

REVIEW

Executionary pathway for apoptosis: lessons from mutant mice

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ABSTRACT

Apoptosis or programmed cell death (PCD) is an evolutionarily conserved cellular process that is essential for normal development and homeostasis of multicellular organisms. Defects in the apoptosis signaling result in many diseases including autoimmune diseases and cancer. The apoptosis signaling pathway was first described genetically in the nematode *Caenorhabditis elegans* which serves as a framework for the more complex apoptotic pathways that exist in mammals. In this review, we will discuss the apoptotic pathways that are emerging in mammals as elucidated by studies of gene-targeted mutant mice.

Key words : *Apoptosis, programmed cell death, caspases, death receptors, mitochondria.*

INTRODUCTION

Apoptosis or programmed cell death (PCD) is a genetically programmed cellular event that is conserved throughout evolution. It is a specialized form of cell death that is essential for normal cellular and embryonic development and tissue homeostasis. Defects in apoptosis underlie many human pathological conditions, including cancer, autoimmune diseases and neurodegenerative disorders.

Apoptosis is characterized by specific biochemical and morphological features that

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culminate in shrinkage of the cell to apoptotic bodies that are engulfed by neighboring macrophages[1]. The process involves disruptions of mitochondrial, cytoplasmic, and nuclear integrity that result in decreased mitochondrial membrane potential, reversal of phosphatidyl serine to the outer side of the plasma membrane, degradation and “laddering” of DNA, and chromatin condensation. As such, apoptosis is distinct from cell death by necrosis or “accidental cell death”, in which stressed cells swell and burst, releasing cellular contents into the immediate microenvironment. These contents trigger an inflammatory response that can result in tissue damage.

The genetic basis of a signaling pathway leading to PCD was initially identified in the nematode *C. elegans*[2]. Since then, our understanding of the mechanisms of PCD has grown exponentially. Through the use of techniques such as the generation of gene-targeted “knockout” mice lacking specific death genes, we have been able to develop mouse models for the elucidation of mammalian PCD pathways in vivo. These mutant animals constitute powerful tools that have the potential to directly influence therapeutic strategies in humans.

Genetic foundation of apoptosis in C. elegans

Four genes in the nematode apoptotic pathway are *ced-3*, *-4*, *-9*, and *egl-1* [*(ced)* cell death abnormal; (*egl*) egg laying defective] (reviewed in[3]). *ced-3* and *ced-4* are the first pro-apoptotic genes identified[4]. CED-3 is a cysteine protease that cleaves proteins at specific aspartic acid residues[5]. CED-4 is autoactivated through oligomerization; the activated CED-4 can then activate CED-3 through proteolytic cleavage. The activities of both CED-4 and CED-3 are regulated by CED-9 and EGL-1. CED-9 interferes with the formation of the CED-4: CED-3 complex required for CED-3 activation by binding CED-4[6]. CED-9 can also bind to EGL-1[7]. Once the induction of apoptosis occurs, EGL-1 disrupts the interaction of CED-9 with CED-4 and promotes CED-3 activation[8]. This appealingly simple pathway, from EGL-1 to CED-9 to CED-4 to CED-3, supplies a framework for the much more complex web of apoptotic pathways found in mammals.

Caspases are the mammalian homologues of *ced-3* and at least 14 have been identified to date. Two putative mammalian homologues of *ced-4* exist: the apoptotic protease-activating factor-1 (Apaf-1)[9] and the Caspase Recruitment Domain4 (CARD4)/Nod-1[10],[11]. The mammalian homologues of *ced-9* are the anti-apoptotic Bcl-2, Bcl-x_L, Bcl-w, and pro-apoptotic Mcl-1 and Bax, Bak and Bok which are members of the Bcl-2 family[12]. Mammalian homologues of *egl-1* are pro-apoptotic BH3-containing members of the Bcl-2 family, including Bik, Bad, Bid, Bim and Bar[13], [14].

Caspases: the key players in apoptotic signaling in mammals

Caspases (cysteiny l aspartate-specific proteinases) are mammalian CED-3 homologues that mediate highly specific cleavage events in dying cells. Caspases are expressed as pro-enzymes containing three domains: an NH₂ terminal pro-domain, a large

subunit (~20 kD), and a small subunit (~10 kD). Activation involves proteolytic processing between domains followed by the association of the large and small subunits into an active heterodimer. The active enzymatic site of these proteases is centered around a cysteine residue contained within a conserved QACxG motif. Once activated, the caspases cleave specific aspartic acid residues of proteins involved in the maintenance of nuclear integrity, such as DNA protein kinase (DNA-PK), and poly-ADP-ribose polymerase (PARP), and proteins involved in cell cycle regulation, such as Rb and P21/WAF1. Cleavage by caspases also activates proteins that destroy cell architecture, such as gelsolin which cleaves actin filaments (reviewed in[15]). Despite many substrates that are identified, the significance and the consequence of many of the cleavage events are not well understood.

There are at least 14 caspases known to date, broadly classified into two subfamilies. The ICE subfamily of proteases, which includes caspase-1, -4, -5 and -11, is primarily involved in processing inflammatory cytokines. The CED-3 subfamily members are known for their roles primarily in apoptosis. Caspase-3 is the prototypical caspase with closest sequence homology and substrate specificity to CED-3. Members of the CED-3 subfamily are further categorized into two groups based on pro-domain characteristics. Caspases in the first group contain long prodomains which facilitate protein-protein interaction and are used to recruit other caspases. For example, caspase-2 and -9 contain the CARD in their prodomains while caspases-8 and -10 have the death effector domain (DED). Through their ability to recruit and activate other caspases, these caspases are thought to play an upstream regulatory role. Caspases in the second group, such as caspase-3, -6, and -7, have short prodomains and are thought to function as downstream “executioner” or effector caspases which cleave substrates resulting in apoptotic cell death (reviewed in[16]).

Complexity of mammalian apoptosis

In contrast to the simple linear pathway in *C. elegans*, there are at least two major pathways of apoptosis in mammals. The “extrinsic” pathway is mediated via specific death receptors (DR) expressed on the cell surface. Protein interaction modules such as the DED and others known as death domains (DDs) are used to assemble receptor signaling complex called death-inducing signaling complex (DISC). These complexes recruit and activate the upstream regulatory enzymes, caspase-8 and -10 (reviewed in[17] and[18]), which in turn activate the executioner protease caspase-3. In contrast, the “intrinsic” or mitochondrial pathway of apoptosis is initiated by events in the mitochondria. In response to apoptotic stimuli, in as yet unclarified mechanism, these organelles release cytochrome c into the cytosol which can, in the presence of dATP, associate with Apaf-1 and activate the upstream protease caspase-9. Caspase-9 in turn activates downstream caspases such as caspase-3[19], [9]. Apoptosis-inducing factor (AIF) is also released from the mitochondria upon apoptotic stimulation. AIF can directly translocate to the nucleus and generate large-scale DNA fragmentation in a

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process which appears to be caspase-independent[20], [21]. Recently, second mitochondria-derived activator of caspase (Smac) or direct IAP binding protein with low pH (DIABLO) has also been shown to be released from the mitochondria[22, 23]. It has been shown to be critical for the activation of apoptosis through binding and inhibition of inhibitors of apoptosis (IAPs) (See Fig 1).

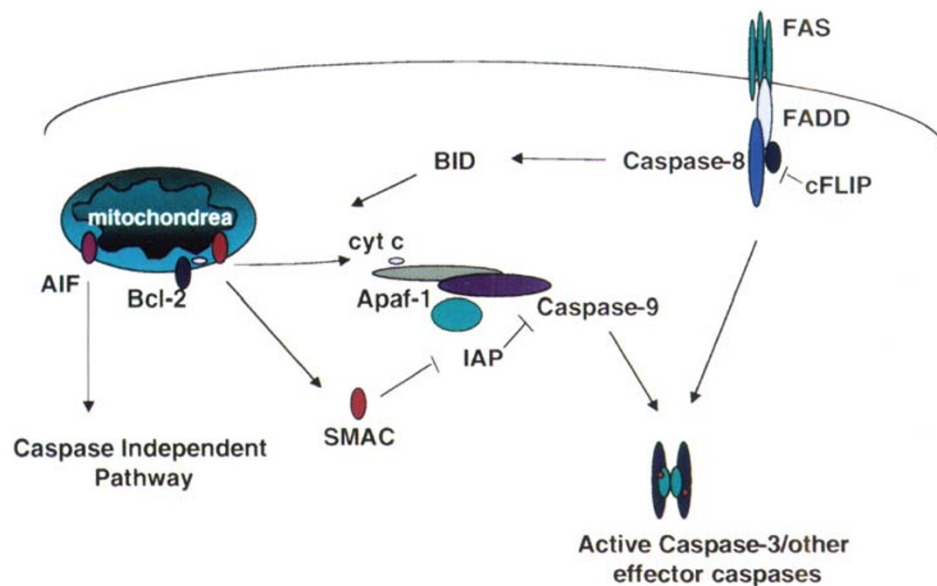


Fig 1. Multiple executionary pathway in mammals

The extrinsic pathway of mammalian PCD

Death receptors constitute a TNF receptor superfamily. The prototypical death receptor is Fas (CD95), and Fas-mediated apoptosis will serve as an illustration of the extrinsic pathway for the purposes of this review. Activation of Fas by its ligand FasL results in rapid recruitment of the DD-containing adaptor protein Fas-Associated Death Domain protein (FADD)/Mort-1[24, 25]. Recently it has been shown that these receptors are preformed trimers that undergo conformational changes upon ligand binding that allow activation signaling to occur[26]. FADD in turn recruits procaspase-8 (also called FLICE (FADD-like IL1- β converting enzyme) or MACH)[27, 28]. Fas, FADD and procaspase-8 come together at the DISC[29], the formation which allows procaspase-8 to oligomerize and activate itself by autocleavage[30]. Activated caspase-8 then cleaves other procaspases, including procaspase-3, -6, and -7[31-33].

DISC also has an inhibitory component, cellular FLICE-inhibitory protein, (cFLIP) [also called CASPER, I-FLICE, CASH, FLAME-1, NRIT, CLARP, or usurpin (reviewed in[18]). Functional cFLIP contains two DED domains and a caspase-like domain which lacks proteolytic enzymatic activity. Rather, cFLIP molecules bind to the DEDs in FADD and inhibit apoptosis by interfering with the recruitment of procaspase-8 to FADD.

PCD is also held in check by the inhibitor of apoptosis proteins (IAPs). IAPs are the Baculovirus IAP Repeat (BIR)-containing proteins which interfere with the apoptotic pathway by binding to TNF Receptor Associated Factor 2 (TRAF2) as well as caspases (reviewed in[34]). These inhibitors are therefore not exclusive to the receptor-mediated pathway but can also inhibit the “intrinsic” pathway as they have the capacity to inhibit caspases. IAPs have also been shown to have ubiquitin protein ligase activity which is triggered upon apoptotic stimulation, resulting in autodegradation of the IAP protein and thereby promoting cell death[35].

The intrinsic pathway of mammalian PCD

In addition to the signals received through cell surface death receptors, cells also sense, in an as yet unknown way, death signals through the mitochondria. The mitochondrial initiating complex consists of cytochrome c, dATP and Apaf-1, known as the apoptosome[36]. The apoptosome facilitates PCD through its ability to recruit and activate caspase-9.

Apaf-1 is a 130 kDa protein composed of three functional domains: a short N-terminal CARD, a central CED-4 homology domain, and a long C-terminal “WD-40 repeat” domain[9]. In the presence of cytochrome c and dATP, Apaf-1 can interact with pro-caspase 9 through their mutual CARDS[19]. Oligomerization of Apaf-1 in the apoptosome triggers autocatalysis of pro-caspase-9 leading to its activation[37]. The WD-40 repeats of Apaf-1 serve an important regulatory function because their deletion results in constitutive processing and activation of caspase-9 independent of cytochrome c and dATP [37]. Other regulatory controls of the intrinsic pathway include the anti-apoptotic Bcl-2 members[38] and Bcl-xL[39], which can block cytochrome c release, and the pro-apoptotic Bcl-2 members, which can promote the release[40], [41].

Lessons from mutant animal models

Much of our present understanding of the apoptotic pathways and their component molecules has come from gain-of-function and loss-of-function studies in animal models. During *C. elegans*' development, 131 cells normally undergo PCD[4]. Loss-of-function mutations in *ced-3*, *-4*, and *egl-1* result in survival of all of the 131 cells, supporting their pro-apoptotic role. However they live a normal life with no obvious deleterious effects. In contrast, *ced-9* mutants die early as a consequence of excessive cell death. Gain-of-function mutations in *ced-9* block all 131 cell death, supporting its anti-apoptotic

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role[42]. In this review, we will discuss the phenotypes of the various mutant mice that have shed light on the roles and importance of individual molecules in the apoptotic pathways.

Analysis of mutants: extrinsic pathway

In mammals, the extrinsic pathway is vital for a normal healthy existence, as is illustrated by the phenotypes of mice mutated in various components of the pathway. *lpr* and *gld* mice, which carry spontaneous mutations of the Fas or FasL genes respectively, show dramatic accumulations of lymphocytes resulting in splenomegaly and lymphadenopathy. They exhibit an autoimmune disorder with pathology that is reminiscent of the human autoimmune disease systemic lupus erythematosus (SLE) [43]. Fas has been shown to be essential for the activation-induced cell death of T cells [44] which may account for the lymphoproliferation and the autoimmunity observed in *lpr* and *gld* mice.

The phenotypes of mice lacking the extrinsic pathway components FADD, caspase-8 or cFLIP have not only provided *in vivo* information about the known apoptotic pathway, but have also revealed previously unknown roles for these genes in other cellular processes[45-47]. FADD^{-/-} and caspase-8^{-/-} cell lines show resistance to death receptor-mediated PCD whereas cFLIP^{-/-} cell lines are more susceptible, as was expected [45, 46]. In addition, all three mutants show abnormalities in heart development, an observation that remains unexplained to date. Even though cFLIP appears to have the opposite function to FADD and caspase-8 in the apoptotic pathway, cFLIP^{-/-} embryos exhibit the same heart anomalies as FADD^{-/-} and caspase-8^{-/-} embryos suggesting that in another as yet unidentified pathway, these molecules cooperate with one another.

Analysis of mutants: caspase-3

Caspase-3 is the executioner caspase that serves as one of the key effectors in both the intrinsic and extrinsic pathways. The most dramatic phenotype observed in caspase-3^{-/-} mice is a defect in embryonic brain development that results in the presence of neuronal supernumerary bodies due to a defect in apoptosis and embryonic lethality of some mice starting at day E12.5[48]. Despite the absence of caspase-3, most caspase-3^{-/-} cells can be induced to undergo normal PCD, implying redundancy of function with other executioner caspases. However, caspase-3 is indispensable for the nuclear changes that occur during apoptosis[49], although these nuclear hallmarks are not in fact required for PCD[50].

The analysis of caspase-3 deficient mice illustrates that components of the death pathways act in a tissue/cell type-specific and stimulus-specific manner. For example, caspase-3^{-/-} embryonic stem (ES) cells die normally following γ -irradiation and heat shock but are resistant to PCD induced by UV-irradiation or hyperosmolarity. caspase-3^{-/-} immature T cells in the thymus are susceptible to Fas-induced cell death but caspase-3^{-/-} mature peripheral T cells are resistant to the same stimulus[49].

In addition, caspase-3^{-/-} surviving adult mice injected with agonistic anti-Fas antibody survive longer than wildtype littermates. We can therefore conclude that caspase-3 is indeed required at least partially in Fas-mediated hepatocyte apoptosis. Interestingly, in the absence of caspase-3, cytochrome c release was also impaired, as well as the cleavage of ‘upstream’ caspases such as caspase-8 and -9. These findings may be explained by the interesting phenomenon that Bcl-2 and Bcl-x_L are cleaved exclusively by caspase-3. We show that caspase-3’s role may be expanded to that of a regulator of apoptosis, in addition to its executionary role supported in the previous study[51].

Analysis of mutants: the intrinsic pathway

caspase-9 deficient mice show a dramatic brain phenotype similar to that seen in caspase-3^{-/-} embryos[52], [53]. Activated caspase-3 is absent from caspase-9^{-/-} brains demonstrating that a linear epistatic process operates in this tissue. Caspase-9 is also required for apoptosis of thymocytes, as determined by the resistance of these cells to PCD induced by dexamethasone or γ -irradiation[53] However, caspase-9 is not required for extrinsic pathways of PCD such as TNF α - or Fas-mediated cell death[53].

Apaf^{-/-} deficient mice exhibit striking cranio-facial abnormalities associated with decreased apoptosis of neuronal cells in the developing brain[54], [55]. The brain defects observed in apaf-1^{-/-} mice are more profound than those observed in either caspase-9^{-/-} or caspase-3^{-/-} mutants, suggesting that Apaf-1 may be involved in apoptotic pathways other than those involving caspase-3 or caspase-9. Mutation of Apaf-1, but not caspase-3 or caspase-9, results in a delay in the removal of the embryonic interdigital webs, and in abnormal eye development. Furthermore, similar to caspase-9 deficiency, apaf-1^{-/-} cells are resistant to a wide variety of apoptotic stimuli, including γ -irradiation, dexamethasone and chemotherapeutic agents but not from death receptor-mediated killing.

Embryos devoid of cytochrome c are not viable beyond midgestation. Nevertheless, development up to this stage appears to progress in a relatively normal but delayed manner[56]. cytochrome c-deficient cultured cells from living embryos display respiratory insufficiency and enhanced anaerobic glycolysis, manifested by slowed growth and rapid acidification of the culture medium. cytochrome c^{-/-} embryonic cells are more resistant to apoptosis induced by UV-irradiation, serum starvation or staurosporin. In contrast, apoptosis in response to TNF α is enhanced in the absence of cytochrome c. This interesting observation suggests a previously unrecognized interaction between the intrinsic and extrinsic pathways, in which a defect in one upregulates the other.

Characterization of caspase-2^{-/-} mice found these mice to have an excess number of germ cells in their ovaries that are resistant to chemotherapeutic agents. On the other hand, caspase-2^{-/-} sympathetic neurons are more sensitive to apoptosis[57]. Caspase-12 has been shown to be required specifically for endoplasmic reticulum (ER) stress mediated apoptosis[58].

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Crosstalk between the two pathways

The seemingly distinct pathways described so far do not always occur independently in vivo. For example, although in some cell types, such as T lymphocytes, ligation of Fas via FADD and caspase-8 can induce effector caspase activation and culminate in cell death without affecting the mitochondria as shown in Apaf-1 and caspase-9 knock-out. However, the mitochondrial pathway is involved upon death receptor triggering in some situations. In the scenario where FADD and caspase-8 are sufficient for Fas-mediated apoptosis[28,27], cytochrome c is not released and effector caspases are directly activated early on[59]. In this mitochondria-independent pathway, cell death is not inhibitable by Bcl-2 but rather by direct caspase inhibitors such as CrmA[59]. However, in other cell types, such as hepatocytes, apoptotic induction by Fas ligation results in cytochrome c release and the process is inhibitable by Bcl-2, implying the involvement of a mitochondrial pathway. One protein that may account for this discrepancy is a BH3 containing Bcl-2 family member called Bid[60,61].

Bid is cleaved by caspase-8 and the cleaved carboxy terminal peptide (tBID) is targeted to the mitochondria where it triggers cytochrome c release[62]. Bid^{-/-} hepatocytes exhibit a defect in cytochrome c release and resistance to apoptosis, thereby illustrating that the mitochondrial pathway is required for amplifying death signals[63].

Other studies show that tBid can somehow cause a change in the conformational state of Bax. This results in oligomerization of Bax and facilitates its insertion into the outer mitochondrial membrane, triggering mitochondrial dysfunction and subsequent cytochrome c release[64]. Another group has shown that tBid is also able to induce a conformational change and oligomerization of Bak[65]. It is postulated that different cell types use different pro-apoptotic Bcl-2 proteins.

Novel roles for caspases

Some recent studies have suggested that caspases may be involved in cellular processes other than apoptosis. Caspase-3 and -8 have been shown to be important for T cell proliferation[66], [67], and caspase-3 may play a role in cell cycle regulation in splenic B cells. In vitro, caspase-3 can cleave numerous proteins involved in variety of pathways, including molecules important for cell cycle and survival signaling[68]. We have shown that caspase-3^{-/-} splenic B cells have an intrinsic capacity to hyperproliferate in the absence of a cell death defect. Indeed, caspase-3^{-/-} B cells have a greater clonogenic capacity and show a higher proportion of cells in S phase than wild type B cells. These data suggest that caspase-3 may be required for cell cycle arrest rather than for apoptosis in certain cell types (Data not shown/Manuscript in preparation).

CONCLUSION

At present, most of our knowledge of PCD signaling pathways is at the cellular level. PCD and its interaction with other processes is much more complex at the organism level. It is imperative that we understand these mechanisms in vivo because so many

human diseases result as a consequence of defects in PCD regulation. Once the cellular mechanisms of cell death are understood *in vivo*, we can devise more effective therapeutic strategies for disease prevention and progression.

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