s media supplemented with low concentrations of EGTA. At such concentrations,

we found that EGTA did not inhibit cell growth strongly (Fig. S4A). In the calcium-depleted

media, the average intracellular GCaMP fluorescence decreased only slightly, ~13% in YE5s

plus 2 mM EGTA and ~3% in the calcium-free EMM5s media (Fig. 8A-B). However, the

fraction of outlier cells decreased significantly under either condition (Fig. 8A-B). Although the

calcium spikes were no longer as visible under either conditions, quantitative image analysis

revealed that the total number of cytokinetic spikes was not reduced eve though their amplitudes

decreased dramatically (Fig. 8C-D and Fig. S4B-E). We concluded that the cytokinetic calcium

spikes can be inhibited by depleting extracellular calcium from the media.

We next examined cytokinesis in the cells whose calcium spikes are inhibited through

time-lapse microscopy. We found that the assembly and maturation of the contractile ring took

slightly longer time in the calcium-free EMM media, compared to the EMM media (Fig. 9A and

C). However, constriction of the contractile ring slowed down much more dramatically (30%,

Fig. 9E and Supplemental Movie S5). Similarly, 1-2 mM EGTA did not alter the assembly and

maturation of the ring significantly (Fig. 9B and D) but the constriction rate was inhibited by

EGTA in a dosage-dependent manner. The rate was down ~50% in YE5s plus 2mM EGTA,

compared to YE5s (Fig. 9E). When EGTA concentration was further increased to 5mM, most

contractile ring (90%, n = 30) disintegrated without completing the constriction (Fig. S5A-B).

We also examined the cell separation when the calcium spikes are inhibited. Surprisingly, the

duration of cell separation remained normal when the extracellular calcium was depleted (Fig.

S5C-D). Although no daughter cells lysed in the calcium-free EMM5s media, 2 mM EGTA

induced more than one third (35 ± 5 %, n = 103) of the daughter cells to lyse following the

separation (Fig. 9G-H). More cells lysed when the concentration of EGTA increased to 5 mM

(data not shown). Further depletion of intracellular calcium by deletion of *pmr1* resulted in 54%

of daughter cells lysed in the calcium-free EMM5s media even though no mutant cells lysed in

the regular EMM5s. Pmr1p is a ER calcium ATPase that replenish the internal storage of

calcium (Cortes et al., 2004). We concluded that the cytokinetic calcium spikes also likely

promote the integrity of separating cells.

Discussion

Overall, we demonstrated that GCaMP-based imaging will allow fission yeast to serve as a

powerful model organism for the study of calcium transients during various cellular processes.

Using this method, we discovered the cytokinetic calcium spikes in this unicellular model

organism for the first time. These calcium transients are similar to those first uncovered in

animal embryos. They likely play a critical role in promoting the ring constriction and the

daughter cell integrity.

GCaMP-based calcium imaging of fission yeast

To our knowledge, our study is the first one to examine the feasibility of GCaMP-based calcium

imaging in fission yeast cells. We are fortunate that fission yeast is quite tolerant to the

expression of this calcium indicator. It does not perturb many cellular processes that we

examined even when it is expressed constitutively. More importantly, GCaMP is also highly

sensitive as a probe for time-lapse microscopy, allowing us to capture calcium transients at

relatively low sampling rate. Compared to synthetic calcium probes, GCaMP exhibited two key

advantages in fission yeast. First, unlike the synthetic probes such as Calcium Green (Chang and

Meng, 1995), the intracellular concentration of GCaMP can be maintained at a constant level

through the homogenous expression. Combined with quantitative microscopy, this largely

eliminated the heterogeneity of fluorescence intensities among cells. In contrast, intracellular

concentration of the synthetic probes can be highly variable among cells. This advantage was

exemplified by the identification of outliers with high calcium among a large number of cells.

The other advantage of GCaMP is its more uniform sub-cellular localization, compared to the

synthetic probes. This allows detection of calcium transients throughout the cytoplasm as well as

the nucleus.

When combined with yeast genetics, GCaMP can be a very versatile tool for studying

calcium signaling. As an example, we constructed both GCaMP and GCaMP-mCherry for this

study, both of which have the potential to be employed in the future studies. We employed

GCaMP not GCaMP-mCherry throughout this study, primarily to preserve the spectrum for

imaging the fluorescence protein markers of cytokinesis. Nevertheless, the tandem reporter can

be used for ratio-metric imaging of intracellular calcium, potentially providing higher signal to

noise ratio. Overall, application of GCaMP shall make fission yeast an excellent model organism

to study calcium transients and homeostasis in non-excitable cells.

Evolutional conservation of the cytokinetic calcium transients

Our study provides fresh evidences for the existence of calcium transients in an unicellular

organism. Similar to the earlier studies (Chang and Meng, 1995; Fluck et al., 1991; Noguchi and

Mabuchi, 2002), we employed live fluorescence microscopy to determine the temporal

correlation between the calcium transients and cytokinesis. With the availability of fluorescence

protein markers of cytokinesis, we now determined the temporal regulation of these spikes with

more precision. In addition to time-lapse microscopy, we sought out an alternative approach to

illustrate this close correlation. We analyzed just snapshots of many GCaMP-expressing cells to

reveal the relationship between cell division and intracellular calcium level. Compared to the

movies, the snapshots examined all cells regardless of their cell-cycle stage, presenting a less

biased view of the potential calcium change in them. Combined with the previous studies of

animal embryos, our results strongly support that a temporal increase of calcium during

cytokinesis may be evolutionally conserved.

The temporal regulation of fission yeast cytokinetic calcium spikes bears strong

similarities to those found in the animal embryos. The constricting spikes of fission yeast are

comparable to the first "furrowing wave" observed in the animal embryos (Fluck et al., 1991)

which initiates just as the cleavage furrow starts to ingress (Chang and Meng, 1995; Noguchi and

Mabuchi, 2002). The separating spikes are comparable to the second "zipping wave" of the

embryos, starts just when the two daughter cells separate (Fluck et al., 1991; Noguchi and

Mabuchi, 2002). Both the calcium waves in the embryos and the spikes in fission yeast can last

for minutes, far longer than other known calcium transients (Jaffe and Creton, 1998). Although

no calcium spikes have been identified yet during budding yeast cytokinesis, Carbo et al.

observed higher frequency of calcium bursts during G1 to S phase transition in synchronized

cells (Carbo et al., 2017), which may be similar to the separating calcium spikes in of fission

yeast. Overall, our study suggests that the regulatory mechanism of calcium during cytokinesis

may have been conserved as well.

There are also key differences between the calcium spikes and those embryonic calcium

waves. Chief among them is the spatial regulation of these calcium transients. In fission yeast,

the calcium spikes propagate globally throughout the cytoplasm and nucleus. In contrast, the

calcium waves of animal embryos are restricted to the cleavage furrow (Chang and Meng, 1995;

Fluck et al., 1991) or its surrounding region (Noguchi and Mabuchi, 2002). This difference may

be due to varying sensitivity of the calcium probes. It is more likely that this distinction is due to

the relatively small size of fission yeast cells, measuring only 4 μm wide and 14 μm long,

compared to the embryos of hundreds of microns. So far, we found no evidence that the

constricting spike of fission yeast triggers the contractile ring constriction, even though there are

evidences that the furrowing waves trigger the ring constriction (Chang and Meng, 1995; Fluck

et al., 1991; Miller et al., 1993). Lastly, the asymmetry and oscillation of the separating spikes

are also unique to fission yeast. The oscillating behavior of the spikes is particularly striking. It

may be linked to the unbalanced turgor pressure between the two daughter cells during the

separation.

Mechanism of the calcium spikes and their potential roles

Although the regulatory mechanism of these cytokinetic calcium spikes remains to be explored,

our data suggests that they likely draw calcium from both the influx and internal release.

Inhibition of the influx through depleting extracellular calcium significantly inhibited the spikes

but not completely. Under such condition, the cytokinetic calcium spikes still occurred albeit

with greatly diminished amplitudes. This points to the ER- or vacuole- stored calcium as the

other likely sources for the spikes. Supporting this hypothesis is our observation that EGTA

exhibited a stronger inhibitory effect to cytokinesis, compared to the calcium-free EMM

medium. EGTA may have chelated the intracellular calcium slowly over long-term incubation.

This is consistent with the lysis of *pmr1Δ* mutant cells in the calcium-free medium. In

comparison, the embryonic calcium waves draw only from the internal store (Chang and Meng,

1995; Miller et al., 1993). It remains unclear what are the ion channels mediating the cytokinetic

calcium transients. Fluck et al. has first proposed the potential role of tension-sensing calcium

channels which can be activated during cytokinesis (Fluck et al., 1991). This remains a feasible

mechanism considering increased membrane tension on the cleavage furrow. Release of internal

calcium could be through the store-operated calcium release (SOCE) channels (Chan et al., 2016;

Chan et al., 2015). Nevertheless, none of these cytokinetic ion channels has been identified.

It remains unknown how the calcium spikes promote the contractile ring constriction and

the daughter cell integrity. One likely explanation is that a transient increase of intracellular

calcium may increase the activity of type II myosin in the ring through the phosphorylation of

myosin regulatory light chain (Chang and Meng, 1995; Fluck et al., 1991). Alternatively, the

spikes could stimulate the septum biosynthesis through activating the calcium-dependent

calcineurin (Cadou et al., 2013; Martin-Garcia et al., 2018; Yoshida et al., 1994). Lastly, the

calcium spikes may adjust the turgor pressure which is required for the separation (Abenza et al.,

2015; Proctor et al., 2012). Further studies will be required to determine the mechanism of

calcium signaling during cytokinesis.

Experimental procedures

Yeast genetics

We followed the standard protocols for yeast cell culture and genetics. Tetrads were dissected

with a Spore+ micromanipulator (Singer, UK). Yeast cells were transformed with the lithium-

acetate method. The pmr1Δ mutant is a commercially available fission yeast deletion library

(Bioneer, South Korea). All the strains used in this study are listed (Table 1).

To construct the GCaMP expressing strain, we amplified the coding sequence from

pCMV-GCaMP6s (Addgene, Plasmid #40753) through PCR using primers P589 and P590. The

DNA fragment was subcloned into a previously described plasmid *pFA6a-KanMX6-Padh1-*

Tadh1 (Chen and Pollard, 2013). The GCaMP6s ORF, together with the Adh1 promoter and

terminator, were integrated into the *leu1* locus through PCR-based homologous recombination

using primers P413 and P414 (Bahler et al., 1998). The resulting strain was confirmed through

both PCR using primers P589 and P416 and the Sanger sequencing using primer P668. The

GCaMP-mCherry strain was constructed through homology-based PCR using primers P647 and

P648. The genome locus was sequenced to confirm the insertion of mCherry coding sequence.

Fluorescence microscopy

For time-lapse microscopy, the cells were first inoculated in liquid YE5s media at 25°C for two

days before they were harvested during the exponential growth phase at a density between

$5*10^6$/ml to $1.0*10^7$/ml. 20 µl of the cell culture were spotted onto the glass-coverslip (#1.5) in a

10 mm petri dish (Cellvis, USA). The coverslip was pre-coated with 50µl of 50-100 µg/ml of

lectin (Sigma, L2380) and allowed to dry for 3-4 hours in a 30°C incubator. After 10 min at

room temperature, most cells attached to the coverslip in the petri dish. 2ml of YE5s media was

then added to the dish before microscopy. When chelation of calcium in media is required, 2ml

of YE5s supplemented with EGTA was added instead.

For the experiments in the calcium-free media, we prepared the synthetic EMM media by

omitting $CaCl_2$ (100µM) and replacing calcium pantothenate (Vitamin B5) with sodium

pantothenate (Cayman Chemical, Cat# NC1435671). The cells were first inoculated in YE5s

media for 1 day, followed by another ~15 hours in EMM5s. After harvesting the exponentially

growing cells, we washed them three times with the calcium-free media before proceeding to

microscopy.

We employed a spinning disk confocal microscope for the time-lapse microscopy. The

microscope is an Olympus IX71 unit equipped with a CSU-X1 spinning disk unit (Yokogawa,

Japan). The motorized stage (ASI, USA) includes a Piezo Z Top plate for acquiring Z-series. The

images were captured on an EMCCD camera (IXON-897, Andor) controlled by iQ3.0 (Andor).

Solid-state lasers of 488 nm and 561 nm were used in the fluorescence microscopy, at power of

no more than 2.5mW. Unless specified, we used a 60x objective lens (Olympus, Plan

Apochromat, NA = 1.40). Typically, a Z-series of 8 slices at an interval of 1 μm was captured at

each time point during the time-lapse microscopy. The temperature was maintained at ~22 ± 2°C

in a designated room for microscopy. To further minimize environmental variations, we typically

imaged both control and experimental groups with randomized order in the same day.

For visualization of septum, we fixed the cells with 4% formaldehyde (EMS, USA)

before staining them with 1 μg/ml of calcofluor white (Sigma, Cat# 18909). For visualization of

nuclei, we fixed the cells with 70% cold ethanol before staining them with 1 μg/ml DAPI

(Roche, Cat# 1023627001). To image fixed cells, we used an Olympus IX81 microscope

equipped with a XM10 camera and a mercury lamp.

To apply osmotic and calcium stresses, we used a CellASIC ONIX2 system (EMD

Millipore), controlled by a desktop computer through the software ONIX (EMD Millipore). To

trap cells in the imaging chamber of a microfluidics plate (Y04C), we injected the exponentially

growing cell culture to the chamber at a pressure of 34.5 KPa for 2 mins, followed by a

continuous infusion of YE5s media at a pressure of 10KPa for another 30 mins. To apply hyper-

osmotic stresses, we injected YE5s plus sorbitol into the chamber at a pressure of 10 KPa.

Similarly, to apply hypo-osmotic stresses, we injected YE5s into the chamber at a pressure of 10

KPa after the chamber had been equilibrated with a continuous infusion of YE5s plus sorbitol for

30 mins.

We captured the kinetics of cytokinetic calcium spikes by taking time-lapse microscopy

of the GCaMP-expressing cells at a frequency of 0.33 frame/sec without the Z-series. A

complete spike is defined as one with a single peak rising from and decaying to the baseline

level. For the constricting spikes, we examined the dividing cells with either two separating

nuclei or a constricting ring.

Image processing and analysis

We quantified the intracellular fluorescence of a GCaMP expressing cell as the total fluorescence

intensity of the cell divided by its area. This value was used to represent the intracellular calcium

level throughout the study. Cells were segmented manually, and their fluorescence intensities

were quantified using average intensity projections of the Z-series. The intracellular fluorescence

of the cells trapped in a microfluidics chamber was quantified by subtracting their combined

fluorescence with the background fluorescence of the camera. Unless specified, we normalized

the intracellular fluorescence of a cell against the baseline value, which is typically defined as the

average of the five lowest values recorded during a time course. The microscopy images

acquired during osmotic or calcium shocks were analyzed by quantifying two regions of interest

in cytoplasm, each measuring 1 µm in diameter. The statistics tests were done using Excel. In

most cases, two-tailed Student T-test was used. Standard deviations (S.D.) are used throughout

the study to represent data deviations.

For quantification of the amplitudes and number of cytokinetic calcium spikes, we first

segmented the dividing cells manually using NIH ImageJ. These videos are first pre-processed

using a median filter to reduce noise. This non-linear filter helps smoothen the images while

preserving high spatiotemporal resolutions. Following the noise removal, the background

fluorescence signal from the cells is eliminated by subtracting the temporal average of each pixel

in every frame. This process enables us to retain most of the flickering events, while eliminating

the background fluorescence to a large extent. The average fluorescence intensity of a cell is then

calculated by applying a binary mask on the images. Binarization of the median filtered images

are performed by using Otsu's thresholding method and applied to the background subtracted

images. The average fluorescence intensity value at each time point is then recorded and used for

analysis. Prior to identifying the peaks, we perform a temporal averaging of the signal, using two

consecutive time points, to eliminate the small fluctuations within the data. Further thresholding

of the signal is performed by nullifying the values lower than the temporal average of the entire

signal. The temporal average is used as one of the criterions to distinguish the peaks from

background fluctuations. This thresholding step removes the variation in the remnant background

fluorescence signal inside the cells. The peaks or local maxima are detected by calculating the

derivative at each point and identifying the points corresponding to the inversion of the

derivatives from positive to negative. The location and average fluorescence intensity values of

each individual peaks are then recorded. The video processing and analysis was performed using

MATLAB (Mathworks, MA).

To measure the rate of the contractile ring closure, we determined the duration of the ring

closure through analyzing the fluorescence kymograph of contractile ring. The closure rate was

then calculated by dividing the initial circumference of the ring before the constriction by the

duration. We used Image J (NIH) and custom-made macros for all the image analyses. The

images were first corrected for X-Y drift using the Image J StackReg plug-in (Thevenaz et al.,

1998) if necessary. Most plots were made with Kaleidagraph (Synergy, PA) except for the 3D

plots which were made with OriginLab (OriginLab, MA). The figures were made with Canvas

(ACDSee Systems).

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QC conceptualized the study, QC and AP designed and carried out the experiments, QC, AP, OS,

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Figure 1

Figure 1. Expression and localization of GCaMP in fission yeast. (A) Bright-field (left) and fluorescence (right) micrographs of the cells expressing GCaMP. The intracellular fluorescence was constant for most cells, except for one outlier (white arrowhead). (B-C) Localization and expression of GCaMP-mCherry. (B) Fluorescence micrographs of the cells expressing GCaMP-mCherry. The insert shows the center slice of the Z-series of a representative cell (magnified, outlined in dashed lines). The intracellular fluorescence of mCherry remained low even in two cells with high GCaMP fluorescence (white arrow heads). GCaMP localized throughout the cytoplasm and nucleus, but it was excluded from the vacuoles. Similar results were found in three biological repeats. (C) A box plot showing the ratio of GCaMP:mCherry fluorescence. The expression is homogenous among the cells, resulting in a near uniform ratio with a few outliners with likely high calcium level. (D-F) Expression of GCaMP did not produce discernable artifacts. Bars graphs compare the GCaMP-expressing cells to the wild type. (D) The width of all cells (left, n > 50) and length (right, n > 50) of dividing cells. No significant differences were found (P > 0.1). (E-F) (D) Mitotic index (n > 1000) and (E) septation index (n > 700). No significant differences were found (P > 0.1). The data was from two biological repeats.

Figure 2

Figure 2. GCaMP responds to the calcium level increase. (A-B) The intracellular GCaMP fluorescence increased with added calcium in the media. (A) Fluorescence micrographs (spectrum colored) of the GCaMP-expressing cells in either YE5s (left) or YE5s supplemented with 30mM $CaCl_2$ (right). Bar represents the intensity scale. (B) Dot plot of fluorescence intensities of the cells (n > 190). Black line represents the average. Added calcium increased the average intracellular fluorescence by ~3 folds. (C-D) Calcium homeostasis of fission yeast cells. 3D-line plots show the time courses of intracellular GCaMP fluorescence in either YE5s (C) or YE5s plus 30mM $CaCl_2$ (D). The time-lapse movies were recorded at a frequency of 1 frame/min. Increased calcium in the media resulted in more frequent fluctuations of the intracellular calcium level. The data was pooled from two biological repeats.

Figure 3

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Figure 3. Osmotic shocks trigger calcium transients. (A-C) Hypo-osmotic shock triggered
calcium transients in fission yeast cells. The GCaMP expressing cells, trapped in a microfluidic
chamber, were first equilibrated in YE5s supplemented with 1.2M sorbitol for 30 mins before the
infusion of YE5s (time zero). **(A)** Time-lapse micrographs (spectrum-colored) of a representative
field in the chamber. The hypo-osmotic shock triggered a synchronized calcium spike in the cells.
(B) Time-lapse micrographs of a representative cell. In the media with 1.2M sorbitol, the cells
appeared shrunk. After the shock (time zero), the cell quickly expanded (asterisk), followed by
calcium transient (right, spectrum-colored). Number: time in minutes. **(C)** 3D-line plots of the time
courses of GCaMP fluorescence in twenty representative cells. The shock triggered synchronized
calcium transients in every cell. Representative results from two biological repeats are shown. **(D-**
F) Average time courses of the intracellular GCaMP fluorescence during either hypo-osmotic (D),
or hyper-osmotic (E), or extracellular calcium (F) shocks applied through microfluidics. These
stimuli triggered distinct change of intracellular calcium. **(D)** The cells were first equilibrated in
YE5s with various concentrations of sorbitol for 30 mins before the infusion of YE5s media (time
zero). The average amplitudes of calcium level increase were proportional to the strength of the
shocks. **(E)** The cells were first equilibrated in YE5s media for 30 mins before the infusion of
YE5s media supplemented with various concentrations of sorbitol (time zero). No significant
change of calcium was triggered by hyper-osmotic shock. **(F)** The cells were equilibrated in YE5s
media for 30 mins before the infusion of YE5s plus 30mM $CaCl_2$ (time zero). The addition of
extracellular calcium triggered a delayed increase of intracellular calcium. All data is pooled from
two biological repeats.
511

Figure 4

Figure 4. Correlation between intracellular calcium level and cell division. The intracellular

calcium level in asynchronized cells, based on the GCaMP fluorescence. Top: cartoon

representations of fission yeast cells at birth, growth, and division. **(A)** A scatter plot showing the

relationship between the intracellular calcium level and the cell length. The calcium level of

most cells was uniform (blue line). Only a few outlier cells (50/407) exhibited elevated (shaded

area, >110% of the average) calcium level. **(B)** A histogram showing the distribution of the cell

length among the outliers. The black line represents the best fit of a bimodal distribution. Two

peak fractions represent the cells of 8-9 μm, and those of 14-15 μm, roughly corresponding to

newly-born and dividing cells respectively. The data is pooled from four biological repeats.

Figure 5

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Figure 5. Calcium spikes accompany the cleavage furrow ingression. (A-B) The calcium
spikes accompanied the contractile ring constriction. **(A)** Left: Time-lapse micrographs of a
dividing cell expressing both Rlc1p-tdTomato (gray, top) and GCaMP (spectrum-colored,
bottom). Number: time in minutes after the start of ring constriction. Interval = 2 mins. Right:
Kymograph of the contractile ring (top) and the GCaMP fluorescence, with the dash line
marking the start of ring constriction. A calcium spike initiated at time zero and peaked at +2
min. Bar represents the relative scale of intracellular calcium level. **(B)** Kymographs of GCaMP
fluorescence in ten representative cells from -30 mins to +30 mins, relative to the initiation of
ring constriction. **(C-D)** Quantitative analysis of the intracellular calcium level during the ring
constriction. Data is pooled from three biological repeats. **(C)** 3-D line plots showing the time
courses of calcium level in representative cells (n = 64). **(D)** Box plots of intracellular calcium
level in the cells shown in C. The number on top indicates the number of observations made at
each time point. The start of ring constriction was accompanied by emergence of calcium spikes.
The average intracellular calcium level started to increase significantly at -4 mins and peaked at
+2 mins, relative to the start of ring constriction.
539

Figure 6

540
Figure 6. Calcium spikes accompany the cell separation. (A-C) Cytokinetic calcium spikes
during cell separation (white arrowhead). Asterisk: the peak of a spike. Bar: the relative scale of
intracellular calcium level. **(A)** Time-lapse micrographs of a cell expressing both GCaMP
(spectrum-colored) and Rlc1p-tdTomato (not shown). Three calcium spikes (asterisk) are
detected in this cell, including one following the start of ring constriction (time zero), the second
one during the ring constriction and the third one after the cell separation (arrowhead). Number:
time in minutes. Interval = 2 mins. **(B)** Time-lapse micrographs of two cells expressing GCaMP
(bottom). Interval = 1 min. Number: time in minutes after the cell separation (time zero,
arrowhead). The spikes appeared either in just one daughter cell (top) or oscillate between two
daughter cells (bottom). Arrowhead: completion of the cell separation. **(C)** Kymograph of
separating spikes in ten representative cells. Dashed box: the separating spikes oscillating
between two daughter cells. **(D-E)** Quantification of the intracellular calcium level during the
separation. The time-lapse movie was captured at a frequency of 1 frame/min. The data is pooled
from four biological repeats. **(D)** 3-D line plots showing the time courses of intracellular calcium
level in the dividing cells (n = 44). A highly synchronized calcium spike followed the cell
separation (time zero) in most cells. **(E)** Box plots of the calcium level of the cells shown in D.
Number of observations at each time point is indicated on the top. The average calcium level
started to increase at 0 min and peaked at +2 min.

Figure 7

Figure 7. Spatiotemporal regulation of cytokinetic calcium spikes. The spikes were captured

at a frequency of one frame every three seconds during the time-lapse microscopy.

Representative constricting (A-C) or separating spikes (D-F) are shown. (A, D) Time-lapse

micrographs (spectrum-colored) of two GCaMP-expressing cells during cytokinesis. Arrowhead:

hots spot of calcium increase in the cytoplasm. Number: time in seconds after the initial rise of

calcium (Time zero). Asterisk: the peak of calcium level. (B, C) Fluorescence kymograph of the

spikes. Calcium increased most significantly at the cell tips, the peri-nuclear area and the cell

division plane during spikes. (C, F) Time course of cytokinetic spikes including the one shown

on the left (thick black line). The cytokinetic calcium spikes are highly heterogeneous with

various amplitudes and lifespans. Data is pooled from three biological repeats.

Figure 8

Figure 8. Depletion of extracellular calcium inhibits the cytokinetic calcium spikes. Either

supplement of EGTA or removal of calcium from the media inhibited the calcium spikes. **(A-B)**

Violin plots showing the intracellular calcium level in the cells. Depletion of calcium from the

media reduced the average intracellular calcium level only slightly (13% by EGTA and 3% by

the calcium-free EMM media) but it greatly reduced the number of outlier cells with high

calcium level. **(C-D)** Bar graphs showing the average amplitude (E) and number (F) of calcium

spikes during the ring constriction (n = 20). The amplitude and number of spikes are quantified

through thresholding using custom-written algorithm as described in the Materials and Methods.

Depletion of extracellular calcium reduced the average amplitude of calcium spikes significantly

(P < 0.001), but not the number of peaks. The data is pooled from two biological repeats.

Figure 9. Temporary depletion of extracellular calcium slows down the contractile ring constriction and leads to daughter cell lysis. (A-F) Temporary depletion of extracellular calcium reduced the ring constriction rate, but it had little effect in the ring assembly and maturation. **(A-B)** Temporary depletion of extracellular calcium slowed down the ring constriction. Time-lapse micrographs of dividing cells expressed both Sad1-GFP (green) and Rlc1p-tdTomato (red). Number: time in minutes after the separation of SPBs (arrowhead). Dash

line marks the outline of cells. **(A)** Cells in either regular EMM5s (top) or the calcium-free

EMM5s media (bottom). **(B)** Cells in either YE5s (top) or YE5s plus 2mM EGTA media

(bottom). **(C, D)** Duration of the ring assembly plus maturation. **(E, F)** The ring constriction

rates. **(G-H)** Temporary depletion of extracellular calcium increased the frequency of cell lysis

among the daughter cells. Time-lapse micrographs of separating cells. Number: time in minutes

after the cell separation. **(G)** The wild-type (top) or *pmr1Δ* (bottom) cells in the calcium-free

EMM5s media (bottom). Although the wild-type cells (n > 100) did not lysis during the

separation, 54% of the mutant cells lysed (asterisk) (n = 57). **(H)** The wild-type cells in either

YE5s (top) or YE5s plus 2mM EGTA (bottom). About 35% of daughter cells lysed after the

separation in the presence of EGTA (n > 100). Data is pooled from two biological repeats.

Supplemental Materials

Supplemental Figure legends:

Figure S1

604 **Figure S1: Expression of GCaMP does not affect fission yeast growth. Related to Fig. 1.**

605 Ten-fold dilution series of both GCaMP-expressing and the wild-type cells. The cells were

606 inoculated at various temperatures on either YE5s (top), or YE5s plus 25mM $CaCl_2$, or YE5s

607 plus 1.2M sorbitol, or YE5s plus 2mM EGTA plates. GCaMP did not affect the yeast growth

608 under any of the conditions. Similar results were obtained in two biological repeats.

Figure S2: Hyper-osmotic shocks do not trigger calcium spikes. Related to Fig. 3. The cells

were equilibrated with YE5s media for 30 mins in the microfluidic chamber before the infusion

of YE5s plus 1.2M sorbitol (time zero). **(A)** Time-lapse micrographs (spectrum-colored) of the

GCaMP expressing cells in a microfluidics chamber. Number: time in minutes after the shock.

Unlike hypo-osmotic shock, hyper-osmotic shock did not trigger synchronized intracellular

calcium spikes. Similar result was obtained in two biological repeats. **(B)** 3D line plots showing

the time courses of the intracellular GCaMP fluorescence in twenty representative cells during

the hyper-osmotic shock.

Figure S3

Figure S3: The calcium spikes in dividing cells. Related to Fig. 4, 5 and 6. (A-C) The
intracellular calcium level increased during late mitosis. **(A)** Time-lapse fluorescence
micrographs of a cell expressing both Sad1-mCherry (red) and GCaMP (green). Number: time in
minutes after the SPB separation (arrowheads). Interval = 4 mins. The intracellular calcium level
peaked at +32 mins (asterisk), at the start of telophase when two SPBs reached a maximum
distance from each other. **(B)** 3D-line plots of the time courses of calcium level in mitotic cells (n
= 36). The calcium level increased significantly starting ~ 30 mins after the separation of SPBs.
(C) Box plots of the calcium level of those cells shown in B. The average calcium level peaked
at +34 min, after the separation of the SPBs. The data was pooled from two biological repeats.
(D-E) The cytokinetic calcium spikes in EMM5s media. 3D-line plots of the calcium level
during the ring constriction (D) and cell separation (E). Similar constricting and separating
spikes are found in these cells, as those inoculated in YE5s.

Figure S4

Figure S4: Depletion of extracellular calcium inhibits the cytokinetic calcium spikes.

Related to Fig. 8. (A) Ten-fold dilution series of the wild-type yeast in YE5s plus 0-2 mM

EGTA. Low concentration of EGTA didn't inhibit cell growth. **(B-E)** 3D-line plots showing the

time courses of the calcium level of the representative cells in either YE5s plus 2mM EGTA (B-

C) or the calcium-free EMM media (D-E) during either the ring closure (B and D) or the cell

separation (C and E). Depletion of extracellular calcium inhibited the cytokinetic calcium spikes.

The data is pooled from two biological repeats.

Figure S5

Figure S5: **Depletion of extracellular calcium slowed down the contractile ring constriction.**

Related to Fig. 9. (A-B) Depletion of calcium by 5 mM EGTA reduces the integrity of the

contractile ring. The cells expressed both Sad1p-GFP (green) and Rlc1p-tdTomato (red). **(A)**

Time-lapse micrographs of a cell in YE5s supplemented with EGTA. The contractile ring failed

to close completely before its disintegration (asterisk). Number: time in mins after the separation

of SPBs (arrowhead). **(B)** Kymographs of two contractile rings that failed to constrict completely

before their disintegration (asterisk). **(C-D)** Box plots showing the duration of cell separation

when extracellular calcium was depleted. The separation was not significantly delayed.

Supplemental Tables:

Table 1: The list of yeast strains

Strains	Genotype	Source
FY527	*h- leu1-32 ura4-D18 his3-D1 ade6-M216*	Lab stock
FY528	*h+ leu1-32 ura4-D18 his3-D1 ade6-M210*	Lab stock
QC-Y941	*h- leu1::KanMX6-Padh1-GCaMP6s*	This study
QC-Y989	*h- sad1-mCherry-NatMX6 leu1::Padh1-GCaMP6s-mCherry*	This study
QC-Y1004	*h+ rlc1-tdTomato-NatMX6 leu1::Padh1-GCaMP6s-mCherry*	This study
QC-Y1068	*h- leu1::Padh1-GCaMP6s-mCherry-ura4+*	This study
QC-Y274	*h- rlcl1-tdTomato-NatMX6 sad1-mGFP-KanMX6*	Lab stock
QC-Y1149	*h? pmr1::kanMX6 rlc1-tdTomato-natMX6 sad1-mEGFP-KanMX6*	This study

Table 2: The list of DNA primers

Prim ers	Sequence
P413	*TTTGGGAGGTGAGGCTTCTACCGCTGATATTACTAAGGCTGTTGTTGAAGA ACTTGAAAAAATTTTGTAAGAATTCGAGCTCGTTTAAAC*
P414	*ATGCGACACAATTAATCAATAAAAATTAGGTCAAATCAAATCATATGAATT TATTCTACATTAAACCCTAGCGGCCGCCAGCTGAAGCTT*
P415	*CCTTCTCCTTCGTTATGGTTTG*
P589	*TTAATTAAC-ATGGGTTCTCATCATCATCATC*
P590	*GGCGCGCC-TCACTTCGCTGTCATCATTTGTAC*
P668	*CAGGGACACGGAAGAAGAAATTA*
P416	*TCATAAGTTGAACGGATGTCG*
P647	*CAGGGAAGCAGACATCGATGGGGATGGTCAGGTAAACTACGAAGAGTTTGTACAAATG ATGACAGCGAAGGTGAGCAAGGGCGAGGAGGA*
P648	*TTTATAACTTATTTAATAATAAAAATCATAAATCATAAGAAATTCGCTTATTTAGAAGTGGCGCGCC CTAGAATTCGAGCTCGTTTAAAC*

Supplemental movies:

Movie S1: The calcium homeostasis of fission yeast reported by GCaMP-based imaging (related to Fig. 2). Time-lapse movie (60 mins) of GCaMP expressing cells in YE5s. Number: time in minutes.

Movie S2: The calcium spikes stimulated by hypo-osmotic shock (related to Fig. 3). Time-lapse movie of the GCaMP expressing cells in a microfluidic chamber. At time zero, the infusing media was switched from YE5s plus 1.2M sorbitol to just YE5s. This hypo-osmotic shock (time zero) triggered synchronized calcium transients. Number: time in minutes.

Movie S3: The constricting calcium spike during cytokinesis (related to Fig. 4). Time-lapse

664 movie of a dividing cell expressing both GCaMP (green) and Rlc1p-tdTomato (red). Number:

665 time in minutes. The calcium spike initiated as the ring constriction started (time zero) and

666 peaked at +2 min. Number: time in minutes.

667 **Movie S4: The separating calcium spike during cytokinesis (related to Fig. 5).** Time-lapse

668 movie of a GCaMP expressing cell during the cell separation. The calcium spike followed the

669 separation (time zero), oscillating between the two daughter cells. Number: time in minutes.

670 **Movie S5: Cytokinesis in the calcium depleted media (related to Fig. 9).** Time-lapse movie of

671 two cells in either YE5s (left) or YE5s plus 2mM EGTA (right). The cells expressed both

672 GCaMP (green) and Rlc1p-tdTomato (red). EGTA inhibited the cytokinetic calcium spikes. The

673 contractile ring constricted more slowly in the presence of EGTA. Number represents time in

674 minutes after the start of the ring constriction (time zero).

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