

Minireview

## Thapsigargin: potent inhibitor of $\text{Ca}^{2+}$ transport ATP-ases of endoplasmic and sarcoplasmic reticulum\*

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Received 17 May, 1993

### GENERAL CHARACTERISTICS

Thapsigargin and thapsigarginin are two major active components found in root extracts of the umbelliferous (*Apiaceae*) plant *Thapsia garganica* L. This root extract has been known from ancient times for its skin irritation effect and was used for treatment of muscles and joint inflammations. Rasmussen *et al.* [1] isolated these two compounds and showed that thapsigargin and thapsigarginin release histamine in rat mast cells. Thapsigargin, the most abundant of the two compounds, is a sesquiterpene lactone whose structure has been studied in detail by NMR, circular dichroism spectroscopy and crystallographic methods [2 - 5] (Fig. 1). Thapsigargin is known to promote tumorigenesis in mouse skin [6] independently of protein kinase C activation or protein phosphatases inhibition [7, 8]. Most recently, Wong *et al.* [9] have shown that thapsigargin-induced tumorigenesis can be caused by rapid inhibition of mRNA translation, protein processing and gene expression. Thapsigargin is a neurotoxin in perinatal rodent brain causing irreversible brain injury [10], and totally blocks the induction of long-term potentiation in rat hippocampal slices [11]. It has been reported that thapsi-

gargin induces acute responses in a large variety of cell types.

Thapsigargin-induced cellular activation appears in all cases to be initiated by a single common event: a rapid and pronounced increase in the concentration of cytosolic free  $\text{Ca}^{2+}$  that occurs *via* a direct discharge of intracellular  $\text{InsP}_3$ -sensitive  $\text{Ca}^{2+}$  stores without hydrolysis of inositolphospholipids [8, 12 - 14]. Since this discovery, the interest in thapsigargin studies has increased, especially when it was established that thapsigargin is a high affinity inhibitor of all the endoplasmic reticulum  $\text{Ca}^{2+}$  transport ATPases tested [15 - 18]. Thus, for its abilities of identification and selective interference with intracellular  $\text{Ca}^{2+}$  pools, thapsigargin became a very convenient experimental tool.

### ACTION OF THAPSIGARGIN ON $\text{Ca}^{2+}$ TRANSPORT ATPases

In 1987 Thastrup *et al.* [12, 13] found that thapsigargin produced elevation of intracellular calcium concentration by emptying  $\text{InsP}_3$ -sensitive stores without generation of phosphoinositides. Three years later Thastrup and coworkers [19] revealed that this effect was due to the inhibition of calcium uptake into the

\*This work was supported by the State Committee for Scientific Research by a Statutable grant to the Nencki Institute and grant No. 406109101 to the authors

<sup>1</sup>Abbreviations: SERCA, sarcoplasmic and endoplasmic reticulum calcium ATPases;  $\text{InsP}_3$ , inositol 1,4,5-trisphosphate

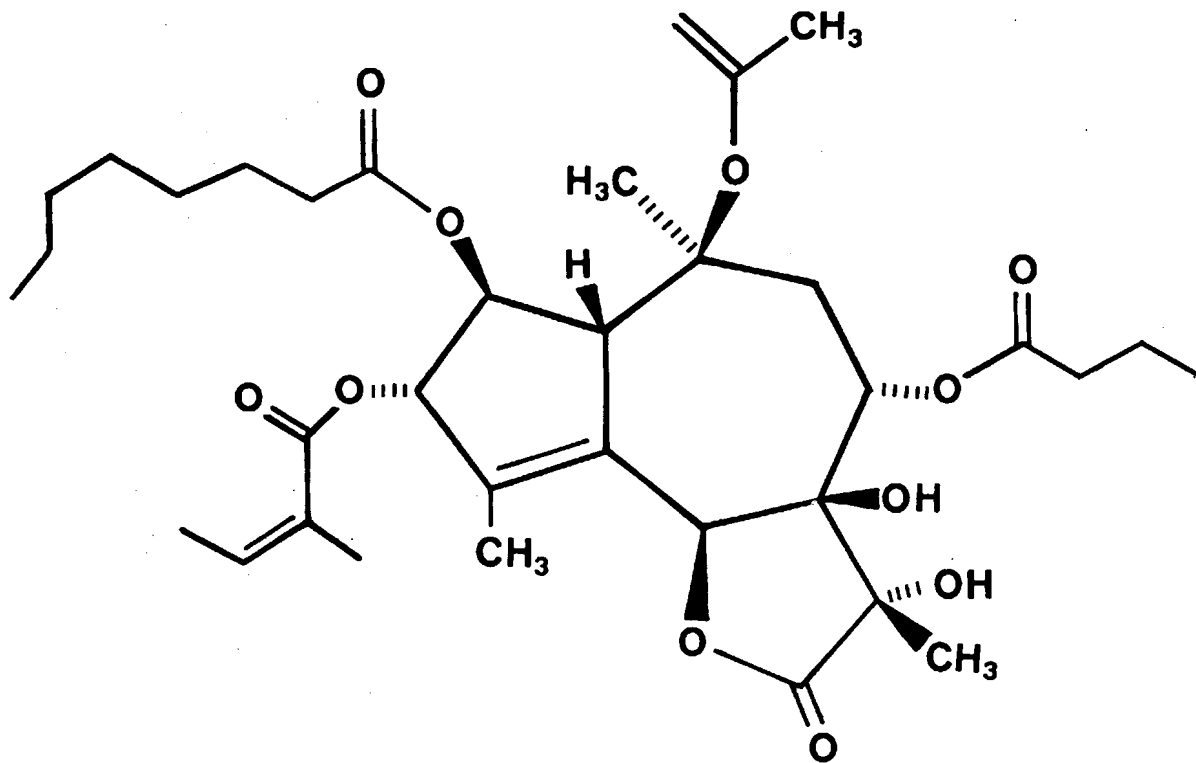


Fig. 1. Structure of thapsigargin.

endoplasmic reticulum by blocking its ATP-dependent calcium pump. In 1991 Lytton *et al.* [17] showed that thapsigargin acted on all sarcoplasmic and endoplasmic reticulum  $\text{Ca}^{2+}$  transport ATPases. The mechanism of this inhibition was explained by Sagara & Inesi [15] in the same year.

#### SERCA ATPases

$\text{Ca}^{2+}$  is loaded into the endoplasmic reticulum by means of intracellular adenosine triphosphate (ATP)-dependent  $\text{Ca}^{2+}$  pumps of the highly homologous SERCA<sup>1</sup> (sarcoplasmic or endoplasmic reticulum  $\text{Ca}^{2+}$ -ATPase) family. Recently three mammalian SERCA genes (SERCA-1, SERCA-2 and SERCA-3) have been identified. It has been found that transcripts of SERCA-1 and SERCA-2 exist in two alternatively spliced forms [20], which produce at least five different protein products. The calcium dependent ATPase whose activity is essential for the  $\text{Ca}^{2+}$  pump's function is a 110 kDa polypeptide chain placed asymmetrically across the endoplasmic reticulum membrane with a large extramembranous region on the cytosolic side [21, 22]. The enzyme binds two  $\text{Ca}^{2+}$  ions and ATP from cytosol and releases the bound  $\text{Ca}^{2+}$  into the lumen of the endoplas-

mic reticulum by utilization of ATP. Calcium binding to the high affinity sites of the ATPase promotes ATP binding and hydrolysis, the enzyme is phosphorylated and a series of conformational changes causes  $\text{Ca}^{2+}$  translocation to the luminal side of the endoplasmic reticulum [23, 24]. The mechanism of action of the  $\text{Ca}^{2+}$ -ATPase may be described according to a model with two major states: E<sub>1</sub> and E<sub>2</sub>, shown in Fig. 2.

As shown in the schematic representation of the endoplasmic reticulum  $\text{Ca}^{2+}$ -ATPase (Fig. 3) the phosphorylation domain (Asp 351) resides in the extramembranous region of the enzyme and the  $\text{Ca}^{2+}$  binding domain resides within the transmembrane region and consists of four helices [22]. It is likely that four helices form a channel for  $\text{Ca}^{2+}$  which is controlled by the phosphorylation site [24].

#### Mechanism of action

The inhibitory effect of thapsigargin on  $\text{Ca}^{2+}$ -ATPase can be observed at very low, subnanomolar concentrations [15] and at various isoforms of SERCA pumps [16,17]. The plasma membrane ATPases and  $\text{Na}^+/\text{K}^+$  ATPases are not inhibited [17]. The dead-end thapsigargin\*ATPase complex is formed when the

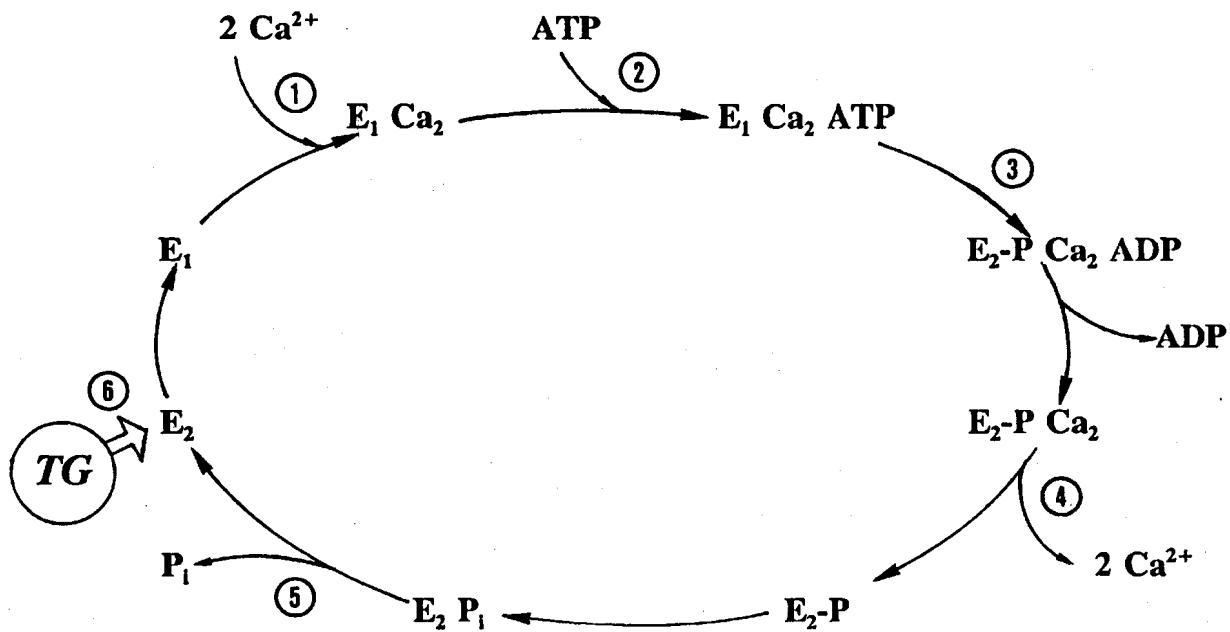


Fig. 2. Calcium pumping cycle of sarcoplasmic and endoplasmic reticulum ATPase.

The binding of the two  $\text{Ca}^{2+}$  from the cytosol (1) to the high affinity sites of  $\text{Ca}^{2+}$ -ATPase ( $\text{E}_1$  state) is strictly required for starting its pumping action. When these sites are occupied ATP binds to the enzyme (2) which is then phosphorylated by a transfer of ATP terminal phosphate (3). The series of conformational changes in the enzyme leads to a translocation of bound  $\text{Ca}^{2+}$  and a 3-order magnitude reduction in affinity for these ions. Finally, calcium dissociates to the luminal spaces of the endoplasmic reticulum (4) and the phosphorylated intermediate is hydrolysed (5). All presented reactions are reversible. Thus, the defined sequence of chemical and vectorial reactions with utilization of free energy from ATP produces ion transition and reduction of the affinity for  $\text{Ca}^{2+}$ . The postulated inhibition of thapsigargin (TG) starts when the enzyme is in the  $\text{E}_2$ ,  $\text{Ca}^{2+}$  free form (6).  $\text{E}_1$  and  $\text{E}_2$  correspond to different conformational forms of  $\text{Ca}^{2+}$ -ATPase. In fact there may be many more conformational changes in the cycle for calcium pumping by the enzyme [23, 80].

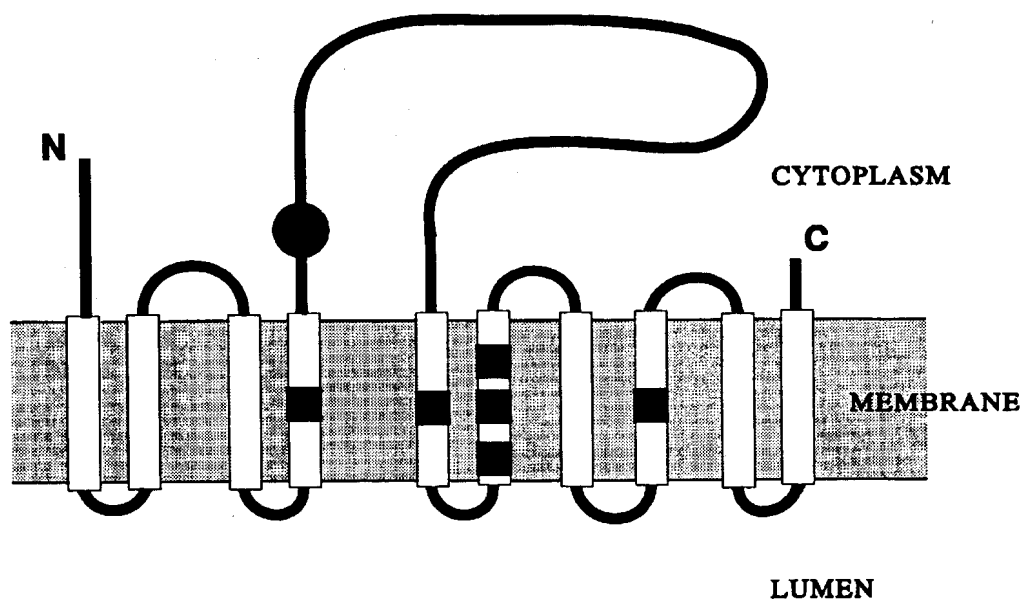


Fig. 3. Schematic representation of SERCA pump.

SERCA pump consists of ten transmembranous domains. Four of them (black squares) are involved in  $\text{Ca}^{2+}$  binding. The largest-cytoplasmic fragment is responsible for ATP binding; the black circle represents the phosphorylation site (Asp 351).

enzyme is in a state without bound  $\text{Ca}^{2+}$ ; it occurs when bound  $\text{Ca}^{2+}$  dissociates as a consequence of ATP utilization during the action of the pump. Thapsigargin produces a global influence on the enzyme resulting in a strong inhibition of  $\text{Ca}^{2+}$  binding and ATPase activity (including ATP binding and partial reactions in the absence of  $\text{Ca}^{2+}$ ) [18, 21, 25]. Thapsigargin is less effective in blocking  $\text{Ca}^{2+}$  binding to the  $E_1$  form of the enzyme, and higher concentrations of free  $\text{Ca}^{2+}$  have a protective effect on  $\text{Ca}^{2+}$  binding to the ATPase [18]. Under these conditions, the phosphorylated enzyme is formed even in the presence of an excess of thapsigargin until the enzyme reaches the  $E_2$  state. In fact,  $\text{Ca}^{2+}$ -ATPase is susceptible to thapsigargin in its  $E_2$ ,  $\text{Ca}^{2+}$ -free form [18, 21]. Thapsigargin acts with the enzyme stoichiometrically of 1 mol of inhibitor per mol of ATPase. Its interactions with  $E_2$  produce a global inhibition of the  $\text{Ca}^{2+}$  transport ATPase activities which is manifested by total blocking of high affinity  $\text{Ca}^{2+}$  binding, loss of  $\text{Ca}^{2+}$  transport, ATP binding and phosphorylated enzyme formation [18, 21]. The reversed action of the ATPase cycle is also inhibited; in the presence of thapsigargin there is a very low (in favorable conditions) phosphorylated enzyme formation from  $\text{P}_i$  [21], and ADP-dependent  $\text{Ca}^{2+}$  efflux from loaded sarcoplasmic reticulum vesicles (with ATP synthesis) is prevented [15, 26].

Thus, thapsigargin is a specific and irreversible blocker of  $\text{Ca}^{2+}$ -ATPases in both muscle (skeletal [15, 18, 21], smooth [27, 28], cardiac [29 - 31]) and nonmuscle cells [17, 19, 32, 33]. The mechanism of the action of this drug on the  $\text{Ca}^{2+}$  pump protein of sarcoplasmic reticulum vesicles purified from rabbit skeletal muscle was, as mentioned above, the subject of studies of Sagara & Inesi [15]. Similar results of  $\text{Ca}^{2+}$ -ATPase blocking by thapsigargin in cardiac myocytes were achieved by Wrzosek *et al.* [29]. The effect of thapsigargin on sarcoplasmic reticulum  $\text{Ca}^{2+}$  content and contractions in single cardiac myocytes of guinea-pig heart was also reported by Lewartowski & Wolska [30]. They showed that thapsigargin completely blocked  $\text{Ca}^{2+}$  uptake by the sarcoplasmic reticulum of these cells. Despite of the sarcoplasmic reticulum depletion of  $\text{Ca}^{2+}$ , the amplitude of electrically stimulated contraction was decreased only by 30%, what suggests that in this type of cells not only sarcoplasmic reticu-

lum, but also sarcolemmal  $\text{Ca}^{2+}$  may activate the contractile system. In vascular smooth muscle [28], thapsigargin inhibits sarcoplasmic reticulum  $\text{Ca}^{2+}$ -ATPase and stimulates calcium entry. Thapsigargin has also been shown to inhibit the endoplasmic reticulum  $\text{Ca}^{2+}$ -ATPase, to mobilize intracellular  $\text{Ca}^{2+}$ , and to activate  $\text{Ca}^{2+}$  entry in nonmuscle cells [14, 19]. It is generally accepted that calcium concentration in the endoplasmic reticulum might depend on two opposed events: active transport of  $\text{Ca}^{2+}$  by the ATPase and its passive leak from the endoplasmic reticulum [19]. Thapsigargin by blocking  $\text{Ca}^{2+}$  uptake may change the equilibrium towards leak from the endoplasmic and sarcoplasmic reticulum in nonmuscle and muscle cells, respectively. The nature of the  $\text{Ca}^{2+}$  leak remains unknown, but it appears to be widespread in eukaryotic cells. Depletion of internal stores is somehow coupled to the entry of extracellular calcium. All of these events lead to a distinct and sustained increase in intracellular  $\text{Ca}^{2+}$  levels in nonmuscle and smooth muscle cells, producing a variety of cellular responses.

#### THAPSIGARGIN AS A NEW MOLECULAR PROBE

Thapsigargin proved to be a valuable tool for the evaluation of many physiological, calcium-related cellular events, such as agonist and hormone activation, immunological response, muscle contraction-relaxation cycle, fertilization, secretion, phospholipid synthesis, intercellular communication and many others. Table 1 presents the areas of research where thapsigargin was engaged to elucidate mechanisms of various cellular events.

Thanks to the discovery and explanation of thapsigargin action on  $\text{Ca}^{2+}$ -ATPases it became clear that the sarcoplasmic and endoplasmic reticulum are the most important  $\text{Ca}^{2+}$  sequestering organelle in the cell. It was also possible to distinguish different calcium pools within this heterogeneous structure. Extracellular signals can trigger elevation of intracellular calcium concentration by inducing its influx from the extracellular space and/or by releasing  $\text{Ca}^{2+}$  from intracellular stores. It has been found that thapsigargin can mimic this action,

Table 1

*Application of thapsigargin as a tool for elucidation of mechanisms of various cellular processes*

Areas of research	References*
Calcium homeostasis and dynamics	
* identification and characterization of intracellular Ca <sup>2+</sup> pools	[34 - 44]
* cytoplasmic Ca <sup>2+</sup> signals after agonist or hormone stimulation	[45 - 52]
* calcium oscillations	[53 - 56]
Intercellular communication	[57]
Secretion	[58 - 62]
Calcium pumps	[21, 29, 63, 64]
Muscle contraction-relaxation cycle	[29 - 31, 65 - 67]
Phosphatidylserine synthesis	[68, 69]
Tumorigenesis	[9, 70 - 73]

\* Literature data published in 1992 and 1993

depleting intracellular InsP<sub>3</sub>-sensitive and thapsigargin-sensitive calcium stores [27, 32].

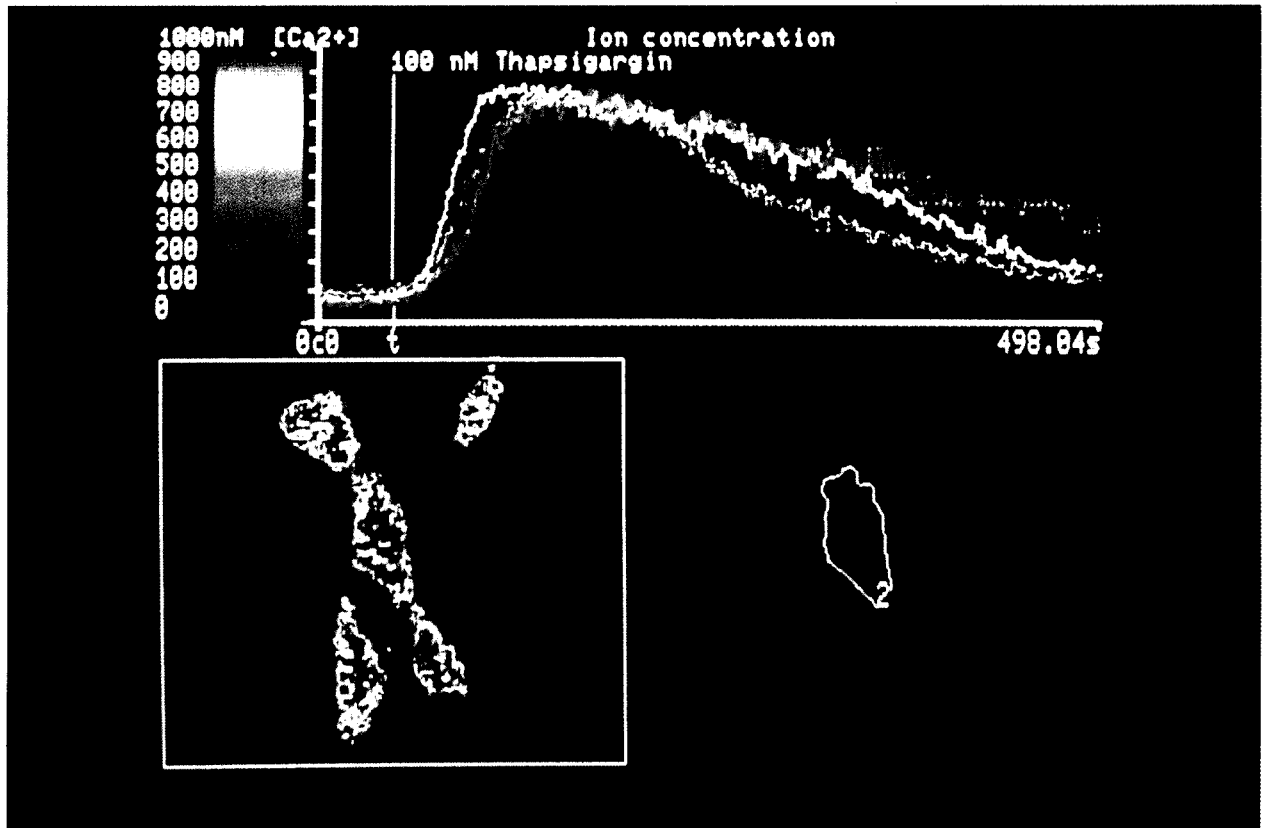
In a variety of nonmuscle cells a relationship between release of Ca<sup>2+</sup> from intracellular stores and increased entry of Ca<sup>2+</sup> across the plasma membrane was observed [14, 33, 74]; the mechanism which triggers this calcium influx remains unknown but two possibilities have been suggested. Some authors believed that the increase of plasma membrane permeability was connected with the elevated Ca<sup>2+</sup> concentration in the cytosol in response to emptying the endoplasmic stores [75, 76]. Others have proposed that the entry pathway of Ca<sup>2+</sup> across the plasma membrane is responsive to the Ca<sup>2+</sup> content of the stores [14, 77]. According to this hypothesis, concentration of the cytosolic Ca<sup>2+</sup> is irrelevant to this process. In 1986, Putney [77] developed a "capacitative" model of Ca<sup>2+</sup> entry and postulated that plasmalemmal permeability is controlled by the degree of filling of an intracellular Ca<sup>2+</sup> storage compartment. Support for this hypothesis has come from experiments in which thapsigargin was used. Inhibition of Ca<sup>2+</sup>-ATPase by thapsigargin produces the leak of calcium from the endoplasmic reticulum. It has been found that the depletion of these intracellular Ca<sup>2+</sup> pools, independent of receptor activation, increases Ca<sup>2+</sup> permeability of the plasma membrane and is responsible for the influx of Ca<sup>2+</sup> into the cell.

The introduction by Gryniewicz *et al.* [78] of fluorescent dyes, whose excitation and

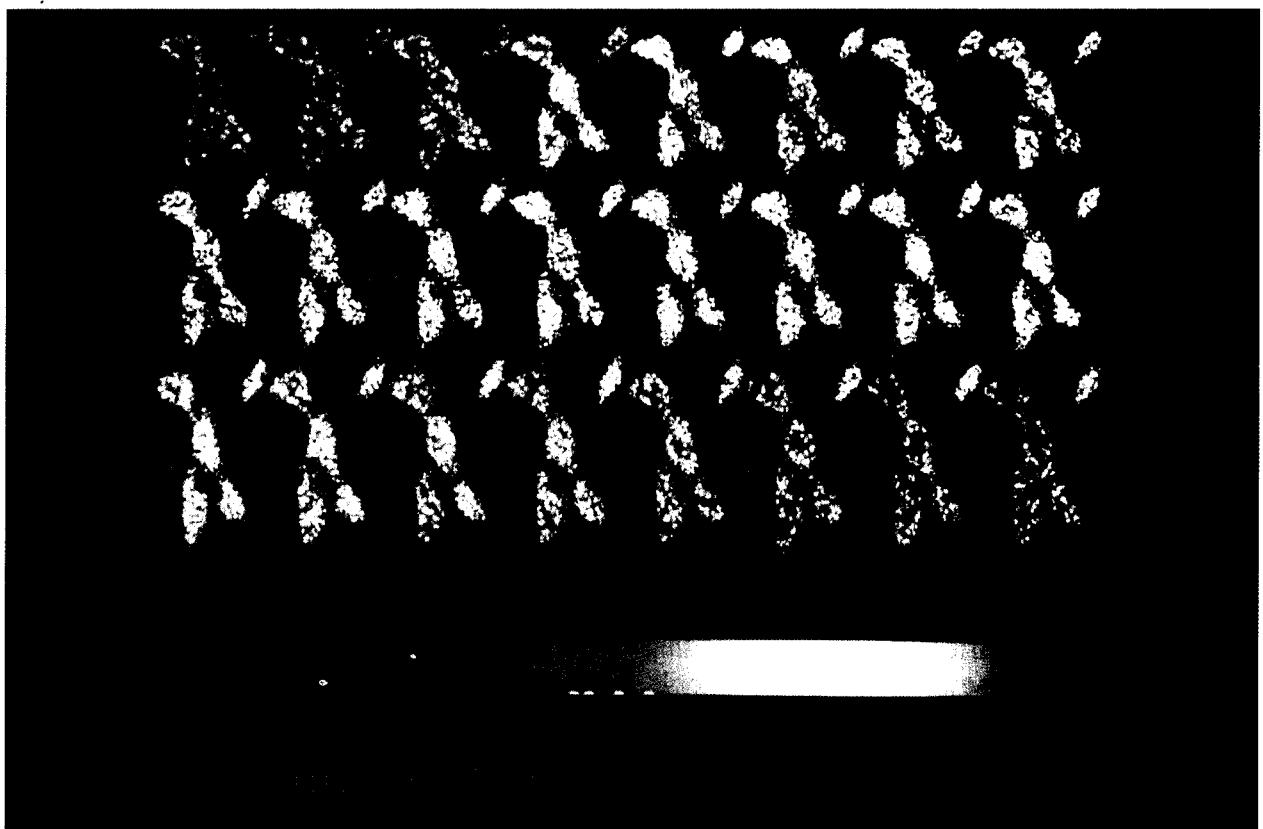
emission properties are altered when they bind Ca<sup>2+</sup>, has greatly enhanced the ability to monitor intracellular Ca<sup>2+</sup> dynamics. The use of fluorescent probes and digital image analysis systems allows for monitoring of intracellular calcium levels in a single, living cell.

Figure 4 illustrates the effect of thapsigargin on intracellular calcium levels in intact glioma C6 cells [68]. In our study the measurements of changes in Ca<sup>2+</sup> level in cytosol after the thapsigargin treatment were performed simultaneously with the determination of the effect of this drug on phosphatidylserine synthesis. We found [68] that thapsigargin, in addition to increasing Ca<sup>2+</sup> concentration in the cytoplasm, strongly diminished phosphatidylserine synthesis. Since the synthesis of this phospholipid depends on Ca<sup>2+</sup> and takes place in the endoplasmic reticulum [79], the inhibitory effect of thapsigargin on this process allowed us to conclude that phosphatidylserine synthesis occurs on the luminal side of this structure and has to be regulated by Ca<sup>2+</sup>-ATPases. Similar results on phosphatidylserine synthesis in Jurkat T lymphocytes were recently presented by Pelassy *et al.* [69].

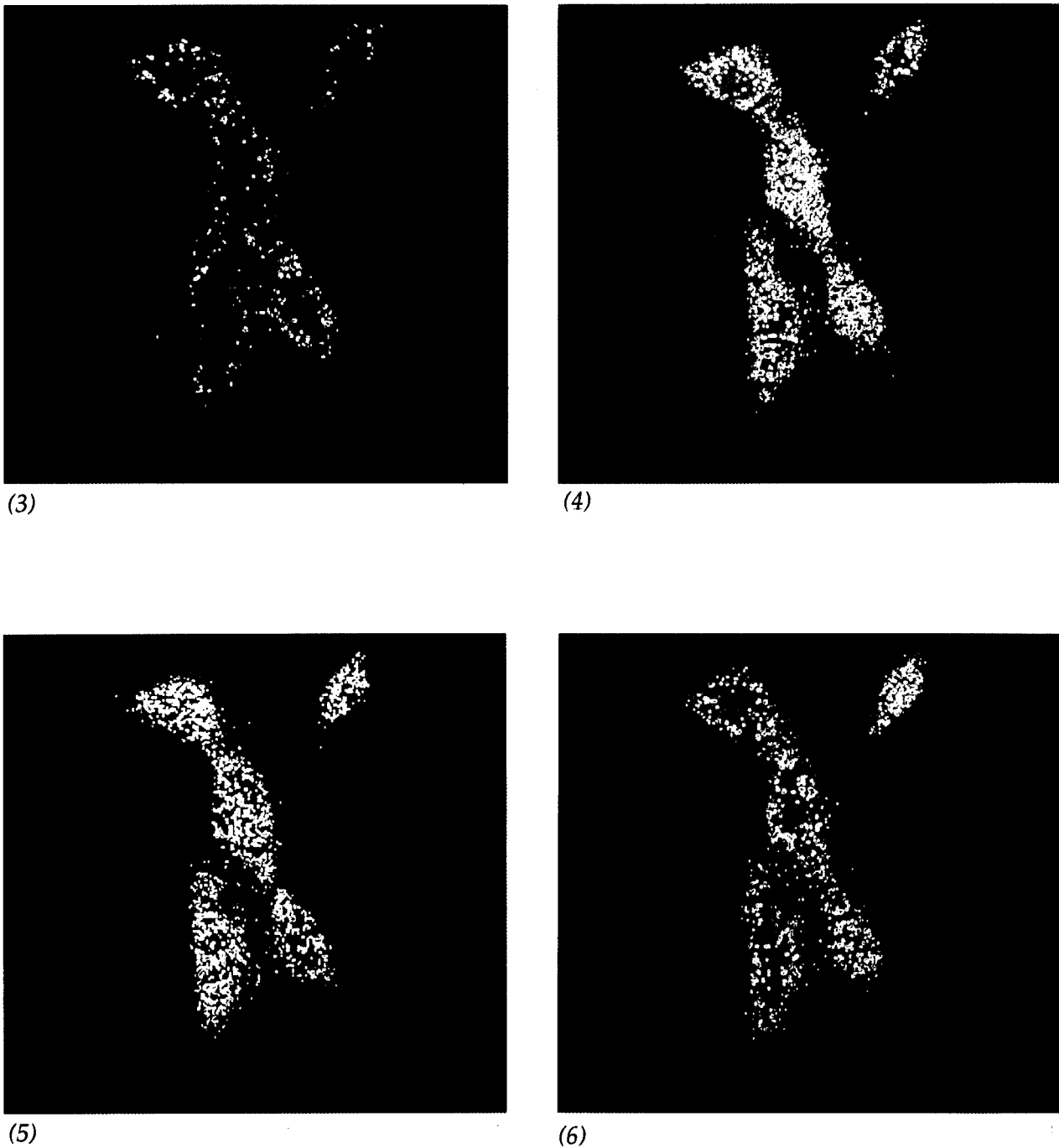
The broad spectrum of research in which thapsigargin is used as a molecular probe to elucidate various intracellular processes (Table 1) is connected with the universal role of Ca<sup>2+</sup> as a second messenger. The fact that thapsigargin is a potent and specific inhibitor of Ca<sup>2+</sup> transport ATPases in both the sarcoplasmic and endoplasmic reticulum revealed the similarity



(1)



(2)



*Fig. 4. Pseudo-color images illustrating the effect of thapsigargin on intracellular calcium levels in intact glioma C6 cells.*

Images show the increase in intracellular calcium concentration after 100 nM thapsigargin treatment. The pseudo-color scale covering the 0 - 1000 nM range is shown. Cells were cultured on glass coverslips and loaded with the intracellular  $\text{Ca}^{2+}$  indicator Fura-2 during 30 min incubation with 1  $\mu\text{M}$  of its acetoxymethyl ester. (1) Shows the action of thapsigargin on a single glioma C6 cell during 500 s. As it is seen, treatment of Fura-2 labelled cells by thapsigargin caused an increase in cytosolic  $\text{Ca}^{2+}$  and after a lag time period of 20 s a sustained  $[\text{Ca}^{2+}]_i$  rise. This changing of the cytosolic calcium levels are represented by curves (1) and by the changing of pseudocolors cf. (2). (3) - (6) Represent Fura-2 loaded, thapsigargin treated cells after subsequent periods of time. All of these pictures originate from an experiment carried out on a Joyce-Loebl Ltd. computerizing image-processing system Magiscan for calcium analysis using Tardis V6.2.

of these organelles which serve as major intracellular  $\text{Ca}^{2+}$  stores in eukaryotic cells. In conclusion, the elucidation of the mechanism of thapsigargin action allows us to gain better insight into the  $\text{Ca}^{2+}$ -related molecular events in the cell.

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