

IP₃ receptors in cell survival and apoptosis: Ca²⁺ release and beyond

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Abstract Inositol 1,4,5-trisphosphate receptors (IP₃Rs) serve to discharge Ca²⁺ from ER stores in response to agonist stimulation. The present review summarizes the role of these receptors in models of Ca²⁺-dependent apoptosis. In particular we focus on the regulation of IP₃Rs by caspase-3 cleavage, cytochrome *c*, anti-apoptotic proteins and Akt kinase. We also address the evidence that some of the effects of IP₃Rs in apoptosis may be independent of their ion-channel function. The role of IP₃Rs in delivering Ca²⁺ to the mitochondria is discussed from the perspective of the factors determining inter-organellar dynamics and the spatial proximity of mitochondria and ER membranes.

Keywords Apoptosis · Calcium · Endoplasmic reticulum · IP₃ · IP₃ receptor · Mitochondria

Abbreviations

NFAT	Nuclear factor of activated T-cells
ER	endoplasmic reticulum
MEF	murine embryo fibroblasts
HIV	Human immunodeficiency virus
STS	staurosporine
Bcl-2	B cell lymphoma 2
Bcl-xL	B cell lymphoma xL
IP ₃ R	Inositol 1,4,5-trisphosphate receptors
RyR	Ryanodine receptor

FasL	Ligand for Fas receptor (CD95)
VDAC	Voltage dependent anion selective channel
[Ca ²⁺] _c	cytoplasmic free [Ca ²⁺]
[Ca ²⁺] _m	mitochondrial matrix free [Ca ²⁺]
OMM	outer mitochondrial membrane
IMM	inner mitochondrial membrane
PTP	permeability transition pore

1 Introduction

IP₃ receptors are ligand-gated channels that function to release Ca²⁺ from intracellular Ca²⁺ stores (predominantly the endoplasmic reticulum (ER)) in response to IP₃ generation initiated by agonist binding to cell-surface receptors (reviewed in Refs. [1–3]). The ensuing Ca²⁺ signal plays a fundamental role in modulating a diverse range of cellular responses including muscle contraction, exocytosis, motility, fertilization, proliferation and gene expression. The increased energy demand required for carrying out these processes is met, in part, from the accumulation of some of the released Ca²⁺ by the mitochondria with the resulting Ca²⁺-dependent activation of key intramitochondrial enzymes linked to ATP production [4, 5]. The efficiency of this process is greatly enhanced by placement of ER and mitochondrial membranes in close proximity [6, 7]. Thus the physiological release of Ca²⁺ through IP₃Rs serves important signaling and “housekeeping” roles in maintaining normal function during agonist-mediated cell activation. However, in the presence of an apoptotic stimulus, the IP₃R-mediated Ca²⁺ release can activate apoptotic pathways by inducing the release from mitochondria of a number of pro-apoptotic factors including cytochrome *c*. The present article is focused on reviewing recent studies that implicate the IP₃R as an important molecular component regulating apoptosis induced by a number

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of different stimuli. In particular, we have emphasized the role of the IP₃R in establishing macromolecular complexes on the surface of the ER membranes and in modulating the linkage between ER and mitochondrial membranes.

2 The pro-apoptotic role of IP₃Rs

There is general agreement in the literature that Ca²⁺ efflux from the ER and Ca²⁺ accumulation into the mitochondria is linked to the effects of various apoptotic stimuli [6–10]. With certain stimuli, such as ligation of T-cell [11, 12] or B-cell [13] receptors, this process clearly involves stimulation of phospholipase-C activity and generation of IP₃. The functional outcome in lymphocytes may result in either enhanced proliferation or apoptosis, depending on the maturation status of the cells or the intensity of the stimulation [14]. Other classic death ligands such as TNF α or Fas can also promote IP₃ generation in some but not all cells [15–17]. Yet other apoptotic stimuli such as ceramide [18], cytotoxic drugs (e.g. STS [19, 20] and cisplatin [21]), or manipulations such as withdrawal of trophic factors [22], can also cause ER Ca²⁺ release. In these cases an initiating role for IP₃ elevation in these processes has not been demonstrated and the ER Ca²⁺ release could be precipitated secondarily to changes in mitochondrial permeability (Fig. 1) or activation of ER stress responses [23, 24]. Thus although a role for ER to mitochondrial redistribution of Ca²⁺ has been implicated in many models of apoptosis, a primary role for IP₃ generation and activation of IP₃Rs in this process has been examined in only a few instances.

The magnitude of the Ca²⁺ translocated from the ER will be determined by the levels of ER Ca²⁺ and the activity of the efflux and influx pathways that cycle Ca²⁺ across ER membranes. The primary influx pathway of the ER are the SERCA pumps. The primary efflux pathway across the ER membranes (in the presence of IP₃) is the IP₃R but additional poorly characterized ‘leak’ pathways are also believed to play important roles (reviewed in Ref. [25]). Figure 1 depicts a number of known apoptotic regulatory feed-back loops that involve Ca²⁺. Examination of this complex regulatory network clearly indicates that IP₃Rs could potentially play a central role in the initiation and/or progression of the apoptotic cascade.

The experimental evidence for an involvement of IP₃Rs in apoptosis have predominantly come from experiments in which the expression level of the receptor protein has been manipulated. There are three isoforms of IP₃Rs that are expressed in differing amounts in various cell types. A functional IP₃R Ca²⁺ channel is composed of tetramers that can be either homotetrameric or, to a lesser extent, heterotetramers of different isoforms [1–3]. In initial studies, it was shown that the levels of the type-III IP₃R protein were in-

creased during apoptosis in T-lymphocytes and that apoptosis could be prevented by antisense-RNA to the type-III but not the type-I isoform [26]. These observations were later replicated in other models of apoptosis [27]. A role for the type-III isoform has also been proposed recently based on selective siRNA knockdown studies of IP₃R isoforms in CHO cells [28]. This preferential role was suggested to be the result of enhanced colocalization of the type-III isoform adjacent to mitochondrial membranes [28]. However, there is also evidence that stable knockdown of the type-I IP₃R in Jurkat T-lymphocytes can inhibit responsiveness to different apoptotic stimuli [29]. siRNA for the type-I IP₃R also diminishes apoptosis in response to FasL in Jurkat lymphocytes [17]. The introduction of constitutively active calcineurin restored the sensitivity of the type-I deficient cells and also restored the ability of the cells to dephosphorylate and generate the pro-apoptotic factor Bad [30], supporting the view that this is one pathway that is downstream of cytosolic Ca²⁺ elevation (c.f. Fig. 1). Immunoblotting of the type-I deficient Jurkat cells also showed decreased levels of the type-II and type-III isoforms [29], so it is difficult to conclude from these particular studies that only the type-I IP₃R plays a unique role. Studies on the T-cells of mice deficient in the type-I IP₃R did not reveal any abnormalities in T-cell function or development [31]. Similarly, mice deleted for both type-II and III IP₃R isoforms showed severe defects in the digestive system but were able to survive with appropriate feeding regimens, indicating that a single IP₃R isoform may be sufficient to sustain many essential signaling requirements [32].

The issue of isoform specificity has also been addressed using chicken B-lymphocyte DT-40 cell lines in which the genes for individual IP₃R isoforms have been selectively deleted by homologous recombination [33]. The loss of any one isoform did not affect B-cell receptor mediated apoptosis whereas the loss of all three isoforms (TKO cells) led to marked inhibition [34]. These data suggest that there is considerable redundancy in the ability of each IP₃R isoform to participate in the apoptotic process. The extent of the inhibition of apoptosis reported for TKO DT-40 cells in various studies has been variable. In the original report, 30% of the TKO cells stimulated with 10 μ g/ml anti-IgM showed an apoptotic phenotype in 24h compared to 74% in WT cells. A similar magnitude of inhibition was also reported by Assefa et al. for both anti-IgM and STS [20]. Others have found a much larger suppression of apoptosis under similar experimental conditions, although the amount of cell death does increase in the TKO cells as exposure to the apoptotic stimulus is prolonged [35]. Overall, the data suggest that IP₃R-dependent apoptotic pathways play a major role, particularly in the early stages of the apoptotic response, but that alternative less efficient apoptotic pathways can be recruited in the absence of IP₃Rs.

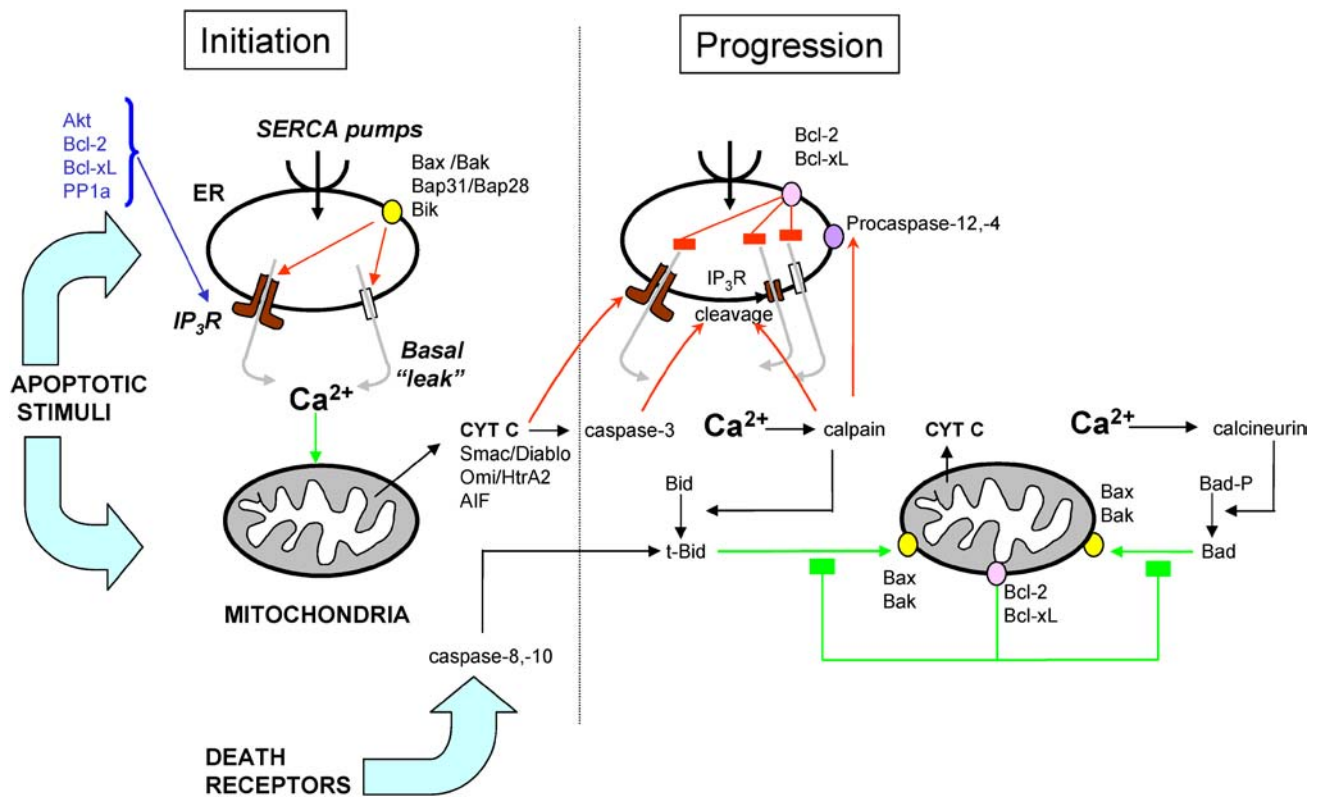


Fig. 1 Ca²⁺ dependent networks regulating apoptosis. The scheme incorporates some of the prevailing hypothesis regarding Ca²⁺-dependent steps in apoptotic cascades. The initiating step induced by divergent stimuli is depicted as an enhanced translocation of Ca²⁺ from the ER to the mitochondria resulting in mitochondrial Ca²⁺ overload provoking the release of a number of pro-apoptotic proteins into the cytoplasm, including cytochrome *c* (CYT C). This intrinsic pathway is activated in response to STS and by many other cytotoxic agents including those that cause ER stress. The extrinsic pathway is activated by cell surface death receptors of the TNF- α superfamily. Downstream pathways which target steps occurring in the ER are depicted in red and those

that target the mitochondria are depicted in green. Arrowheads and bars indicate activation and inhibitory steps respectively. Enhanced Ca²⁺ efflux from the ER can be promoted by cytochrome *c* directly activating IP₃R or indirectly by caspase or calpain cleavage of IP₃R generating "leaky" ion channels. The Ca²⁺ leak pathways in the ER membrane can be positively or negatively modulated by a number of Bcl-2 family of proteins. Proteins that bind to IP₃R but whose functional effects have not been firmly established are shown in blue. Ca²⁺ dependent steps that target the mitochondria include calpain cleavage of Bid and the calcineurin dependent dephosphorylation of Bad. For additional details see text

3 Molecular mechanism(s) of IP₃R function in apoptosis

In addition to the direct role of IP₃R in the initiation of apoptosis by providing a conduit for ER-to-mitochondria Ca²⁺ transfer, there are several additional feedback mechanisms that have been proposed which allow IP₃R to play a role in amplifying Ca²⁺ dependent apoptotic pathways. These include the cleavage of the receptor by caspases (and possibly calpains) which is proposed to provide an enhanced ER Ca²⁺ leak pathway, and the direct binding and regulation of the IP₃R Ca²⁺ channel by several key players in apoptosis including cytochrome *c* and the anti-apoptotic proteins Bcl-2 and Bcl-xL. In addition, the IP₃R can be directly phosphorylated by Akt kinase. Current information on these regulatory mechanisms are summarized below.

3.1 Regulation by proteolytic cleavage

Mikoshiha and colleagues first demonstrated that the type-I IP₃R was cleaved by caspase-3 in Jurkat T-cells in response to STS or activation of the Fas pathway [36]. Cleavage *in vitro* of the cerebellum IP₃R was reported to inactivate IP₃-mediated Ca²⁺ release. The basic observation of caspase-3 mediated IP₃R proteolysis has since been confirmed in Jurkat T-cells [37], DT-40 B lymphocytes [20], SH-SY5Y neuroblastoma cells [38] and PC-12 cells [39]. The cleavage site was localized to aspartate 1892 within a DEVD caspase-3 proteolytic site. Proteolysis would be expected to generate a 95 kDa C-terminal fragment containing the six transmembrane domains that form the ion channel, and a 215 kDa N-terminal fragment that includes the IP₃ binding region of the receptor (Fig. 2(A)). Although the 95 kDa C-terminal band is experimentally observed [20, 36–38], the status of the

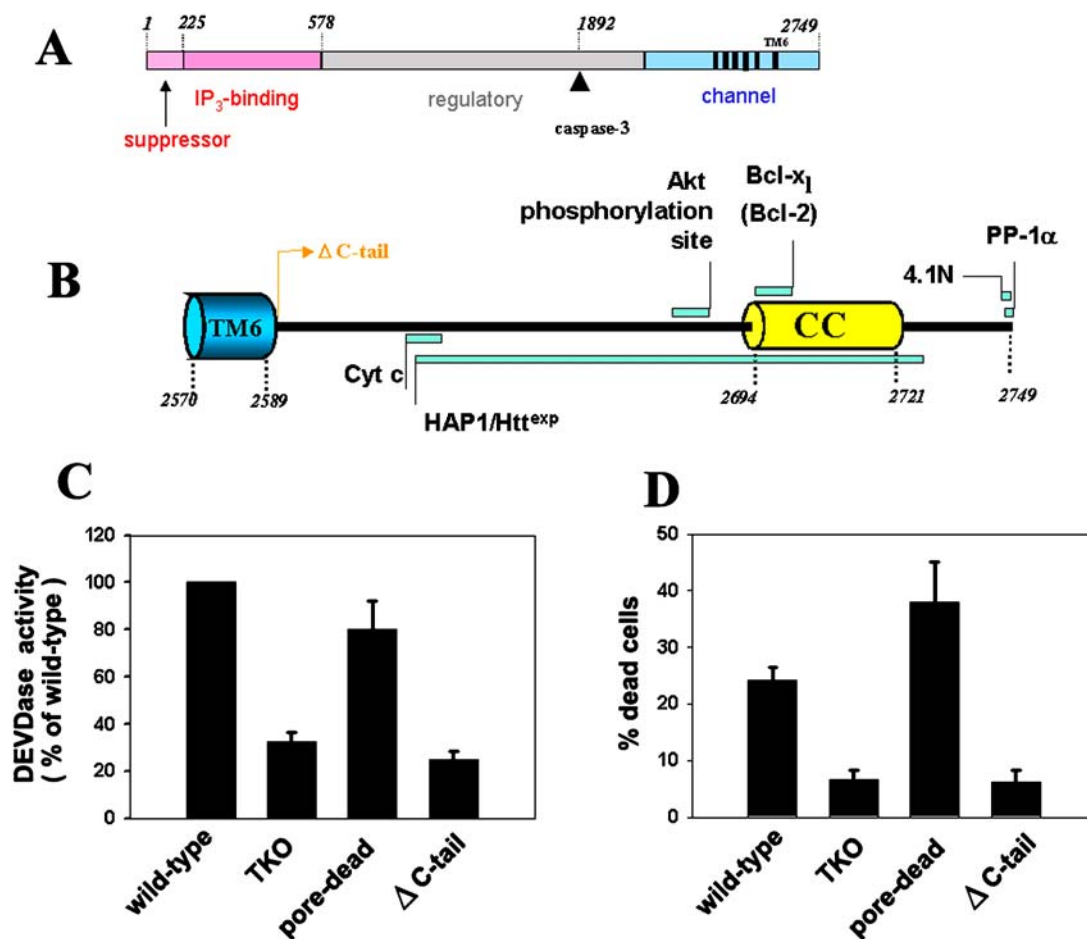


Fig. 2 Domain organization and protein/protein associations in the C-terminal tail of IP₃Rs. *Panel A* shows the domain organization of IP₃Rs with the N-terminal binding domain, C-terminal channel domain and intervening regulatory domain. The small region at the N-terminus which acts to suppress IP₃ binding and the predominant site of caspase-3 cleavage is also shown. *Panel B* shows in more detail the 159 amino acid, cytosol-exposed, C-terminal tail which contains interaction sites for several proteins that are key regulators of apoptosis. Abbreviations are CC, coiled-coil domain; TM6, transmembrane domain 6; PP-1α protein phosphatase-1α. Other abbreviations are defined in the text. *Panel C*: DT-40 chicken B-lymphocytes containing or lacking all 3 IP₃R isoforms (wild-type and TKO respectively) were incubated with

0.5 μM STS for 6 h and caspase-3 activity was measured in cell lysates with a fluorometric assay measuring the hydrolysis of DEVD-AMC. The data was compared to stable DT-40 cell lines made in the TKO background which contained the D2550A pore-dead mutant or a construct of the type-I IP₃R missing the 159 amino-acid tail (ΔC-tail; see Panel B). *Panel D*: The conditions were the same as for Panel D except that cell viability was measured with an MTS assay. The data in Panels D & E are the mean + S.E.M. of 3–5 independent experiments (M.T. Khan & S.K. Joseph, unpublished). Technical details of the preparation of stable DT-40 cell lines, cell culture and caspase-3 assay conditions are as described in Ref. [96]

N-terminal segment of the receptor after caspase-3 cleavage *in vivo* has not been examined. At least *in vitro*, the N-terminal segment formed is smaller than the expected 215 kDa, suggesting that caspase-3 may cleave at multiple sites [40]. Each IP₃R isoform contains multiple DXXD consensus caspase-3 cleavage sites, although the DEVD site at position 1892 is unique to the type-I isoform. In the initial study using Jurkat T-cells only the type-I isoform was reported to be a substrate [36]. However, in another study also using Jurkat T-cells, all 3 isoforms were degraded in response to TNFα stimulation but only the type-I and type-II IP₃R degradation was sensitive to the pan-caspase inhibitor z-VAD-fmk whereas type-III IP₃R degradation was sensitive to a calpain inhibitor [37].

Two key questions regarding the caspase-3 cleavage of IP₃R receptors remain to be addressed. Is proteolysis of IP₃Rs an essential step in the pathway of Ca²⁺-dependent apoptosis and what are the functional consequences of generating cleaved forms of the IP₃R? MCF-7 breast carcinoma cells which lack caspase-3 do not degrade IP₃Rs in response to TNFα or STS [36]. This indicates that IP₃R cleavage cannot be mediated by other effector caspases present in MCF-7 cells, including caspase-7, which has the same substrate specificity as caspase-3. Yet, MCF-7 cells undergo Ca²⁺-dependent apoptosis [41] and show a number of features common to other cells, including a Bcl-2 mediated decrease in ER Ca²⁺ [42, 43]. Boehning et al. observed minimal cleavage of IP₃Rs during the early stages of apoptosis induction by

STS in HeLa or PC12 cells when compared to the more rapid and extensive cleavage of established substrates such as poly (ADP-ribose) polymerase [39]. However, complete loss of IP₃R was observed at 24–48 h and this loss was blocked by caspase-3 inhibitors. Thus IP₃R cleavage may reflect a very distal event in apoptosis/necrosis that may be associated with an inactivation of Ca²⁺ channel function [36].

The strongest evidence in favor of an essential role of caspase-3 proteolysis comes from the finding that a non-cleavable DEVD mutant, in contrast to the wild-type type-I IP₃R, was unable to support apoptosis when introduced into the TKO DT-40 cells [20]. In this study, the addition of STS to wild-type cells caused a prolonged increase of cytosolic free Ca²⁺ which was blocked by caspase-3 inhibition. This increase was absent in the DEVD mutant. The interpretation of the data is consistent with the view that caspase-3 cleaves the IP₃R and the resulting 95 kDa “channel-only” fragment is responsible for the persistent cytosolic Ca²⁺ increase. The GFP-tagged version of the 95 kDa “channel-only” fragment induces an ER Ca²⁺ leak when expressed in COS-7 or HeLa cells [44] and this has lent support to the idea that the “channel-only” IP₃R fragment is a constitutively leaky Ca²⁺ channel (reviewed in Ref. [45]). However, expression of the untagged version of the 95 kDa “channel-only” domain in TKO DT-40 cells does not by itself induce ER Ca²⁺ depletion [20] (Bhanumathy & Joseph; unpublished observations). An enhanced leak in the presence of an apoptotic stimulus could arise if the truncated channel is gated open by some change induced by the stimulus e.g. alteration in the channel thiol redox state as a consequence of increased ROS generation. Alternatively, the “channel-only” or the N-terminal fragment could indirectly regulate the activity of another Ca²⁺ leak pathway in the ER. If the caspase-3 cleavage of IP₃R were critical for causing the Ca²⁺ changes observed in the cytosol and ER in response to various apoptotic stimuli, then it would be expected that these changes would be blocked by caspase-3 inhibitors. This is indeed the case for the STS-induced Ca²⁺ changes reported by Assefa et al. [20] but this is not a general finding. For example, the prolonged oscillations in cytosolic Ca²⁺ induced by STS in HeLa cells [39] or the STS-mediated depletion of ER Ca²⁺ in PC-3 prostate adenocarcinoma cells [19] are both insensitive to caspase-3 inhibitors. Thus although the data for caspase-3 IP₃R cleavage appears strong in the DT-40 B-cell model, a general involvement of caspase-3 cleavage of IP₃R in the early stages of the apoptotic cascade and a role for leaky channels formed from proteolytic fragments remain to be firmly established.

Activation of calpains have been observed in many models of apoptosis [46] and calpains have also been reported to cleave IP₃R [47, 48]. The Ca²⁺ dependent cleavage of the type-I IP₃R in cerebellum membranes produced C-terminal fragments of 95 and 130 kDa [47]. However, in those models

of apoptosis where a degradation of IP₃R can be observed, the inclusion of calpain inhibitors does not prevent IP₃R proteolysis ([36, 39]; but c.f. [37]). Even if calpain activation is not important for IP₃R degradation, there is strong evidence that activation of calpain plays important roles in regulating Ca²⁺-dependent steps involved in apoptosis. For example μ -calpain is involved in the cleavage and activation of *t*-Bid which, in turn, plays a prominent role in the release of pro-apoptotic factors from the mitochondria [49, 50]. In addition the stimulation of calpain activity has been linked to the activation of ER-associated caspase-12 [51], the activation of calcineurin [52] and the elevation of cytosolic Ca²⁺ as a result of the cleavage of the plasma membrane Na⁺/Ca²⁺ exchanger [53].

3.2 Regulation by cytochrome *c*

Interactions with cytochrome *c* were detected in a yeast two-hybrid screen using the C-terminal segment of the type-I IP₃R as bait [39]. The IP₃R Ca²⁺ channel exhibits a biphasic dependence on [Ca²⁺] with low concentrations of Ca²⁺ stimulating and high concentrations of Ca²⁺ inhibiting channel activity [54]. The functional effect of cytochrome *c* on the channel was to prevent the inhibitory effect of Ca²⁺ thereby disrupting the normal mechanism that acts to shut off the channel. The uptake of Ca²⁺ into mitochondria can also indirectly influence local feedback effects of Ca²⁺ on IP₃R [55, 56]. The exact mechanism by which Ca²⁺ activates or inhibits the IP₃R Ca²⁺ channel is not well understood. However, it is of interest that the site of cytochrome *c* binding on the IP₃R has been localized to a 16 amino-acid stretch (2621–2636) present within the short 160 amino acid C-terminal tail of the receptor [57]. This segment projects into the cytosol from the end of the transmembrane domain helix 6 which lines the ion-channel pore [58]. The C-terminal tail is also the binding site for Bcl-xL, protein phosphatase-1 α and contains the phosphorylation site for Akt kinase (Fig. 2(B)). It has been assumed that these effector proteins must be able to easily access the cytosol-exposed C-terminal segment in the intact cell. The 3D conformation of the IP₃R has been reconstructed from EM studies but the limited resolution of the structures does not allow the location of the C-terminal tail to be identified (reviewed in Ref. [58]). The presence of Ca²⁺ causes a large conformational change in the IP₃R that has been visualized in EM as a transition from a “square” to a “wind-mill” structure [59]. In the native receptor, this conformational change could also alter the accessibility of the C-terminal tail to cytochrome *c* and to the other molecules that bind to this region. The finding that cytochrome *c* has its largest functional effect at concentrations of Ca²⁺ > 1 μ M is consistent with this idea [39].

Analysis of ER enriched membrane fractions in cells undergoing apoptosis indicates that cytochrome *c* translocates

from the mitochondria to ER membranes [39]. Under these conditions, the 100,000×*g* cytosol fraction appears devoid of cytochrome *c* except in DT-40 cells lacking IP₃Rs [39]. This has led to the proposal that cytochrome *c* released from the mitochondria is sequestered at ER membranes by high-affinity binding to IP₃Rs. Recently, a translocation of cytochrome *c* to the ER was also noted in the hippocampus after ischemia/reperfusion injury [60]. However, a principal function of cytochrome *c* is to associate with Apaf-1 and facilitate the formations of the apoptosome that ultimately leads to the activation of caspases. Since these complexes are formed in the cytosol, a minimal amount of cytochrome *c* must be released into this compartment. This has been confirmed in many studies which have used digitonin-permeabilization (e.g. [61]) or single cell imaging of GFP-cytochrome *c* [62].

Evaluation of the mitochondrial release of cytochrome *c* in single cells indicate that, once initiated, the process is rapid (<5 min) and encompasses all the mitochondria in the cell [62]. The cytochrome *c*/IP₃R interaction has been suggested to co-ordinate the global release of cytochrome *c*. It is proposed that a partial release of cytochrome *c* would activate Ca²⁺ release from adjacent IP₃Rs that would feed back on the mitochondria to promote further permeabilization and additional cytochrome *c* release [39, 63]. A requirement for this feed-back loop is supported by studies which have employed the 16 amino-acid peptide encoding the highly conserved cytochrome *c* binding motif from IP₃Rs. Addition of a cell-permeant version of the peptide exerts dominant negative effects and inhibited apoptosis induced by either STS or FasL, presumably by competing for cytochrome *c* binding to the IP₃R [57].

In summary, the evidence points to an important role for the interaction of cytochrome *c* with IP₃Rs in the regulation of apoptosis. Further characterization of the cytochrome *c* effects on Ca²⁺ regulation of the IP₃R at the single channel level and identification of the sites on cytochrome *c* that are relevant for IP₃R interaction should provide additional mechanistic insights. An absolute requirement of IP₃Rs for global cytochrome *c* release would predict that cytochrome *c* release would be severely inhibited in the DT 40 TKO cell line from which IP₃Rs are absent. Although cytochrome *c* translocation to the ER membrane fraction is eliminated, a release into the cytosol is still observed in TKO cells stimulated with anti-IgM [39]. Thus, alternative compensatory mechanisms must be in place to facilitate limited mitochondrial permeabilization in the absence of IP₃Rs.

3.3 Regulation by pro- and anti-apoptotic proteins

It is increasingly recognized that a number of the key pro and anti-apoptotic Bcl-2 family of proteins in the cell are present in ER membranes and exert important functional effects from this subcellular location (reviewed in Refs. [23]). In partic-

ular, the anti-apoptotic proteins Bcl-2 / Bcl-xL and the pro-apoptotic proteins Bax/Bak have been found to alter the Ca²⁺ permeability of ER membranes [64–66]. Anti-apoptotic proteins can potentially attenuate the pro-apoptotic role of IP₃Rs in two distinct ways. Firstly, overexpression of Bcl-xL causes a dramatic decrease in IP₃R expression in a number of T-cell lines [67]. However, this has not been a general finding in cells over-expressing Bcl-2 anti-apoptotic proteins [35, 43, 68]. Secondly, the extent of agonist-mediated Ca²⁺ release from the ER is diminished when anti-apoptotic proteins are over-expressed. Although this is a fairly consistent observation in the literature [42, 43, 68–70] the underlying mechanism remains controversial. Several lines of evidence support the view that the anti-apoptotic proteins function to inhibit IP₃R channel activity. This has been documented by measuring the response of permeabilized cells over-expressing Bcl-2 to added IP₃ [43, 68, 70], or intact cells to a permeant form of IP₃ [68], or by monitoring IP₃R channel activity in planar lipid bilayers in response to recombinant Bcl-2 [68]. None of these studies found any changes in the content of ER Ca²⁺ stores induced by Bcl-2 overexpression. However, other studies have shown that Bcl-2 expression can induce a leak of Ca²⁺ from ER stores that would be expected to diminish the amount of Ca²⁺ available for release by apoptotic stimuli (reviewed in Ref. [64]). Recently White et al. [35] have reported that Bcl-xL activates IP₃Rs by allowing channels to remain open even at the very low IP₃ concentrations expected to be present under unstimulated conditions. They have made the interesting proposal that the enhanced leak of ER Ca²⁺ promoted by anti-apoptotic proteins, under some conditions, may just be a manifestation of an activation of IP₃R channels by endogenous levels of IP₃. This was supported by the finding that stable over-expression of Bcl-xL induced a decrease in the ER Ca²⁺ store in wild-type but not in the TKO IP₃R-deficient DT-40 cells [35].

At present the question of whether the Bcl-2 family of anti-apoptotic proteins inhibits or activates the IP₃R channel awaits further clarification. Some of the discrepant data may be related to the different type of complexes that Bcl-2/Bcl-xL could potentially form with IP₃Rs. Both proteins have been shown to bind to all three IP₃R isoforms using co-immunoprecipitation and fusion protein pull-down assays [35, 68, 71]. The site of interaction with IP₃Rs is in the C-terminal tail and has been further localized to a region in the proximal portion of a coiled-coil domain that may be critical for channel assembly and function (Kevin Foskett, personal communication) ([72]; c.f. Fig. 2(B)). The interactions with this site have been presumed to be direct but a recent report argues that only the type-I IP₃R isoform can associate with Bcl-2 and that this association is indirect, involving a ternary complex with protein phosphatase-1α (PP-1α) [43]. PP-1α associates with a site on the C-terminal tail and inhibits IP₃R channel activity [73] (Fig. 2(B)). The

binding of Bcl-2 to PP1 α has also been observed in other experimental systems [74]. Another protein phosphatase, calcineurin, is also reported to form a ternary complex with Bcl-2 and IP₃Rs [75]. This raises the issue that the modulation of IP₃R function by anti-apoptotic proteins could be related to alterations in the phosphorylation state of the IP₃R and that the direct versus indirect modes of association of anti-apoptotic proteins with the receptor could have different functional effects on the IP₃R Ca²⁺ channel. In addition, the assumption that Bcl-2 and Bcl-xL function in exactly the same manner may not be warranted [76].

IP₃ receptors are phosphoproteins and have long been recognized as excellent substrates for A and G- kinases [3, 77, 78]. The major phosphorylation sites for A-kinase on the type-I IP₃R are serines 1589 and 1755 and the phosphorylation in most cases produces a potentiation of IP₃-mediated Ca²⁺ release [79, 80]. A notable exception is vascular smooth muscle where phosphorylation by G-kinase inhibits IP₃R-mediated Ca²⁺ release and promotes relaxation [81]. IP₃Rs form macromolecular complexes with both A-kinase, G-kinase and protein phosphatases [81–83]. The role of the pro-apoptotic proteins Bax and Bak in regulating ER Ca²⁺ dynamics have been studied in MEFs isolated from Bax(–/–)/Bak(–/–) mice [84]. These cells had a lower steady-state Ca²⁺ content in the ER, an enhanced basal leak of Ca²⁺ and were resistant to Ca²⁺ dependent apoptotic stimuli. These effects were attributed to the presence of unrestrained Bcl-2 in ER membranes. It was also noted that the type-I IP₃R in the Bax/Bak deficient cells showed increased phosphorylation at serine 1755 [71]. siRNA suppression of Bcl-2 or the type-I IP₃R (but not the type-III IP₃R) reduced the basal Ca²⁺ leak and partially restored the ER Ca²⁺ content of the Bax/Bak deficient cells. From this data it was suggested that the intrinsic leak of Ca²⁺ from the ER is related to the degree of phosphorylation of the IP₃R, with “hyperphosphorylation” of the receptor generating a leaky channel [71]. The exact mechanism by which unrestrained Bcl-2 activity may bring about enhanced phosphorylation of the type-I IP₃R was not defined but the ability of Bcl-2 to recruit calcineurin to the IP₃R was considered a possibility. However, enhanced calcineurin association would be expected to decrease (not increase) the phosphorylation state of the protein. Indeed, IP₃Rs have a reduced phosphorylation state in a T-cell line overexpressing Bcl-2 [85]. There is also no evidence that increasing the phosphorylation state of the receptor makes IP₃R channels intrinsically leaky. In contrast, there is clear evidence that increased phosphorylation at the A-kinase sites can markedly increase the IP₃ sensitivity of the channel [79, 80]. Therefore, the increased Ca²⁺ “leak” seen in the ER of Bax/Bak deficient cells could reflect an enhanced responsiveness of IP₃Rs to ambient levels of IP₃.

Although Bax and Bak are partly localized to ER membranes, there have been no reports that they directly interact with IP₃Rs. Therefore, their effects on ER Ca²⁺ handling are assumed to be entirely due to modulating the levels of IP₃R-associated anti-apoptotic proteins. A mutant form of Bcl-2 which is unable to bind pro-apoptotic proteins does not reduce the levels of ER Ca²⁺ stores [86]. Interestingly, a mutant form of Bcl-x1 with impaired binding to Bax or Bak, but normal binding to BH3-only Bcl-2 family of proteins, continued to decrease ER Ca²⁺ stores [86]. It is unclear if any of these Bcl-2/Bcl-xL mutants have an altered ability to interact with IP₃Rs. The role of the BH3-only family of proteins in regulating ER Ca²⁺ dynamics has not been extensively explored. The addition of *t*-Bid had no effects by itself on IP₃R channel activity but did reverse the stimulatory effects of Bcl-xL [35]. The ER localized BH3-only protein Bik can also induce an ER Ca²⁺ leak and the effect requires the presence of Bax or Bak [87]. The Bcl-2 associated protein-31 (Bap31) is a polytopic integral membrane protein located in the ER which can be cleaved during apoptosis by caspase-8. The cleaved product p21 enhances ER Ca²⁺ leak and triggers apoptosis [88]. Whether any of these pathways of ER Ca²⁺ depletion also involves IP₃Rs or reflect separate mechanisms remains to be established.

An increased prevalence of neural stem cells has been noted in the adult brains of mice deficient in either Bax and/or Bak [89, 90]. Recently, it was reported that the stem cells in culture from Bax-deficient mice are more resistant to Ca²⁺-dependent apoptosis and have a lower ER Ca²⁺ content than those from wild-type mice. siRNA knockdown of the type-I IP₃R in wild-type neural precursor cells mimics the phenotype of the precursor cells from Bax deficient mice [89]. Further studies of the interactions of IP₃Rs with anti-apoptotic proteins and molecular identification of the pathways that constitute an ER Ca²⁺ leak should provide important advances in our understanding of the changes occurring in ER Ca²⁺ handling during apoptosis.

3.4 Regulation by Akt-kinase

In addition to the phosphorylation by A-kinase and G-kinase discussed above, the IP₃R is also a substrate for several other protein kinases including calmodulin dependent protein kinase II [91], protein kinase-C [92], cdc2/cyclin B1 [93] and MAP kinases [94]. From the context of apoptosis, a particularly relevant observation is that the IP₃R is a substrate for Akt kinase [95]. This enzyme has a key role in promoting cell survival utilizing many independent mechanisms. These include regulating the activity of fork-head and NF- κ B transcription factors and inactivation of the pro-apoptotic protein Bad (reviewed in Refs. [96, 97]). The phosphorylation of the type-I IP₃R occurs in the C-terminal tail at a single consensus Akt phosphorylation site (serine 2618) that is present in all

three IP₃R isoforms (Fig. 2(B)). IP₃R phosphorylation was demonstrated *in vitro* and *in vivo* in response to insulin activation of the PI-3 kinase pathway [94]. Some transformed cells which have a constitutively active Akt kinase, such as the prostate carcinoma cell line LnCaP, also have basally phosphorylated IP₃Rs which can be suppressed by the PI-3 kinase inhibitor LY294002 [95]. Only a small proportion of the total cellular pool of IP₃Rs appears to be a substrate for Akt kinase. This may be related to the fact that Akt kinase is normally activated in the plasma membrane and that subsequent translocation to the ER surface is limited and/or that the phosphorylation site in the C-terminal tail in the native protein is not readily accessible to the kinase. Potentially, changes in the phosphorylation state of a small pool of IP₃Rs at a critical location could still exert large effects on the initiation of the apoptotic cascade. The functional consequences of the phosphorylation have been studied by making phosphomimic mutants of the Akt phosphorylation site and examining their flux properties when expressed in TKO DT-40 cells or in COS cell microsomes. The data indicate that phosphorylation does not alter the flux properties of the channel [95]. A number of studies have shown that a requirement for IP₃Rs in obtaining optimal rates of apoptosis can be observed by monitoring the levels of caspase-3 activation. Thus, TKO DT-40 cells [20, 98] and T-cells deficient in the type-I IP₃R [99] have low rates of caspase-3 activation in response to an apoptotic stimulus when compared to the wild-type cells. In TKO DT-40 cells a non-phosphorylatable S2618A mutant was able to support almost 3.5-fold higher caspase-3 activation than the phosphomimic S2618E which behaves similarly to wild-type receptors [95]. This suggests that there is some protective role of Akt phosphorylation of IP₃Rs that is independent of any inhibition of Ca²⁺ channel activity. Possibly, this could involve altered interactions of other regulatory proteins with the IP₃R such as those occurring in the C-terminal tail (Fig. 2(B)) but the nature of these alterations remain to be identified.

4 IP₃Rs as a macromolecular signaling complex at ER membranes

At the time of writing 55 proteins are reported as associating with IP₃Rs based on co-immunoprecipitation, yeast two-hybrid and fusion protein studies (see Ref. [3] for a partial list). The most recently identified interaction is with a novel protein called DANGER, which in contrast to cytochrome *c*, enhances the inhibitory effect of Ca²⁺ on IP₃-mediated Ca²⁺ release [100]. The multiplicity of associations displayed by the IP₃R has led several authors to propose that the receptor functions as a macromolecular signaling complex on the ER membrane [3, 101]. A role for IP₃Rs independent of their ability to mobilize intracellular stores has been observed

previously. TKO DT-40 cells lack both agonist-mediated intracellular Ca²⁺ release and Ca²⁺ entry across the plasma membrane [102]. However, agonist-mediated Ca²⁺ entry can be restored by transfection with a functionally inactive IP₃R containing a point mutation (D2550A) in the selectivity filter of the ion channel pore, but not by IP₃R mutants defective in IP₃ binding [102]. Figures 2(D) and (E) show the effect of STS on caspase-3 activation and cell viability of stable cell lines containing the D2550A mutant. Surprisingly, this mutant functions to support STS-induced apoptosis. By contrast, another functionally inactive mutant in which the C-terminal tail was deleted [72] was unable to support apoptosis in DT-40 cells. Assefa et al. also showed that a functionally inactive mutant containing a 225 amino-acid deletion from the N-terminus was able to support caspase-3 activation in DT-40 cells [20]. The interpretation of this result was that the N-terminal deletion mutant could still be proteolytically cleaved by caspase-3 to generate the 95 kDa “channel-only” fragment, and would provide the ER Ca²⁺ leak necessary to initiate the apoptotic cascade. In our hands the “channel-only” domain containing the pore-inactivating D2550A mutant was also able to support STS-induced apoptosis in DT-40 cells (Khan & Joseph, data not shown).

It can be anticipated that multiple and redundant pathways to apoptosis exist in a cell. Functional IP₃Rs may provide one of several pathways of Ca²⁺ delivery to the mitochondria. A correlation between enhanced IP₃ sensitivity of IP₃Rs and STS sensitivity to apoptosis has indeed been noted in DT-40 cell lines [98]. However, the results also suggest that IP₃Rs can serve functions in apoptosis that go beyond simply supplying Ca²⁺ to sensitize the mitochondria to apoptotic insults. In the absence of functional IP₃Rs, this role could be subserved by Ca²⁺ entry across the plasma membrane. There is ample evidence for microdomains in which plasma membrane Ca²⁺ entry and mitochondrial Ca²⁺ uptake are coupled (reviewed in Ref. [103]). Recent studies suggest that a few IP₃Rs in the plasma membrane can themselves contribute substantially to Ca²⁺ entry in lymphocytes, but this activity requires a functional pore [104]. While the ion-channel function of IP₃Rs appears dispensable for apoptosis when plasma membrane Ca²⁺ entry mechanisms are operative, the C-terminal tail of the receptor still appears to be essential (Fig. 2(C) and (D)). This could reflect the necessity for interaction with one or more of the known proteins targeting this domain (Fig. 2(B)). It cannot presently be excluded that there are additional interactors remaining to be identified. As pointed out above, cell-permeant peptides encoding the IP₃R cytochrome *c* binding site can inhibit cell death [57]. Further mapping studies with additional C-terminal deletion mutants, a search for additional proteins targeting the C-terminal tail and investigation of the relative accessibility of this region of the native receptor should all provide insights into the ion-channel independent role of IP₃Rs.

5 Diseases linked to aberrant apoptotic signaling by IP₃Rs

Recent studies of several diseases linked to defects in apoptosis and Ca²⁺ signaling have converged on examining IP₃Rs as a possible molecular target. Four examples are briefly discussed. They are the HIV-1 associated Nef protein, presenilins, the polyglutamine repeat protein huntingtin, and the NF-κB signaling pathway.

Nef is an accessory protein encoded by the HIV-1 & 2 virus which functions to maintain infected T-cells in an activated state and to induce apoptosis in non-infected cells of the immune system [105]. Manninen and Saksela [106] found that Jurkat cells transfected with Nef showed a robust activation of NFAT which was dependent on Ca²⁺ entry across the plasma membrane. The authors also found that NFAT activation was attenuated in Jurkat cells deficient in the type-I IP₃R, that Nef and the IP₃R were co-immunoprecipitated, and that Nef did not modify the content of IP₃-sensitive stores [106]. It was concluded that Nef did not affect the channel function of IP₃Rs but altered its coupling to the plasma membrane Ca²⁺ entry mechanism(s) such that Ca²⁺ entry occurred independently of store depletion. Interestingly, another HIV encoded protein, HIV-Tat, whose neurotoxic effects are thought to contribute to AIDS related dementia also promotes apoptosis [107]. In this case the mechanism appears to involve a rapid mobilization of IP₃-sensitive stores resulting from an increased generation of IP₃ [108].

Presenilin-1 (PS-1) is a predominantly ER localized protein involved in the proteolysis of amyloid precursor protein. Mutations in PS-1 are associated with the majority of cases of early onset familial Alzheimer's disease (FAD). There have been a number of studies showing that cultured cells or primary neurons expressing mutant PS-1 display an enhanced response to caged-IP₃ with no changes in the expression level of IP₃Rs [109, 110]. This has led to the hypothesis that mutant PS-1 can cause an over-filling of IP₃-sensitive Ca²⁺ stores by an unidentified mechanism, and that an enhanced Ca²⁺ release may underlie a number of the pathological effects associated with Alzheimers disease [109, 110]. The expression of ryanodine receptors is increased in PS-1 expressing transgenic mouse models [110]. Recently, the mechanism of enhanced IP₃-mediated Ca²⁺ release has been suggested to be indirect, and to involve recruitment of ryanodine receptor channels activated by calcium-induced Ca²⁺ release [111]. Another possible explanation offered by Tu et al. is based on their observations that the wild-type (but not the mutant) presenilins can form low conductance, non-selective ion channels in planar lipid bilayers [112]. They suggest that wild-type PS may contribute to the basal ER Ca²⁺ leak and that the absence of this leak with mutant PS may cause the observed overfilling of Ca²⁺ stores. Although, this would be in agreement with the bulk of the experimental data, it should

also be pointed out that data inconsistent with an enhanced ER Ca²⁺ leak mediated by PS-1 has been reported by Kasri et al. [113].

Presenilin-2 (PS-2) is closely related to PS-1 (67% identical) and has similar effects on the over-filling of ER Ca²⁺ stores [114]. Both normal and mutant PS-2 are pro-apoptotic proteins [115]. Proteolytic processing of PS-2 by caspase-3 generates a short 22 amino-acid peptide from the C-terminal fragment that has been shown to induce apoptosis when introduced into a prostate adenocarcinoma cell line [116]. This peptide depleted intracellular Ca²⁺ stores, co-immunoprecipitated with the type-I IP₃R, and increased IP₃R channel activity. The site of interaction with the IP₃R has not been identified. The corresponding peptide from PS-1 was inactive. Thus PS-1 and PS-2 appear to possess both similar and different modalities for augmenting IP₃-mediated Ca²⁺ release.

Huntington's disease is caused by a polyglutamine expansion in the N-terminal region of huntingtin (Htt^{exp}), a 350 kDa cytosolic protein. Both wild type Htt and the mutant Htt^{exp} can associate with the type-I IP₃R by direct interactions, as well as the formation of a ternary complex involving the Huntington associated protein HAP1A [117]. A high-affinity binding site was localized to the extreme C-terminal region of the IP₃R [117] (Fig. 2(B)). The addition of Htt^{exp}, but not Htt, caused an activation of IP₃R channel activity measured with suboptimal IP₃ concentrations in planar lipid bilayers [117, 118]. Repetitive glutamate stimulation of medium spiny neurons derived from a mouse model of Huntington's disease caused a persistent elevation of cytosolic Ca²⁺ that was associated with release of cytochrome *c* and apoptosis [119]. These changes were absent in the neurons derived from control mice. These studies have led to a model in which the activation of IP₃R by Htt^{exp}, or its proteolytically derived fragments, contributes to enhanced apoptosis and neuronal cell loss [119].

NF-κB is a transcription factor whose function has been implicated in a wide range of diseases including immune disorders, cancer and inflammatory conditions [120, 121]. NF-κB signaling functions to favor survival by regulating the expression of key proteins such as Bcl-2, inhibitor of apoptosis proteins and manganese superoxide dismutase [122]. Mobilization of intracellular Ca²⁺ stores can activate NF-κB by indirect mechanisms that are not fully understood. For example Glazner et al. [123] reported that depletion of ER stores with IP₃ or thapsigargin caused the release of a protein factor that could activate NF-κB when added to cytosol extracts. More recently, it has been shown that NF-κB activation can decrease type-I IP₃R expression probably through NF-κB-sensitive promoter elements present in the type-I IP₃R gene [123]. Thus a feed-back loop exists in which Ca²⁺ can activate NF-κB which in turn can suppress the Ca²⁺ signal by reducing IP₃R expression. The reduction of IP₃R expression

is believed to be one mechanism that would contribute to the pro-survival function of NF- κ B [123].

6 Mechanisms underlying a pro-apoptotic shift in the IP₃R-mitochondrial Ca²⁺ transfer

In the 1990s, a breakthrough discovery revealed that permeabilization of the outer mitochondrial membrane (OMM) and the ensuing release of cytochrome *c* and other intermembrane space factors to the cytosol represents the commitment steps in a variety of apoptotic paradigms [124, 125]. The possible role of the IP₃-induced mitochondrial matrix [Ca²⁺]_m ([Ca²⁺]_m) signal in the control of apoptosis has become a subject of immense interest since uptake of Ca²⁺ by mitochondria is one of the most effective triggers of the mitochondrial permeability transition [126]. First, IP₃R-mediated [Ca²⁺]_m spikes were shown to cooperate with some apoptotic stimuli to induce mitochondrial membrane permeabilization and apoptosis [127]. Subsequently, apoptotic stimuli were shown to target IP₃R-mediated Ca²⁺ mobilization, evoking a [Ca²⁺]_m signal that leads to mitochondrial membrane permeabilization [128]. These two mechanisms i.e. the coincidence of IP₃R-mediated [Ca²⁺]_m signal with some forms of mitochondrial stress and the augmentation and prolongation of IP₃R-mediated Ca²⁺ transfer to the mitochondria seem to represent the main pathways for inducing cell death by the IP₃R-linked [Ca²⁺]_m signal.

Upon exposure of mitochondria to trains of Ca²⁺ pulses (micromolar or larger), Ca²⁺ uptake triggers opening of the permeability transition pore (PTP) and in turn, permeabilization of both the outer and inner mitochondrial membranes [126]. Permeation of solutes (with molecular masses below approximately 1500 Da) across the PTP may cause mitochondrial swelling and rupture of the OMM by itself. However, in several Ca²⁺-induced apoptosis paradigms the release of intermembrane space proteins to the cytosol occurs in the absence of large scale swelling, indicating that the PTP also recruits alternative mechanisms of OMM permeabilization. The Ca²⁺-induced PTP opening can be promoted by increasing either the size of the individual pulses or the frequency of the pulsing, illustrating possible mechanisms for a pro-death shift during [Ca²⁺]_c oscillations [129]. Furthermore, a prolonged, submicromolar Ca²⁺ increase may evoke gradual sensitization of the mitochondrial Ca²⁺ uptake sites resulting in a slower and more sustained stimulation of mitochondrial Ca²⁺ accumulation [130, 131]. Over a long time, this may lead to mitochondrial Ca²⁺ overloading and PTP-dependent permeabilization of mitochondria. Therefore, an enhancement of the ER-mitochondrial transfer may cause a pro-death shift in calcium signaling both during IP₃-induced [Ca²⁺]_c spikes/oscillations or as a result of a slow and persistent leak from the ER. Although the PTP

opening has been considered as the central mechanism for the [Ca²⁺]_m-dependent mitochondrial membrane permeabilization and cell death, the diversity of mitochondrial Ca²⁺ effectors (reviewed in Refs. [9, 132]) suggests the possible involvement of additional mechanisms. The factors determining the extent of the Ca²⁺ transfer from IP₃Rs to the mitochondria are discussed below.

6.1 Ca²⁺ storage in the ER

During [Ca²⁺]_c oscillations, mitochondrial Ca²⁺ uptake is closely coupled to the IP₃R-dependent Ca²⁺ release that is driven by the [Ca²⁺]_c gradient between the ER lumen and the cytosol. Studies evaluating the connection between the ER Ca²⁺ storage and cell survival have reported that fluctuations of ER luminal [Ca²⁺]_{ER} ([Ca²⁺]_{ER}) in a narrow range (approx. 30%) cause massive changes in apoptosis. A clue to the mechanism of this effect is that in cells where the apoptotic agents or Bcl-2 family proteins induced moderate differences in the [Ca²⁺]_{ER} and in the IP₃-linked [Ca²⁺]_c rise, relatively large changes occurred in the [Ca²⁺]_m signal. For example, Scorrano et al. reported that in the Bax/Bak knockout cells the IP₃-linked [Ca²⁺]_c signal was partially inhibited but the [Ca²⁺]_m signal was essentially abolished [84]. Quantitative analysis of ER loading and IP₃-induced mitochondrial Ca²⁺ transport indicates a supralinear relationship between these parameters (Csordás and Hajnóczky, unpublished). This complex relationship seems to reflect the sensitization of the mitochondrial Ca²⁺ uptake sites exposed to the high [Ca²⁺]_c microdomains close to the IP₃Rs. Patches of calreticulin, a high capacity Ca²⁺ binding protein in the ER lumen and IP₃Rs were shown in close association with mitochondria along cell processes in cultured rat oligodendrocytes, illustrating the morphological basis for a local interaction between ER Ca²⁺ storage, IP₃R-mediated Ca²⁺ release and mitochondrial Ca²⁺ uptake [133]. Local ER Ca²⁺ filling also seems to be supported by concentration of SERCA pumps in proximity to these sites [134]. Interestingly, overexpression of calreticulin in HEK-293 cells led to an increase in ER Ca²⁺ storage and in the IP₃-linked [Ca²⁺]_c signal but caused a decrease in the concomitant [Ca²⁺]_m signal [135]. However, the calreticulin-overexpressing cells also showed low $\Delta\Psi_m$, indicating that a decrease in the driving force was responsible for the attenuated mitochondrial Ca²⁺ uptake. Overexpression of the Ca²⁺ binding C domain of calreticulin was sufficient to reproduce the effect of the full length calreticulin. It was suggested that a prolonged increase in Ca²⁺ turnover between ER and mitochondria evoked mitochondrial injury and in turn, promoted apoptosis in the calreticulin-overexpressing cells [135]. Thus, an increase in the ER Ca²⁺ storage may support mitochondrial membrane permeabilization by both rapidly and slowly acting mechanisms.

6.2 IP₃R-dependent Ca²⁺ permeation across the ER membrane

During stimulation with varying concentrations of IP₃-linked agonists or IP₃, the [Ca²⁺]_c transients are usually associated with a [Ca²⁺]_m rise that is in proportion to the magnitude of the [Ca²⁺]_c signal. However, quantitation of the [Ca²⁺]_c and [Ca²⁺]_m responses has revealed a rightward shifted IP₃ dose-response for the [Ca²⁺]_m signal [136]. Furthermore, the [Ca²⁺]_m signal was disproportionately suppressed by overexpression of IP₃-binding proteins that serve as IP₃ buffers [137] and was also small when complete mobilization of the ER Ca²⁺ store was achieved slowly by a gradual increase in [IP₃] [136]. These data indicated that optimal activation of the rapid Ca²⁺ transfer to the mitochondria is dependent on concurrent activation of the IP₃Rs during Ca²⁺ mobilization. Therefore, any factors that alter the IP₃/Ca²⁺ sensitivity, Ca²⁺ permeability, or the number of available IP₃Rs are expected to affect the efficacy of the [Ca²⁺]_c signal propagation to the mitochondria. However, among the numerous factors that have been described to interact with the IP₃Rs (see Section 4), only Bcl-2/Bcl-xL has been evaluated for modulation of the IP₃-linked [Ca²⁺]_m signal or the mitochondrial calcium content. Both overexpression of Bcl-2 [69] and knock-out of Bax and Bak, which results in a relative abundance of Bcl-2/Bcl-xL [84] have been reported to cause suppression of both the [Ca²⁺]_c and [Ca²⁺]_m signal evoked by IP₃-linked agonists. In these cases, attenuation of Ca²⁺ release through the IP₃R accounts for the suboptimal Ca²⁺ delivery to the mitochondria. The decrease in the Ca²⁺ release may reflect either a Bcl-2-mediated inhibition of the IP₃R Ca²⁺ channel [68] or reduction in the ER Ca²⁺ load [69]. Notably, in the study that described Bcl-xL-dependent sensitization of the IP₃R to IP₃ [35], the IP₃R-mediated Ca²⁺ leakage from the ER was proposed to support a permanent Ca²⁺ transfer to the mitochondria, stimulating oxidative metabolism. Based on the amount of the FCCP-releasable Ca²⁺, an increase in mitochondrial Ca²⁺ storage has also been reported in unstimulated Bcl-2-overexpressing cells [138]. Generally, prolonged mitochondrial Ca²⁺ storage would enhance the vulnerability of the mitochondria but the mitochondria localized fraction of Bcl-2/Bcl-xL in the overexpression model exerts protection against membrane permeabilization.

As described earlier (Section 2), the siRNA-mediated depletion of the type-III IP₃R isoform had a more marked effect on the [Ca²⁺]_m signal and bile-acid or STS-induced apoptosis in CHO cells than depletion of the other isoforms [28]. These results suggested that type-III IP₃Rs preferentially transmit Ca²⁺ to the mitochondria at least in some cell types and that the maximal strength of the IP₃R-dependent [Ca²⁺]_m signaling is needed for the apoptotic cascade to prevail. Localization studies indicated that the type-III isoform may be preferentially localized adjacent to mitochondria

[28]. Hence, another potentially important factor dictating the magnitude of the [Ca²⁺]_m signal may be the amount and spatial localization of individual IP₃R isoforms.

6.3 Strategic positioning of mitochondria close to the sites of IP₃R-mediated Ca²⁺ release

Mitochondria are distributed at various subcellular locations and show functional heterogeneity but in most paradigms, at least a fraction of mitochondria rapidly respond to IP₃-mediated Ca²⁺ mobilization [139–141]. The bulk [Ca²⁺]_c rise seems to be too low to activate the uniporter [142, 143], although a “rapid mode” of mitochondrial Ca²⁺ uptake has been claimed to respond to submicromolar [Ca²⁺]_c increases [144]. A compelling piece of evidence for the role of a local IP₃R-mitochondrial Ca²⁺ transfer in the rapid mitochondrial Ca²⁺ uptake is the observation that buffering of the global [Ca²⁺]_c signal by Ca²⁺ chelators (EGTA or BAPTA 200 μM) fails to abolish the IP₃-induced [Ca²⁺]_m signal [136, 145]. While the bulk [Ca²⁺]_c rises to approx. 1 μM, the [Ca²⁺]_c at a 100 nm distance from an open IP₃R channel is estimated to reach ≥10 μM and in the immediate vicinity of the IP₃R (10 nm distance) would attain ≥100 μM [146]. Since the mitochondrial Ca²⁺ uptake is greatly facilitated in the 10 μM [Ca²⁺]_c range [142] the uniporter-mediated Ca²⁺ current would be expected to steeply increase in the micromolar to millimolar [Ca²⁺]_c range [147]. In addition, the distance between the IP₃R and the uniporter is expected to affect the relay of the IP₃-dependent [Ca²⁺]_c signal to the mitochondria. When bifunctional linkers (5 nm in length) that anchor ER to the OMM were transiently overexpressed in RBL-2H3 cells in order to narrow the ER-mitochondrial gap, an enhancement of the IP₃R-mitochondrial Ca²⁺ transfer and an increased sensitivity to ER stress-induced apoptosis was observed [148].

In light of the above data, a change in mitochondrial distribution might be expected to cause a change in the IP₃R-mediated Ca²⁺ transfer to the mitochondria. However, the movement of mitochondria to the perinuclear region, induced by overexpression of proteins that control mitochondrial morphology and anchorage did not seem to affect the IP₃-linked [Ca²⁺]_m signal [149, 150]. In this case, mitochondria were clearly separated from the peripheral domains of the ER. However, mitochondria could form interactions with the copious perinuclear ER to sustain the IP₃-dependent [Ca²⁺]_m signal. Also, formation of a physical linkage between ER and mitochondria (see below) could still hold some ER domains close even during repositioning of the mitochondria.

6.4 Physical linkage between ER and mitochondria

The existence of anchorage sites between ER and mitochondria have been suggested based on co-sedimentation of ER

particles with mitochondria and electron microscopic observations of close associations between mitochondria and ER vesicles [151, 152]. Recent studies of the ER-mitochondrial interface by electron tomography and by limited proteolysis provided evidence for direct protein linkage (tethers) between both rough and smooth ER and mitochondria. The tethers range in size from 6 to 15 nm at the smooth ER, and 19 to 30 nm at the rough ER [148]. The variations in the tether length indicate diversity of the protein components that may be involved in the linkage. The specific tether-forming proteins are unknown. Tethers originated at the rough ER appear close to the ribosomes, raising the possible role of proteins that leave the ER to enter the mitochondria. In addition, several ER or mitochondrial membrane or membrane-associated proteins have been considered. Participation of the IP₃R in the formation of the links would localize the Ca²⁺ release site to the anchorage points of the ER-mitochondrial interface. Indeed, the IP₃R seems to act as an anchor for >50 proteins (see Section 4) and show high density at the ER-mitochondrial associations [133, 153]. Formation of a possible ER-mitochondrial “Ca²⁺ tunnel” has been supported by data on a mitochondria-associated macromolecular complex that would include IP₃R and VDAC [154]. Furthermore, overexpression of VDAC has been shown to shorten the coupling time between IP₃R-mediated [Ca²⁺]_c rise and [Ca²⁺]_m elevation and led to an increased susceptibility to a Ca²⁺-dependent form of apoptosis [155]. The IP₃R domains that may participate in ER-mitochondrial linkages are unknown. Szabadkai et al. [154] have described an unexpected stimulatory effect of the ligand binding domain of the IP₃R (LBD_{IP₃R}) on the IP₃-linked [Ca²⁺]_m rise in HeLa cells when overexpressed in the cytoplasm or membrane-anchored either to the ER or OMM. It remains to be seen whether these constructs enhance the physical coupling between ER and mitochondria. Also, the cytosolic LBD_{IP₃R} caused suppression of the IP₃-dependent [Ca²⁺]_c and [Ca²⁺]_m signals in COS-7 cells, which seems to reflect the IP₃-buffering effect of the construct [137]. If the LBD_{IP₃R} increases both IP₃ buffering and the ER-mitochondrial coupling, depending on the strength of each effect, the [Ca²⁺]_m signal may be either increased or suppressed. Ryanodine receptors (RyRs) also provide for rapid Ca²⁺ transfer to the mitochondria, supporting the possibility that an IP₃R-like domain for interaction with the mitochondria may also be present in the RyR. The IP₃R and RyRs share the highest sequence homology within the N-terminal and transmembrane regions [156]. Conservation in the transmembrane region may be due to their common conductance of Ca²⁺ ions, whereas the homology within the N terminus is unclear since RyR lacks the ability to bind IP₃. Structural analysis and structure-based sequence analysis of IP₃R and RyRs suggest that the N-terminal portion of RyRs is comprised of two β-trefoil domains in a similar manner as IP₃R [157]. IP₃R knockouts have also been

used to determine the role of IP₃R in the coupling of ER to the mitochondria. Electron microscopy and tomography studies have shown the presence of tight ER-mitochondrial associations and tethers in both IP₃R-expressing and IP₃R-deficient TKO DT40 cells [148]. Thus, an IP₃R- and VDAC-containing complex may support both the functional and structural aspects of the ER-mitochondrial interface but it has no exclusive role in formation of the linker.

In the past years, several ER- or mitochondria-resident proteins have been shown to be important for maintaining the spatial relationship between ER and mitochondria and hence, have also been implicated as possible ER-mitochondrial linking elements. These proteins include dynamin-related protein (DLP-1/DRP1-1; [158, 159]), tumor autocrine motility factor receptor (AMF-R; [160]) and PACS-2 (phosphofurin acidic cluster sorting protein-2) and BAP31 (B cell antigen receptor-associated protein 31) [161]. Overexpression of DLP-1/DRP-1 that mediates OMM fission has been reported to cause attenuation of the IP₃R-mediated [Ca²⁺]_m signal evoked by suboptimal stimulation. Attenuation of the [Ca²⁺]_m signal was also associated with suppression of ceramide-induced apoptosis, whereas other apoptotic mechanisms were augmented [162]. However, in this case mitochondrial fragmentation seems to hinder the spreading of the [Ca²⁺]_m rise among the mitochondria rather than the Ca²⁺ transfer between ER and mitochondria [162]. Immunofluorescence studies of the smooth ER resident AMF-R revealed a domain of the ER that shows Ca²⁺ dependent associations with the mitochondria, illuminating a possible mechanism for the recruitment of mitochondrial Ca²⁺ uptake to the sites of Ca²⁺ release [160]. Further investigation is needed however, to determine the significance of this mechanism for the Ca²⁺ transfer to the mitochondria and for the control of apoptosis. Similar to the DLP-1/DRP-1 overexpression, depletion of the ER/cytoplasmic sorting protein PACS-2 has been reported to cause mitochondrial fragmentation and localization in the perinuclear area and suppression of apoptosis [161]. The PACS-2-depletion induced mitochondrial fragmentation required the cleavage of BAP31 by caspase-8 in the ER to yield p20. In turn, p20 would cause ER Ca²⁺ release and ER-mitochondrial Ca²⁺ transfer to facilitate DLP1/DRP1-mediated mitochondrial fission [23]. A similar mechanism has also been reported for another ER-linked protein, the BH3-only Bik [163]. Thus several ER factors converge on the DLP-1/DRP-1-mediated mitochondrial fission to evoke a change in the relative distribution of mitochondria and ER and to impact on cell survival. The change in mitochondrial distribution is expected to yield some rearrangements of the ER-mitochondrial interface, but the area of the ER-mitochondrial interface does not necessarily decrease and its organization does not necessarily change. At least, Ca²⁺ transfer at the ER-mitochondrial interface is

needed to activate DLP-1/DRP-1 and seems to be sustained in some situations where mitochondria have changed their distribution. Most studies of ER-mitochondrial coupling described conditions that caused increased separation of ER from mitochondria. In a recent study, a narrowing of the ER-mitochondrial gap and an increase in the area of the interface was observed in cells exposed to ER stress conditions (tunicamycin treatment and serum starvation) [148]. It has been speculated that these changes support mitochondrial Ca^{2+} uptake during slow ER Ca^{2+} release that normally would be excluded from the mitochondria, making mitochondria more prone to Ca^{2+} overloading and PTP opening. Although a change in the gross ER and mitochondrial morphology did not seem occur, the role of tethers in shortening the ER-mitochondrial gap needs further investigation. Identifying the tether forming proteins and imaging the tethers in live cells, where the ER and mitochondria also move and undergo fusion-fission, will offer clues about how the ER-mitochondrial interface is controlled.

6.5 Dynamic aspects of the ER-mitochondrial distribution

In many cell types, mitochondria are highly dynamic organelles that undergo fusion and fission and display both short and long range movements. Mitochondrial fragmentation has been observed as an obligatory early event in several apoptotic paradigms [164, 165]. Mitochondrial fragmentation may be both a cause and a consequence of a change in the IP_3R -mitochondrial calcium transfer (reviewed in Ref. [166]). Recently, mitochondrial and ER motility has also been implicated in the control of ER-mitochondrial calcium signaling. Mitochondria are commonly bound and move along microtubules utilizing kinesin and dynein motor proteins but in other instances mitochondrial association with microfilament and intermediate filament tracks have also been documented [167]. ER morphology is less dependent on the cytoskeletal structures [168] but domains of the ER are also linked to and may move along the microtubules or microfilaments. However, it has not been determined whether the mitochondrial and ER movements are coordinated with each other. The ER-mitochondrial tethers may allow ER stacks to piggy-back on moving mitochondria and vice versa. Alternatively, rapid turnover of the tether-forming structures may ensure movement of mitochondria along the ER while a permanent linkage is maintained. Pretreatment of H9c2 cells with a microtubule-disrupting agent, to inhibit the mitochondrial movements, resulted in an improvement in IP_3R induced $[\text{Ca}^{2+}]_m$ rise although the $[\text{Ca}^{2+}]_c$ increase was not enhanced [169]. Thus organellar movements may interfere with some aspects of the local ER-mitochondrial Ca^{2+} transfer. Recently, $[\text{Ca}^{2+}]_c$ elevations have been shown to cause mitochondrial motility inhibition in several cell types [169–171] and a homeostatic circuit has been proposed whereby

mitochondria are retained at the sites of the IP_3 -induced Ca^{2+} release to enhance the local ATP supply and Ca^{2+} buffering and become free to move after the decay of the $[\text{Ca}^{2+}]_c$ rise [169]. The relevance of mitochondrial motility in the context of apoptosis has not yet been explored. Nevertheless, when cells were treated with NO or H_2O_2 to induce cell death, suppression of the mitochondrial motility was observed as an early event ([172]; Saotome and Hajnóczky, unpublished). Furthermore, several studies support a link between impairment of the mitochondrial-cytoskeletal interactions and cell death [173].

7 Conclusions and future perspectives

A considerable redundancy is apparent in apoptotic pathways and this has been a complicating feature of attempts to therapeutically modify this process in the treatment of diseases such as cancer or neurodegenerative disorders. An important goal has been the identification of critical control points in apoptotic pathways that could potentially be used as drug targets. The present review has summarized the data indicating that IP_3Rs may be critical regulators of apoptosis triggered by stimuli that engage either ER and/or mitochondrial mechanisms. IP_3Rs may serve not only to funnel Ca^{2+} into the mitochondria but may have other effects on apoptosis that are independent of their ion-channel function, such as their role as adaptor proteins. Since IP_3Rs also have critical house-keeping roles in cell survival an indiscriminate targeting of these proteins would seem undesirable. The selective targeting of spatially and functionally distinct pools of IP_3Rs would be more profitable, but this would require much more knowledge of the full range of IP_3R interactions with other proteins. Another critical control factor is the spatial relationship and dynamics of ER and mitochondrial membranes. The possibility of manipulating the apoptotic process at this level awaits the challenging task of identifying the molecular machinery involved in inter-organellar dynamics.

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