



# IP<sub>3</sub> receptors and their intimate liaisons

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Inositol 1,4,5-trisphosphate receptors (IP<sub>3</sub>Rs) are intracellular Ca<sup>2+</sup> channels. They allow cell-surface receptors that stimulate IP<sub>3</sub> formation to evoke rapid Ca<sup>2+</sup> release from the endoplasmic reticulum (ER). IP<sub>3</sub>Rs initiate local and global cytosolic Ca<sup>2+</sup> signals, they deliver Ca<sup>2+</sup> selectively to other organelles including mitochondria and lysosomes, and, by depleting the ER of Ca<sup>2+</sup>, they control store-operated Ca<sup>2+</sup> entry (SOCE). We consider two areas where recent work highlights the importance of liaisons between IP<sub>3</sub>Rs and other intracellular membranes. Interactions between IP<sub>3</sub>Rs and lysosomes illustrate striking parallels with the relationships between IP<sub>3</sub>Rs and mitochondria. In each case, the ER concentrates Ca<sup>2+</sup> from the cytosol and then delivers it through IP<sub>3</sub>Rs to a low-affinity Ca<sup>2+</sup>-uptake system in a juxtaposed organelle. Evidence that only immobile IP<sub>3</sub>Rs parked alongside the sites where SOCE occurs are 'licensed' to evoke Ca<sup>2+</sup> release suggests a mechanism whereby local depletion of the ER may activate SOCE without compromising other Ca<sup>2+</sup>-dependent ER functions.

## Addresses

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## Introduction

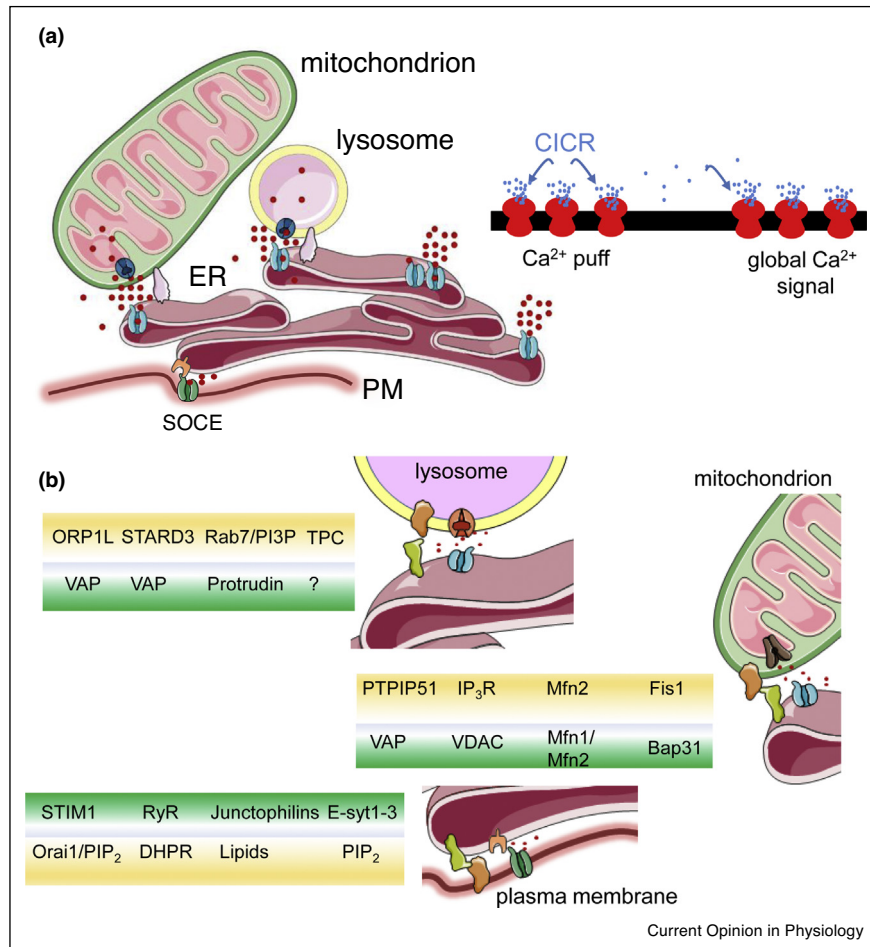
Inositol 1,4,5-trisphosphate receptors (IP<sub>3</sub>Rs) are intracellular Ca<sup>2+</sup> channels that are almost ubiquitously expressed in the endoplasmic reticulum (ER) of animal cells [1]. The discovery, in 1983, that IP<sub>3</sub> stimulates Ca<sup>2+</sup> release from the ER [2] prompted analyses of both the mechanisms linking IP<sub>3</sub> binding to channel opening, and of the organization and functional significance of the resulting Ca<sup>2+</sup> signals. The former established that IP<sub>3</sub>Rs

are large-conductance Ca<sup>2+</sup>-permeable channels [3], opening of which allows Ca<sup>2+</sup> to flow rapidly from the ER. The opening requires binding of IP<sub>3</sub> to all four subunits of an IP<sub>3</sub>R [4<sup>\*</sup>], which then primes the IP<sub>3</sub>R to bind the Ca<sup>2+</sup> that triggers channel opening [3,5]. Structural studies of IP<sub>3</sub>R fragments [6–9] and, more recently, of the complete tetrameric IP<sub>3</sub>R [10<sup>\*\*</sup>,11<sup>\*\*</sup>,12,13<sup>\*</sup>] revealed the initial conformational changes evoked by IP<sub>3</sub> and they are beginning to suggest mechanisms by which they are transduced into rearranged Ca<sup>2+</sup>-binding sites and thence channel opening.

In addressing the functional consequences of IP<sub>3</sub>R activation, most attention focussed on the increase in cytosolic free Ca<sup>2+</sup> concentration ([Ca<sup>2+</sup>]<sub>c</sub>). A recurrent theme has been that as stimulus intensities increase, IP<sub>3</sub>-evoked cytosolic signals progress from local events, 'Ca<sup>2+</sup> puffs' that report the coordinated opening of a few IP<sub>3</sub>Rs within a cluster [14], to global increases in [Ca<sup>2+</sup>]<sub>c</sub> that invade the entire cell [15]. It is assumed that co-regulation of IP<sub>3</sub>Rs by IP<sub>3</sub> and Ca<sup>2+</sup> allows this hierarchical recruitment, with the concentration of IP<sub>3</sub> and spacing of IP<sub>3</sub>Rs tuning the sensitivity of an amplification mechanism (Ca<sup>2+</sup>-induced Ca<sup>2+</sup> release, CICR) [15]. Evidence that all three mammalian IP<sub>3</sub>R subtypes can elicit rather similar Ca<sup>2+</sup> puffs [16,17] has further fuelled speculation that the small clusters of IP<sub>3</sub>Rs from which Ca<sup>2+</sup> puffs originate may be the building blocks of all IP<sub>3</sub>-evoked cytosolic Ca<sup>2+</sup> signals. But that consensus was recently challenged by evidence that global Ca<sup>2+</sup> signals appear not to be built from underlying Ca<sup>2+</sup> puffs [18<sup>\*\*</sup>]. Another puzzle is that while most IP<sub>3</sub>Rs appear to be mobile [19–21], Ca<sup>2+</sup> puffs occur repeatedly at relatively few fixed subcellular locations [22,23]. We address this enigma in the first part of this short review.

Two further observations establish the importance of IP<sub>3</sub>Rs as Ca<sup>2+</sup>-regulating hubs beyond their ability to evoke increases in [Ca<sup>2+</sup>]<sub>c</sub> (Figure 1a). Firstly, IP<sub>3</sub>Rs are the usual route through which physiological stimuli evoke the substantial decrease in ER luminal [Ca<sup>2+</sup>] that activates store-operated Ca<sup>2+</sup> entry (SOCE) [24–28]. The core mechanisms are now clear, with store depletion causing stromal interaction molecule 1 (STIM1) to unfurl cytosolic domains that reach across a narrow gap between the ER and plasma membrane (PM) to capture Orai channels and trigger their opening [29]. It is not, however, clear how the substantial loss of ER Ca<sup>2+</sup> needed to activate STIM1 occurs without compromising other Ca<sup>2+</sup>-dependent ER functions. We speculate on a

Figure 1



IP<sub>3</sub>R<sub>s</sub> regulate different Ca<sup>2+</sup> signals. **(a)** By controlling release of Ca<sup>2+</sup> from the ER, IP<sub>3</sub>R<sub>s</sub> are gateways to different Ca<sup>2+</sup> signals. They can selectively deliver Ca<sup>2+</sup> from the ER to the cytosol, mitochondria or lysosomes, and by decreasing the [Ca<sup>2+</sup>]<sub>ER</sub> they lead to activation of SOCE. Within the cytosol, co-regulation of IP<sub>3</sub>R<sub>s</sub> by IP<sub>3</sub> and Ca<sup>2+</sup> allows regenerative recruitment of IP<sub>3</sub>R activity by CICR. At low concentrations of IP<sub>3</sub>, this CICR is restricted to small clusters of IP<sub>3</sub>R<sub>s</sub> which evoke Ca<sup>2+</sup> puffs. At higher IP<sub>3</sub> concentrations, the regenerative activity propagates across the cell to give a global increase in [Ca<sup>2+</sup>]<sub>c</sub>. **(b)** Some of the candidates proposed to assemble the MCS wherein SOCE occurs, or where Ca<sup>2+</sup> is selectively transferred from IP<sub>3</sub>R<sub>s</sub> in the ER to mitochondria or lysosomes. ORP1L, oxysterol-binding protein-related protein 1L [71]; STARD3, StAR-related lipid transfer domain-3 [72]; P13P, phosphatidylinositol 3-phosphate [65]; TPC, two-pore channel [70]; VAP, vesicle-associated membrane protein-associated proteins; PTPIP51, protein tyrosine phosphatase-interacting protein 51; Mfn1/2, mitofusin 1/2 [73,74]; Fis1, fission 1 homologue [75]; VDAC voltage-dependent anion channel [76]; Bap31, an ER integral membrane protein; RyR, ryanodine receptor; JP, junctophilins [77]; E-Syt, extended synaptotagmins [78]; PIP<sub>2</sub>, phosphatidylinositol 4,5-bisphosphate; DHPR, dihydropyridine receptor.

possible mechanism in light of recent evidence that the IP<sub>3</sub>R<sub>s</sub> that evoke Ca<sup>2+</sup> puffs sit alongside the sites where SOCE occurs [30,31<sup>••</sup>]. Secondly, IP<sub>3</sub>R<sub>s</sub> can selectively deliver Ca<sup>2+</sup> at high concentrations to the mitochondrial surface [32]. This can both regulate mitochondrial mobility [33] and allow Ca<sup>2+</sup> uptake by the low-affinity mitochondrial Ca<sup>2+</sup> uniporter (MCU) [34], with many functional consequences including stimulation of oxidative phosphorylation [35]. The intimate contact is maintained by tether proteins that hold the ER and the outer mitochondrial membrane less than 25 nm apart [36<sup>••</sup>], scarcely any further than the cytosolic dimensions of an

IP<sub>3</sub>R [10<sup>••</sup>]. Among the fifty or so proteins associated with these membrane contact sites (MCS) [35], the relative importance of candidate tethers is unclear, perhaps reflecting some redundancy (Figure 1b). But a key observation is that IP<sub>3</sub>R<sub>s</sub> may themselves form tethers independent of their ability to release Ca<sup>2+</sup> [36<sup>••</sup>]. There is a parallel between SOCE and these synapse-like contacts between ER and mitochondria, in that Ca<sup>2+</sup> channels (Orai or IP<sub>3</sub>R) contribute not only to the Ca<sup>2+</sup> flux within the MCS, but also to its assembly (Figure 1b). Many reviews describe the architecture and function of ER-mitochondrial contacts [34,35,37]. Here, we consider

evidence that there may be a similar association of lysosomes with IP<sub>3</sub>Rs in the ER.

### Licensed IP<sub>3</sub>Rs liaise with SOCE

Ca<sup>2+</sup> puffs repeatedly initiate at rather few fixed sites within a cell [22,23], yet most IP<sub>3</sub>Rs are mobile, at least when overexpressed for live-cell imaging of tagged proteins [19–21]. In cells with endogenous IP<sub>3</sub>R tagged with EGFP using gene editing, most IP<sub>3</sub>Rs form loose clusters, typically comprising about eight IP<sub>3</sub>Rs [31<sup>••</sup>]. Most of these clusters (~70%) are mobile, but Ca<sup>2+</sup> puffs, whether evoked by IP<sub>3</sub> provided by endogenous pathways or uniformly throughout the cytosol by photolysis of caged-IP<sub>3</sub>, invariably occur at immobile clusters and only those located within ER immediately beneath the PM. Hence, only a tiny fraction of the several thousand IP<sub>3</sub>Rs within a cell are ‘licensed’ to evoke Ca<sup>2+</sup> puffs. The licensing mechanism has yet to be defined. It is, however, noteworthy that the licensed IP<sub>3</sub>Rs are parked immediately alongside the MCS where SOCE occurs, although IP<sub>3</sub>Rs are not themselves required to assemble the SOCE MCS [31<sup>••</sup>]. This additional level of IP<sub>3</sub>R regulation, licensing, suggests possible explanations for the enigmas described earlier.

Firstly, ‘licensing’ suggests a regulatory mechanism that both immobilizes some IP<sub>3</sub>Rs and permits them to evoke Ca<sup>2+</sup> puffs. Mobile IP<sub>3</sub>Rs do not evoke Ca<sup>2+</sup> puffs. Since the mobile and immobile pools of IP<sub>3</sub>Rs do not freely mix, at least over tens of minutes [31<sup>••</sup>], the function of most IP<sub>3</sub>Rs, the mobile ones and those remote from the PM, remains mysterious. They may provide a pool from which IP<sub>3</sub>Rs can be drawn for licensing; they may have different roles in Ca<sup>2+</sup> signalling, for example in selectively delivering Ca<sup>2+</sup> to mitochondria or lysosomes (many of which do not lie immediately beneath the PM); or they may have roles unrelated to Ca<sup>2+</sup> signalling. The latter would be consistent with many different proteins, not all obviously related to Ca<sup>2+</sup> signalling, associating with IP<sub>3</sub>Rs [38].

Secondly, the location of licensed IP<sub>3</sub>Rs alongside the MCS where STIM1 accumulates after store depletion [31<sup>••</sup>] suggests a mechanism whereby activation of SOCE by physiological stimuli, namely those that evoke IP<sub>3</sub> formation, might activate STIM1, and thereby SOCE, without substantially depleting the entire ER of Ca<sup>2+</sup> [30,39]. We speculate that an ER structure with a short neck and flattened head, somewhat reminiscent of a dendritic spine, might allow licensed IP<sub>3</sub>Rs to locally and substantially deplete ER within a SOCE MCS of Ca<sup>2+</sup> (Figure 2). Each of these junctions might then function as a ‘digital switch’, such that STIM1 within the depleted ER would be fully active after stimulation of associated IP<sub>3</sub>Rs. Graded activation of SOCE within a cell would come from recruitment of each MCS, and without global depletion of ER Ca<sup>2+</sup> (Figure 2b). Additional

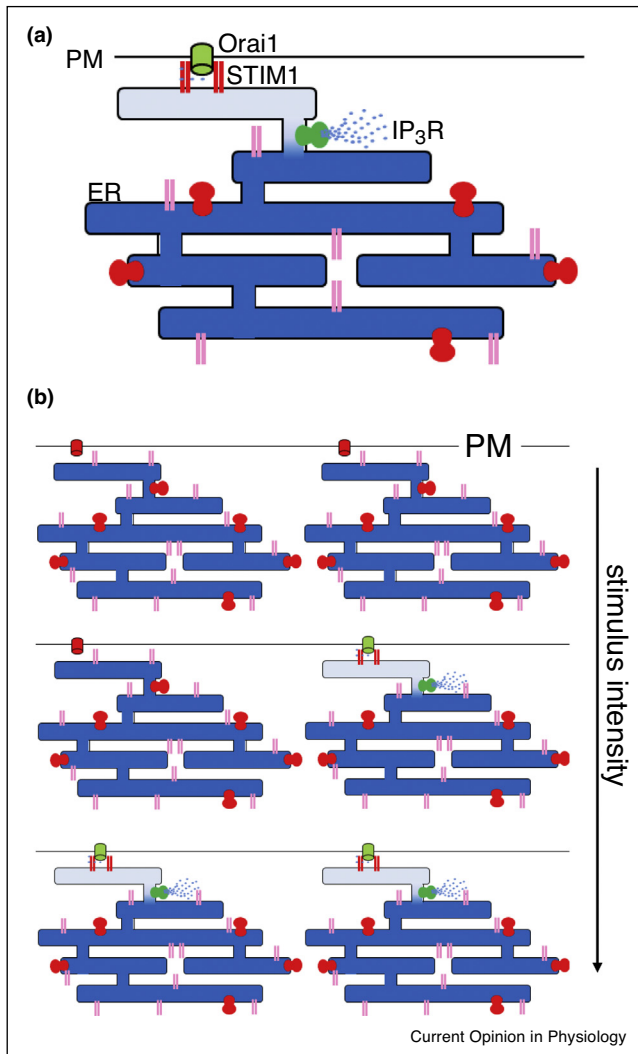
mechanisms, notably hetero-oligomerization of STIM2 with STIM1, might tune the Ca<sup>2+</sup> sensitivity of Orai1 activation to respond to less than complete store emptying by combining the greater Ca<sup>2+</sup> sensitivity of STIM2 [25] and its enhanced affinity for phosphatidylinositol 4,5-bisphosphate (PIP<sub>2</sub>) [40], with the greater efficacy of STIM1 in activating Orai1 [41<sup>••</sup>].

### IP<sub>3</sub>R and lysosomes – more intimate liaisons

Lysosomes, with their luminal cocktail of hydrolytic enzymes and acidic pH, are best known as degradative organelles [42,43], but there is also evidence that they make important contributions to intracellular Ca<sup>2+</sup> handling [44]. Experimental analyses of the latter are compromised by the hostile luminal environment of the lysosome, which constrains use of protein-based indicators and demands meticulous pH-corrections to the properties of classical Ca<sup>2+</sup> indicators. An ingenious construct that tethers conventional pH and Ca<sup>2+</sup> indicators to a DNA backbone may provide the much-needed tool to effectively measure [Ca<sup>2+</sup>] within lysosomes [45<sup>••</sup>]. At present, reliable measures of luminal free [Ca<sup>2+</sup>] for lysosomes are scarce, but they suggest a concentration comparable (350–600 μM) to that of the ER [45<sup>••</sup>,46,47], although lysosomes enclose a much smaller intracellular volume (perhaps 500-times less than that of the ER) [48]. There is, however, persuasive evidence that lysosomes express Ca<sup>2+</sup>-permeable channels, including two-pore channels (TPC1-3), transient receptor potential mucolipin (TRPML1-3) and type 2 purinoceptors (P2X<sub>4</sub>) [44,49], and that Ca<sup>2+</sup> released from lysosomes regulates membrane fission and fusion within the endocytic pathway [42,50–52]. Indeed, Ca<sup>2+</sup> release through TPCs is required for trafficking of Ebola virus through late stages of the endocytic pathway [53]. In addition to this local Ca<sup>2+</sup> signalling within the endocytic pathway, lysosomes may also initiate larger cytosolic Ca<sup>2+</sup> signals when Ca<sup>2+</sup> released through lysosomal channels, triggers Ca<sup>2+</sup>-induced Ca<sup>2+</sup> release from the ER through IP<sub>3</sub>Rs or ryanodine receptors [54,55].

The means by which lysosomes accumulate Ca<sup>2+</sup> is unknown. Considerable evidence, including an increase in lysosomal pH after addition of Ca<sup>2+</sup> [56], and loss of lysosomal Ca<sup>2+</sup> after dissipation of the lysosomal pH gradient [46,47,57,58], suggest an important role for luminal H<sup>+</sup>, but there is no evidence that placental mammals express a lysosomal Ca<sup>2+</sup>-H<sup>+</sup> exchanger (CAX) [59]. Several reviews consider alternative lysosomal pH-dependent mechanisms for Ca<sup>2+</sup> uptake [44,45<sup>••</sup>,49], and a recent report suggests a possible role for a P-type ATPase (ATP13A2), distinct from those that mediate Ca<sup>2+</sup> transport across other biological membranes, in mediating lysosomal Ca<sup>2+</sup> uptake [45<sup>••</sup>]. This is intriguing because mutations in ATP13A2 are associated with a juvenile form of Parkinson’s disease [60] and defective autophagy [61], but there is presently no direct evidence that

Figure 2



Local regulation of SOCE by licensed IP<sub>3</sub>Rs. **(a)** Licensed clusters of IP<sub>3</sub>Rs (green) parked alongside SOCE-MCS may allow substantial loss of Ca<sup>2+</sup> from ER close to the PM, while negligibly affecting the much larger volume of ER deeper in the cell. The large local decrease in ER [Ca<sup>2+</sup>] may then locally activate STIM1, and thereby the Ca<sup>2+</sup> channel, Orai1, that mediates SOCE. Active and inactive STIM1 dimers are shown in red and pink, respectively. **(b)** Digital recruitment of all-or-nothing SOCE-MCS may allow graded activation of SOCE in cells.

ATP13A2 is a Ca<sup>2+</sup> pump. The limited evidence available suggests that whatever the lysosomal Ca<sup>2+</sup> uptake mechanism, it probably has relatively low affinity for Ca<sup>2+</sup> [62]. There is, therefore, a problem similar to that faced by mitochondria and their MCU, namely the organelles acquire Ca<sup>2+</sup> from the cytosol, where the global [Ca<sup>2+</sup>]<sub>c</sub> is too low to fuel the low-affinity uptake systems. For mitochondria, the solution is provided by having the ER, with its high-affinity Ca<sup>2+</sup> pump (the SR/ER Ca<sup>2+</sup>-ATPase, SERCA), concentrate Ca<sup>2+</sup> from the cytosol and then squirt it back at MCU through large-conductance IP<sub>3</sub>Rs

(Figures 1b and 3 a). Recent evidence suggests a similar scheme for lysosomes.

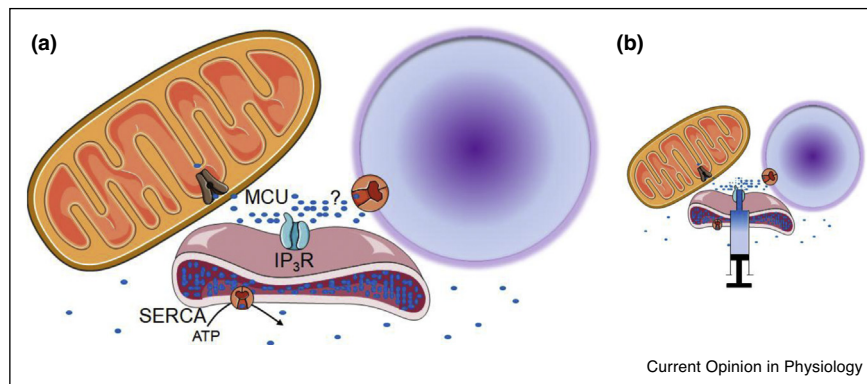
The first evidence that lysosomes selectively sequester Ca<sup>2+</sup> released by IP<sub>3</sub>Rs came from cells in which cytosolic Ca<sup>2+</sup> signals evoked by receptors that stimulate IP<sub>3</sub> formation were potentiated when lysosomes were perturbed. SOCE-evoked signals were unaffected [63,64<sup>\*\*</sup>]. Furthermore, a low-affinity Ca<sup>2+</sup>-sensor expressed on the cytosolic surface of lysosomes, reported large Ca<sup>2+</sup> signals in response to IP<sub>3</sub>-evoked Ca<sup>2+</sup> release, but not in response to SOCE [64<sup>\*\*</sup>]. Close contacts (MCS) between ER and lysosomes are well documented [65–67], and recent evidence suggests that these contacts form preferentially at ER populated with IP<sub>3</sub>Rs [64<sup>\*\*</sup>]. Another study, which used cytosolic Ca<sup>2+</sup> sensors targeted to a lysosomal Ca<sup>2+</sup> channel to determine lysosomal Ca<sup>2+</sup> content, concluded that Ca<sup>2+</sup> released from the ER through IP<sub>3</sub>Rs is required to fuel lysosomal Ca<sup>2+</sup> uptake [68<sup>\*</sup>]. There are, however, some concerns in that the sensor used had an affinity for Ca<sup>2+</sup> that was too high to selectively report local Ca<sup>2+</sup> signals, and there was some reliance on glycyl-L-phenylalanine 2-naphthylamide (GPN) to release Ca<sup>2+</sup> from lysosomes, when recent work shows that GPN evokes Ca<sup>2+</sup> release from the ER [69<sup>\*</sup>]. The studies concur, however, in suggesting that Ca<sup>2+</sup> uptake by lysosomes is fuelled by Ca<sup>2+</sup> release from the ER [63,64<sup>\*\*</sup>,68<sup>\*</sup>], and in suggesting that the lysosomal pH gradient is not required for that Ca<sup>2+</sup> uptake [64<sup>\*\*</sup>,68<sup>\*</sup>], although it is required to maintain the ER-lysosome MCS [64<sup>\*\*</sup>].

Hence, intimate liaisons between ER and lysosomes allow a two-way traffic of Ca<sup>2+</sup> between them. Ca<sup>2+</sup> release through lysosomal Ca<sup>2+</sup> channels is amplified by CICR from ER channels [54,55]; and the ER, through IP<sub>3</sub>Rs [64<sup>\*\*</sup>,68<sup>\*</sup>] and other leak channels [64<sup>\*\*</sup>] provides Ca<sup>2+</sup> for a low-affinity lysosomal Ca<sup>2+</sup>-uptake system. The latter provides a close analogy with MCU in mitochondria. In both cases, the ER works like a piston, concentrating Ca<sup>2+</sup> from the cytosol into the ER lumen and then delivering it rapidly through a large-conductance channel, the IP<sub>3</sub>R, at sufficient local concentration to allow the organelle to sequester it with a low-affinity uptake system (Figure 3b). An unresolved question is whether the mechanisms that license IP<sub>3</sub>Rs to evoke cytosolic Ca<sup>2+</sup> puffs serve also to license the IP<sub>3</sub>Rs that deliver Ca<sup>2+</sup> to lysosomes and mitochondria.

We noted earlier that amongst the many proteins that may contribute to MCS between ER and mitochondria, IP<sub>3</sub>R fulfilled a structural role in assembling the junction [36<sup>\*\*</sup>]. At SOCE MCS too, the Orai Ca<sup>2+</sup> channel may contribute to assembly of the junction. This 'hijacking' of Ca<sup>2+</sup> channels to assemble the MCS within which they function may extend also to the ER-lysosome MCS, because although IP<sub>3</sub>Rs are not required for their assembly, there is evidence that TPC1 regulates assembly of the MCS between ER and endosomes [70].



Figure 3



Delivering cytosolic Ca<sup>2+</sup> to other organelles via the ER. **(a)** High-affinity Ca<sup>2+</sup> pumps (SERCA) allow the ER to concentrate Ca<sup>2+</sup> from the cytosol, where its concentration (~100 nM) is too low to allow sequestration by low-affinity uptake systems in mitochondria (MCU) or lysosomes (unknown, ?). When they open, IP<sub>3</sub>Rs, with their large single-channel conductances, allow Ca<sup>2+</sup> to be delivered from the ER to the organellar uptake systems at sufficient concentration to support their activity. **(b)** We can regard the ER as a piston that concentrates Ca<sup>2+</sup> from the cytosol before squirting it at organelles within enclosed MCS.

## Concluding remarks

IP<sub>3</sub>Rs provide an essential link between receptors in the PM that transduce extracellular stimuli into an increase in intracellular IP<sub>3</sub> concentration, and the redistribution of Ca<sup>2+</sup> across biological membranes. By rapidly releasing Ca<sup>2+</sup> from the ER, IP<sub>3</sub>Rs selectively deliver Ca<sup>2+</sup> to the cytosol or other organelles, and they activate SOCE (Figure 1). ‘Licensed’ IP<sub>3</sub>Rs parked alongside SOCE-MCS may allow local regulation of SOCE without compromising other ER functions (Figure 2). Comparisons of ER contacts with lysosomes and mitochondria suggest a general scheme, wherein the ER behaves like a piston, concentrating Ca<sup>2+</sup> from the cytosol, where its concentration is very low, and then rapidly releasing it through large-conductance IP<sub>3</sub>Rs into enclosed MCS, where the high local [Ca<sup>2+</sup>] is sufficient to fuel Ca<sup>2+</sup> uptake by the low-affinity Ca<sup>2+</sup> uptake systems of the organelles (Figure 3).

## Conflict of interest statement

Nothing declared.

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