

Apoptosis and the Cell Cycle

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Abstract In this review, we consider apoptosis as a process intimately linked to the cell cycle. There are several reasons for thinking of apoptosis as a cell cycle phenomenon. First, within the organism, apoptosis is almost exclusively found in proliferating tissues. Second, artificial manipulation of the cell cycle can either prevent or potentiate apoptosis, depending on the point of arrest. Data from such studies have suggested that molecules acting late in G1 are required for apoptosis. Since passage through late G1 into S phase in mammalian cells is known to be regulated by p53 and by activation of cyclin-dependent kinases, we also examine recent studies linking these molecules to the apoptotic pathway. © 1995 Wiley-Liss, Inc.

Key words: DNA replication, chromatin condensation, premature mitosis, cyclin A, p53, cyclin-dependent kinases

Not all cell deaths are accidental. In animal development, proper shape and function require cell death as well as cell division. For example, fingers and toes are formed from the death of cells in the interdigital tissue [Kerr et al., 1987; Novack and Korsmeyer, 1994]. The elaboration of even more complex structures also depends on cell death. Formation of the T and B cell repertoires of the immune system is dependent on negative selection (elimination of nonreactive or self-reactive cells) as much as positive selection [Cohen, 1991]. Similar forces are at work in shaping the nervous system [Raff, 1992]. In development of the nematode *Caenorhabditis elegans*, exactly 131 cells die according to a precisely regulated genetic program [Yuan and Horvitz, 1990]. Cell death is also necessary as a protective mechanism. Examples include the directed lysis of infected cells, foreign cells, or incipient neoplasms by the immune system [Bishop and Whiting, 1983; Duke, 1991; Squier and Cohen, 1994] and the autolysis of immune cells themselves following damage by γ radiation [Umansky, 1991]. It is striking that in these various cases of cell death, there exist features sufficiently unique as to constitute a well-defined cytological entity, known as apoptosis: the form of cell death in which the cell is de-

stroyed from within [Kerr et al., 1972; Wyllie et al., 1980]. A cardinal feature of apoptotic cells is the appearance of highly condensed chromatin, segregated into sharply defined bodies within an intact nuclear envelope. Often, the DNA is digested by endonucleases, making detection of DNA fragments a commonly used assay for apoptosis [Arends et al., 1990]. The cell shrinks and condenses, fragmenting into multiple, membrane-bound bodies (apoptotic bodies) which are eventually engulfed by surrounding cells—thus removed without inflammation and attendant damage to surrounding tissue. It is this constellation of features that is referred to as apoptosis.

The thought that multicellular organisms have evolved genetic mechanisms for promoting cell death presents something of a puzzle—how might genes be selected that have death as a phenotype? One way is by linking cell death to cell proliferation. Whereas in yeast essentially all proliferation controls ultimately feed into a single gene, *cdc2* in *S. pombe* or *CDC28* in *S. cerevisiae*, in multicellular eukaryotes cell cycle control is split up amongst a dozen *cdc2/CDC28* homologs, whose products are referred to as cyclin-dependent kinases (CDKs) because of their dependence on a cyclin subunit for full activity and cell cycle regulation. If independent activation of several CDK-cyclin complexes were required for cell division to proceed, activation of more or fewer than the proper number could be fatal. Depending then on the physiological environment, withholding activating stimuli (hor-

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mones, growth factors, cytokines) or supplying inappropriate stimuli (cell contact, antigen binding, receptor ligation) might lead to incorrect timing of CDK activation. In this scenario, the "death genes" are then actually a subset of the genes whose normal function is to mediate cell proliferation, and apoptosis is an incomplete, abnormal cell division [Ucker, 1991]. Such a hypothesis suggests that it might be possible to induce apoptosis by premature activation of CDKs by chemical or other artificial means. In yeast, mutations in regulatory genes impinging on *cdc2/CDC28* can lead to premature activation of *cdc2/CDC28*, which brings about a "mitotic catastrophe": a premature mitosis in which chromatin condensation, spindle formation, and abortive division is disastrous for the cell. In mammalian cells, the morphological similarities between apoptosis and premature mitosis are quite suggestive of a role for CDKs in apoptosis [Rubin et al., 1993; Sen and D'Incalci, 1992; Ucker, 1991]; in fact, such a role has recently been demonstrated [Meikrantz et al., 1994; Shi et al., 1994]. In this review, we discuss apoptosis as a process intimately linked to the cell cycle.

APOPTOSIS AS A CELL CYCLE PHENOMENON Association With Proliferation

In vivo, apoptosis is detected primarily in proliferating tissues [Kerr and Harmon, 1991; Wyllie et al., 1980]. Anywhere in the body where cells are dividing, apoptotic bodies are to be found: at low frequency in self-renewing tissues (like intestinal crypt) [Potten, 1977], in epithelium of the adrenal cortex [Wyllie et al., 1973a,b], in the differentiating spermatogonium [Allan et al., 1987], and in the germinal centers of lymph nodes [Swartzendruber and Congdon, 1963]. Apoptosis becomes particularly evident after periods of rapid proliferation, and is thus detected in mammary tissue following weaning [Marti et al., 1994; Walker et al., 1989], in the endometrium at estrus [Gerschenson and Rotello, 1991; Sandow et al., 1979; Otsuki et al., 1994], during ovarian follicular atresia [O'Shea et al., 1978], and in numerous other cases [e.g., Kerr and Harmon, 1991; Kerr et al., 1972; Ledda-Columbano and Columbano, 1991; Wyllie et al., 1980]. As might be expected, apoptosis is a prominent feature in malignancies [Moore, 1987] and occurs at increased frequency in preneoplastic tissues [Ledda-Columbano and Columbano, 1991]. In these cases, apoptosis is thought to serve a compensatory role: balancing an increase in cell

number due to proliferation with cell loss due to death. The role of apoptosis in maintaining tissue homeostasis is well-illustrated in regression of liver tissue following lead nitrate- or cyproterone acetate-induced hyperplasia [Bursch et al., 1985; Columbano et al., 1985]. Administration of either mitogen leads to an increase in organ size due to massive cell proliferation. Upon withdrawal of the mitogen, the hyperplastic tissue regresses and the organ returns to normal size. Regression is entirely due to apoptotic death.

Both positive and negative signals trigger cells to stop proliferating and undergo apoptosis. Positive signals include antigen receptor ligation of immature lymphocytes [Green and Scott, 1994], thyroxine-induced regression of larval components in amphibian metamorphosis [Yoshizato, 1989], glucocorticoid-induced thymocyte death [Wyllie, 1980], and absorption of the Müllerian structures in the male fetus in response to Müllerian-inhibiting substance [Glucksmann, 1951; Saunders, 1966]. Negative signaling comprises withdrawal of necessary growth factors. For example, sympathetic neurons depend on a continuous supply of nerve growth factor from the cells they innervate [Raff, 1992; Rubin et al., 1993], while cells in the ventral prostate rapidly undergo apoptosis if testosterone is removed by castration [reviewed in Buttyan, 1991]. Survival factor dependence is also found in tumor cell lines. For example, mouse myeloma lines will undergo apoptosis following withdrawal of IL-2 and IL-6 [Colotta et al., 1992]. Interestingly, apoptosis induced by factor withdrawal is associated with induction of proliferation-associated genes [Buttyan et al., 1988; Colotta et al., 1992]. This topic is discussed in greater detail below.

Cell Cycle Arrest and Apoptosis

Ultimately, the signals described above must impinge on the cell cycle if proliferation is to be halted and cells are to die. How this is accomplished requires understanding where in the cell cycle this intervention occurs. We consider two types of analysis: 1) cell cycle effects of physiological mediators of apoptosis, and 2) effects of artificially imposed cell cycle synchrony on apoptosis induced by other agents.

1. Cell cycle phase-specific death induced by physiological mediators of apoptosis. The WEHI-231 line is a murine B cell lymphoma which retains many of the characteristics of immature B cells [Boyd and Schrader, 1981]. It displays surface IgM but not IgD, and

does not secrete immunoglobulin. When treated with anti-IgM antibodies, mimicking antigen-induced surface IgM cross-linking, these cells die by apoptosis after approximately 24–48 h [Benhamou et al., 1990; Hasbold and Klaus, 1990]. Flow cytometry revealed anti-IgM-treated cells were blocked from entering S phase: they ceased incorporating [³H]thymidine into DNA [Scott et al., 1985] and did not phosphorylate pRB [Maheswaran et al., 1991; Warner et al., 1992]. Growth arrest in G1 was apparent within 12 h, well before the morphological changes characteristic of apoptosis had occurred. Antibody treatment of G1 cells, obtained by centrifugal elutriation, prevented entry into S phase with arrest very close to the G1-S border [Scott et al., 1986].

T cell hybridomas are also sensitive to activation-induced death following ligation of their antigen receptors [Green and Scott, 1994]. Following stimulation, they rapidly accumulate in G1 prior to the appearance of an apoptotic morphology [Ashwell et al., 1987]. The ability of antisense *c-myc* oligonucleotides to halt growth and prevent apoptosis in such cells [Shi et al., 1992] implies a requirement for molecules required for continued proliferation/initiation of DNA replication (see below). Similarly, exposure of lymphocytic lines to dexamethasone, a model agent for glucocorticoid-induced deletion of T cells/thymocytes, induces G1 arrest prior to morphological evidence of killing [Harmon et al., 1979].

The bursa of Fabricius, the primary site of B cell development in birds, contains a cohort of stem cells that gives rise to a rapidly proliferating population of lymphoblasts with IgM on their surface. Bursal stem cells and their progeny have been shown to be very sensitive to apoptosis induced by mechanical disruption of cell-cell contact in short term culture of bursal follicles as well as by γ radiation and other DNA damaging agents [Neiman et al., 1991]. Apoptosis is apparent within the first hour following tissue dispersion, and occurs primarily in S phase cells: fluorescence-activated cell sorting based on DNA content followed by agarose gel electrophoresis revealed that DNA degradation was occurring primarily in S phase cells. Apoptotic cells were positive for several molecular markers associated with S phase of the cell cycle [Neiman et al., 1994].

Since many growth factors act as progression factors [Pardee, 1989], promoting passage from G1 to S phase of the cell cycle, it is expected that

apoptosis resulting from factor withdrawal correlates with arrest in G1. This is the case when ND7 cells, prepared by fusing rat dorsal root ganglion neurons with a neuroblastoma line, are transferred to serum-free medium [Howard et al., 1993] and when a variety of hematopoietic cell lines are deprived of particular interleukins [Nuñez et al., 1990; Colotta et al., 1992]. An exception to this generalization is the FVA cell line (murine erythroid hematopoietic progenitor cells infected with the anemia-inducing strain of Friend virus). These cells undergo apoptosis when deprived of erythropoietin [Kelley et al., 1992; Koury and Bondurant, 1990]. Cell loss occurs in the G1 and S phases of the cell cycle, but without apparent growth arrest [Kelley et al., 1994].

These data seem to place entry into apoptosis around the time of entry into S phase. Cells may halt abruptly in G1, without entering S, or may actually show evidence of having begun DNA replication. Since there are always exceptions to any generalization, we note that a human B cell lymphoma analogous to the murine WEHI-231 line seems to undergo activation-induced apoptosis from G2 rather than G1 [Ishigami et al., 1992].

2. Cell cycle arrest modifies the response to apoptosis-inducing agents. Cytotoxic lymphocytes (CTL) kill target cells by mechanisms that independently disrupt the cell membrane and induce apoptosis [Greenberg and Litchfield, in press; Squier and Cohen, 1994]. Inhibitors of replication fork-associated enzymes prevented DNA fragmentation and apoptosis in target cells, but not membrane lysis, suggesting that apoptosis was dependent on cell proliferation [Nishioka and Welsh, 1992]. In fact, target 3T3 fibroblasts could be made refractory to CTL-induced DNA fragmentation by serum deprivation. Infection of these quiescent 3T3 cells with herpes simplex virus 1 forced induction of DNA synthesis and restored susceptibility to CTL-induced apoptosis [Nishioka and Welsh, 1994].

Proliferating, mature T lymphocytes can be induced to undergo apoptosis after ligation of their antigen receptor [Boehme and Lenardo, 1993]. When proliferating A.E7 cells, a nontransformed CD4⁺ T_H1 clone, were stimulated via the T cell receptor (TCR) by anti-CD3 ϵ antibodies or via irradiated splenocytes and antigen, cells died by apoptosis. The extent of cell loss was directly proportional to the extent of proliferation. When A.E7 cells were incubated with cell cycle arrest agents prior to TCR ligation, cells that were blocked in G1 by mimosine, deferoxamine, or

dibutyryl cAMP were resistant to apoptosis, while cells arrested at G1/S by aphidicolin or in early S by excess thymidine were sensitized. G1 arrest did not simply select for a resistant population of cells, since release from arrest rapidly led to S phase and apoptosis. Like anti-IgM-mediated apoptosis of WEHI-231 cells, molecules acting in late G1/early S seem to be implicated in induction of apoptosis.

Another physiological inducer of apoptosis which has received careful cell cycle scrutiny is TNF- α . A variety of data indicate that TNF- α induces apoptosis in G1 in L929 cells [Darzynkiewicz et al., 1984; Kirstein et al., 1986; van de Loosdrecht et al., 1993]. However, G1 arrest with TGF- β , TNF- α , IL-1 β , and IL-6 did not render cells more sensitive to TNF- α . Instead, cells were protected against TNF- α -induced cytotoxicity [Belizario and Dinarello, 1991]. This suggests that these agents may arrest cells at a point in G1 prior to the synthesis of TNF-sensitivity factors. Studies with human epithelial cells suggest that TNF sensitivity appears late in G1. Human HeLa cells are normally resistant to TNF- α , but, like numerous other cell types, can be sensitized to TNF- α by cotreatment with cycloheximide [Wallach, 1984], an agent which arrests HeLa cells late in G1, near the G1-S boundary. Cell cycle arrest in S phase with hydroxyurea was as effective at sensitizing HeLa cells as cycloheximide, suggesting a requirement for molecules acting at the time of the G1 to S transition in TNF- α -induced apoptosis [Meikrantz et al., 1994].

S phase arrest also potentiates apoptosis induced in HeLa cells by numerous agents with a wide variety of pharmacological activities, including staurosporine, 6-dimethylaminopurine, okadaic acid, caffeine, and γ -radiation [Meikrantz et al., 1994; Meikrantz and Schlegel, unpublished observations]. At the doses used, the majority of these agents have little or no effect on HeLa cell proliferation. In the case of staurosporine, arrest earlier in the cell cycle with lovastatin, which causes a reversible G1 arrest [Jakobisiak et al., 1991; Keyomarsi et al., 1991], did not potentiate apoptosis. Thus, a wide variety of apoptosis-inducing agents seem to require molecules synthesized in late G1/early S.

Apoptosis Induced by p53

Transfection of wild type p53 into transfected cells reduces colony forming ability [Baker et al., 1990; Isaacs et al., 1991; Johnson et al., 1991; Takahashi et al., 1992] and invari-

ably leads to growth arrest in G1 [Diller et al., 1990; Martinez et al., 1991; Mercer et al., 1990; Michalovitz et al., 1990]. In murine myeloid leukemic lines, which typically lack p53, restoration of the wild type gene results in apoptosis [Ryan et al., 1993; Yonish-Rouach et al., 1991, 1993]. By transfecting M1 cells, a mouse myeloid leukemia line, or DP16-1 MEL cells, a Friend virus-induced murine erythroleukemia line, with a temperature-sensitive p53, cell cycle effects of p53-induced apoptosis were studied by shifting cells to the permissive temperature [Ryan et al., 1993; Yonish-Rouach et al., 1993]. In both cases, cells in G1 were preferentially susceptible to p53-induced killing. Treatment of the transfected M1 cells with IL-6, which arrests these cells in G1 [Resnitzsky and Kimchi, 1991; Resnitzsky et al., 1992], prevented apoptosis. However, TGF- β , another G1-arrest agent, failed to protect against p53-induced apoptosis; instead, in the presence of TGF- β , p53-dependent death was accelerated [Yonish-Rouach et al., 1993]. One explanation for these results is that IL-6 and TGF- β arrest at different points in G1, before and after synthesis of molecules required for p53-dependent apoptosis. Density-dependent growth arrest of transfected DP16-1 cells also blocked p53-dependent apoptosis, while cells released from the block died rapidly after shifting to the permissive temperature. Cells synchronized in G1 with mimosine or by isoleucine deprivation underwent apoptosis much more quickly than density-arrested cells. If arrested cells were released and allowed to progress out of G1 before shifting to the permissive temperature, apoptosis did not occur until cells had returned to G1 [Ryan et al., 1993]. The different response rates of density-arrested versus mimosine or isoleucine deprivation arrested cells again indicates that molecules required for p53-dependent apoptosis are synthesized at a later point in G1.

Apoptosis Induced by Perturbation of DNA Metabolism

Agents that block DNA replication, induce DNA damage, interfere with DNA topology, or block the segregation of chromosomes are lethal to the cell. Perhaps unsurprisingly, the majority of these agents have been found to induce apoptosis [see the tabulation in Sen and D'Incalci, 1992]. In some cases apoptosis occurred with little or no cell cycle specificity [Cotter et al., 1992]. In other cases, careful flow cytometric analysis revealed cell cycle differences in toxicity

and susceptibility. For example, HL-60 cells, a human promyelocytic leukemia line, were preferentially affected in G1 by nitrogen mustard; cells progressing through S phase were more susceptible to hydroxyurea, camptothecin, and teniposide; while G2/M cells preferentially underwent apoptosis in response to γ -radiation [Gorczyca et al., 1993]. These data raise two important points. First, it is possible that some agents directly activate apoptosis regardless of the cell cycle stage. Second, it is possible that cell cycle-related differences in response to such agents tells less about pathways of apoptosis induction than about the effect of the cell cycle stage on the severity of damage or the capacity to repair particular lesions [Gorczyca et al., 1993].

Apoptosis in Quiescent Cells

Several studies have reported apoptosis in quiescent cells. One such model system is involution of the prostate following androgen withdrawal, a process that has been studied extensively in the rat following castration. Within days of castration, the terminally differentiated secretory epithelial cells which line the ducts of the gland begin to undergo massive apoptosis. These cells, which compose as much as 85% of the cell population of the prostate, are lost within a week after castration [Lesser and Bruchofsky, 1973; Kerr and Searle, 1973], making this is an excellent tissue for studying the biochemistry and molecular biology of apoptosis. A number of apoptosis-specific proteins and "testosterone-repressed" messages have been identified [see Buttyan, 1991 for review]. Most interesting was the sequential induction of *c-fos* and *c-myc*, since induction of these genes, particularly in this sequence, is usually associated with the transition from quiescence to proliferative growth [Muller et al., 1984; Thompson et al., 1986], including regrowth of the prostate after androgen replacement [Katz et al., 1989]. This finding suggests that the quiescent acinar epithelial cells were being induced to reenter a defective cell cycle [Colombel et al., 1992] that ends in apoptosis rather than continuing DNA replication. In fact, cyclin A is expressed in regressing prostate, suggesting that these cells progress to a point quite late in G1 (R. Buttyan, personal communication). A similar early induction of *c-fos*, *junB*, *junD*, and *c-jun* is seen during involution of mammary epithelium at weaning [Marti et al., 1994] and during apoptosis induced by growth

factor withdrawal from a myeloid leukemia line [Colotta et al., 1992].

Normally, *c-myc* is down-regulated during the transition from proliferative growth to quiescence; in fact, down regulation occurs in a cell cycle-independent manner as soon as mitogen is withdrawn [Dean et al., 1986; Waters et al., 1991]. When *c-myc* is expressed ectopically in Rat-1 fibroblasts or in primary rat embryo fibroblasts during serum withdrawal, proliferative growth is maintained and cells undergo apoptosis [Evan et al., 1992]. Similarly, apoptosis can be induced in serum-starved, quiescent Rat-1a fibroblasts following expression of *myc* from the Zn²⁺-inducible metallothionein promoter [Hoang et al., 1994]. Since *myc* expression stimulates transcription (either directly or indirectly) of a number of genes acting late in G1 that are necessary for the G1 to S transition [Buchou et al., 1993; Hoang et al., 1994; Jansen-Dürr et al., 1993; Kim et al., 1994; Shibuya et al., 1992], ectopic expression of *c-myc* may force cells into a late G1-like state of preparedness for DNA replication or apoptosis.

A similar interpretation may apply to induction of apoptosis in quiescent cells by adenovirus E1A. Adenovirus E1A induces apoptosis when expressed in the absence of E1B 19K, a second adenovirus protein [Pilder et al., 1984; Rao et al., 1992; Subramanian et al., 1984; White et al., 1984a,b, 1992]. Infection of normal rat kidney cells with an adenovirus producing only the E1A protein and lacking the E1B region caused apoptosis only when the cells were growth arrested, either by growth to confluence or by serum starvation [Mymryk et al., 1994]. Like *myc*, E1A activates transcription of a number of genes required for DNA replication [Buchou et al., 1993; Moran, 1993; Nevins, 1992; Whyte et al., 1988]. In fact, E1A mutants that no longer promote DNA synthesis in rat kidney cells are also defective for inducing apoptosis [Howe et al., 1990].

In general, it appears that cells must progress to late G1 of the cell cycle for apoptosis to occur. Arrest prior to this stage delays or blocks apoptosis while arrest after this promotes apoptosis. What is significant about this particular moment in the cell cycle? First of all, this is the p53 restriction point. In normal cells, passage through this point of the cell cycle is at least partially regulated by p53, and it has become increasingly evident that many types of apoptosis-inducing agents are p53-dependent. What is

the role of p53 in apoptosis? How is the role of p53 in apoptosis related to its role as a cell cycle regulator? Secondly, this is the cell cycle stage often referred to as the "R point" in mammalian cells [Pardee, 1974] or "start" in yeast [Hartwell et al., 1974], the stage where cells become irreversibly committed to DNA replication. In organisms from yeast to man, passage through this point requires activation of specific CDKs. Are there restriction point-related CDKs whose activation is necessary for apoptosis to occur? We shall take up these questions in turn.

APOPTOSIS AND THE G1 RESTRICTION POINT: p53 AND p21

The idea that p53 acts as a "guardian of the genome" [Lane, 1992], monitoring a restriction point that requires cells to repair their DNA damage before entering S phase or else face apoptosis, is consistent with the finding that cells or cell lines derived from p53-deficient animals fail to undergo apoptosis in response to DNA damaging agents [Clarke et al., 1993, 1994; Lowe et al., 1993a,b; McCarthy et al., 1994; Merritt et al., 1994]. Loss of p53 function would then lead to fixation of genetic abnormalities resulting from DNA replication in the presence of DNA damage. If p53 were to act by eliminating cells with compromised DNA, the integrity of the organism would be maintained, although at the expense of cell loss due to apoptosis. Unlike the unicellular yeasts, which lack p53, this is a viable option for a multicellular organism. The proliferation of multiple malignancies in p53^{-/-} mice could result from the persistent proliferation of cells with DNA damage. The ability of p53 to induce apoptosis on its own after re-introduction into p53-deficient cells, and its ability to limit the growth of existing neoplasms through apoptosis [Liu et al., 1994; Radinsky et al., 1994; Symonds et al., 1994], demonstrates a role for p53 in limiting neoplastic growth via induction of apoptosis. How p53 accomplishes its growth-limiting function must be related to its ties to the cell cycle and to control of CDK activation and inactivation, for it is via CDKs that the cell cycle progresses.

Cyclins are proteins whose levels rise and fall in an orderly way during the course of the cell cycle. They are positive regulatory subunits for the CDKs, conferring upon them substrate and cell cycle specificity. In addition to binding of the cyclin subunit, CDKs are regulated by activating and inactivating phosphorylations and by

association of inhibitory CDK-interacting proteins (CIPs). These interactions are summarized schematically in Figure 1, which depicts an idealized CDK-cyclin complex. Events required to prepare the cell for entry into S phase are mediated by the D and E type cyclins [Sherr, 1994]. Principal among these events is phosphorylation of the product of the retinoblastoma susceptibility gene, pRB. Unphosphorylated pRB, present throughout most of G1, acts as a damper to cell proliferation by sequestering E2F transcription factors, which are necessary for transcription of genes required for entry into S phase. Phosphorylation of pRB by cyclin D- and cyclin E-associated CDKs releases the transcription factors and allows proliferation to advance. Ultimately, commencement of DNA replication requires activation of CDK-cyclin A complexes, which occurs at the time of the G1-S transition, a process described in more detail below.

Although identified (as *CIP1*) in a two-hybrid screen for proteins interacting with cdk2 [Harper et al., 1993] and purified (as CAP20) from mouse cells by virtue of its association with an inactive population of cdk2 [Gu et al., 1993], the p21 CIP has been shown to bind to and inactivate in vitro virtually every member of the CDK-cyclin family [Harper et al., 1993; Xiong et al., 1993]. Clues as to its physiological function come from how it was identified in two other systems. First, p21 was discovered as a senescent cell-derived inhibitor of DNA replication (*sd1*) in human diploid fibroblasts by screening cDNAs prepared from senescent human diploid fibroblasts for their ability to inhibit [³H]thymidine uptake upon transient overexpression in young, dividing fibroblasts [Noda et al., 1994]. Second, a connection to p53-dependent inhibition of growth was realized when p21 was discovered independently (under the name *WAF1*, for wild type p53-activated fragment 1) as a message whose transcription was entirely dependent on wild type p53 during p53-mediated growth arrest [El-Deiry et al., 1993]. p21 is induced in p53-dependent G1 arrest which follows γ irradiation of normal human diploid fibroblasts, resulting in inhibition of cdk2-cyclin E kinase activity, preventing phosphorylation of pRB, and thereby blocking entry into S phase [Dulić et al., 1994]. Similarly, in transformed cells (wild type p53-expressing colorectal carcinoma lines RKO and HCT-116), doxorubicin (Adriamycin) treatment induced p21 expression and blocked cdk2-cyclin E kinase activity. There was no induction of p21

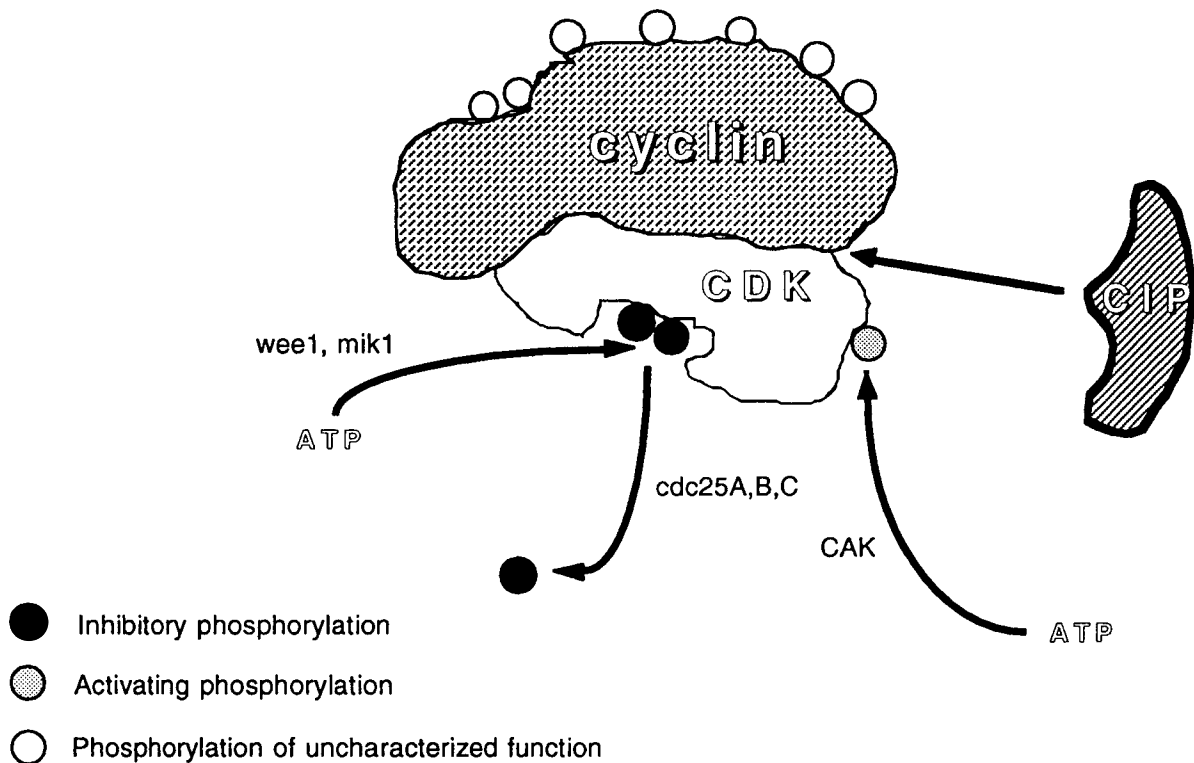


Fig. 1. Activation and inactivation of a model CDK-cyclin complex.

and little or no decrease in cyclin E-associated kinase activity following similar treatment of p53-deficient tumor lines treated with doxorubicin [El-Deiry et al., 1994]. While induction of p21 by serum- or growth factor-stimulation of quiescent cells [Noda et al., 1994] is independent of p53, induction of p21 by DNA damaging agents is completely dependent on p53 [Michieli et al., 1994]. Expression of p21 in a number of tumor-derived cell lines was able to suppress tumor growth. Expression of the p21 gene in the sense but not the antisense orientation resulted in a 10- to 20-fold decrease in colony-forming ability in p53-deficient tumor lines (the colon tumor line SW480, the brain tumor line DEL, and the lung adenocarcinoma line H1299 [El-Deiry et al., 1993]). Upon exposure to γ radiation, tumor cells containing wild type p53 induced p21 and underwent apoptosis, while p21 and apoptosis were lacking in p53-deficient cells [El-Deiry et al., 1994]. Thus, induction of p21 and consequent inhibition of CDKs may be a first step in p53-dependent apoptosis.

Interestingly, p53 and p21 also appear to induce a cell cycle arrest via mechanisms that are independent of CDK-cyclin complexes. SV40 DNA replication can be supported in an in vitro

system containing T antigen and human DNA replication proteins [Waga and Stillman, 1994]. Addition of excess p21 to the reaction caused an accumulation of early DNA replication intermediates [Waga et al., 1994], suggesting that p21 was interfering with DNA elongation but not initiation/melting of origins of replication. Addition of the p16 CIP was without effect. The p21 inhibition could be overcome by addition of excess PCNA (proliferating cell nuclear antigen), a subunit required by DNA polymerase δ . This inhibition took place in the absence of detectable cyclins or CDKs, and involved formation of a complex between p21 and PCNA [Waga et al., 1994]. p53 itself physically interacts with a protein required for DNA replication, replication factor A (RPA, also referred to in the literature as RF-A or RP-A), a single-stranded DNA binding protein complex required for DNA replication [Fang and Newport, 1993; Waga and Stillman, 1994]. Binding of p53 to RPA blocks DNA replication [Dutta et al., 1993; He et al., 1993; Li and Botchan, 1993]. This may explain the observation that UV-induced apoptosis in a T antigen-immortalized pituitary (somatotrophic) line expressing a temperature sensitive p53 mutant occurs in a p53-dependent manner (i.e., only at

the permissive temperature), but in the absence of p53-dependent transcription/translation (i.e., in the presence of actinomycin D or cycloheximide). Since shifting to the permissive temperature resulted in translocation of p53 to the nucleus, the authors propose that p53 affects apoptosis via direct participation in DNA repair or DNA degradation [Caelles et al., 1994].

Regardless of mechanism, the data strongly suggest that the first steps on the pathway of p53-dependent apoptosis are a cell cycle block at the beginning of S phase. In this light, it is interesting to note that cell cycle arrest is also a feature of several p53-independent pathways of apoptosis, for example, the activation-induced death of WEHI-231 cells, dexamethasone deletion of thymocytes, and TNF- α cytotoxicity. Furthermore, S phase arrest was able to promote apoptosis by p53-independent apoptosis-inducing agents in a T cell hybridoma [Boehme and Lenardo, 1993] and in HeLa cells, which are human papillomavirus positive and thus lack functional p53 [Meikrantz et al., 1994; Meikrantz and Schlegel, unpublished observations]. HeLa cells are normally very resistant to γ radiation-induced apoptosis. When HeLa cells are arrested in S phase, however, this resistance is overcome. We suggest that S phase arrest by agents such as hydroxyurea, thymidine, and aphidicolin chemically impose the cell cycle arrest normally mediated via p53. This immediately suggests that transient overexpression of p21—i.e., imposition of a molecular block to proliferation as opposed to a chemical one—will substitute for chemical agents in sensitizing cells to apoptosis inducers. Evidence that p21 is actually required for apoptosis comes from the observation that antisense constructs of p21 block p53-mediated apoptosis (J.C. Barrett, personal communication).

Growth arrest by p53 cannot be the only trigger for apoptosis, as can be inferred from the different responses to γ radiation of transformed and nontransformed cells expressing wild type p53. Nontransformed cells typically show only growth arrest, while transformed cells undergo apoptosis [Dulić et al., 1994; El-Deiry et al., 1994]. Primary fibroblasts from p53^{+/+} mice do not display a significant apoptotic response to γ radiation unless they are transformed by a combination of oncogenes (*E1A* and *H-ras*) [Lowe et al., 1993a]. These findings strongly suggest that an additional proliferation-associated step is required for apoptosis. Cell prolifera-

tion and passage through the G1 restriction point requires activation of specific CDKs, particularly cyclin E- and cyclin A-dependent protein kinases [Dou et al., 1993]. We have seen that inactivation of CDK-cyclin E complexes is associated with p53-dependent growth arrest and apoptosis; now we shall examine a role for activation of CDK-cyclin A complexes in apoptosis.

APOPTOSIS AND THE G1 RESTRICTION POINT: ACTIVATION OF CDKS

Activation of cdk2-cyclin A is required for the transition from G1 to S. Cdk2 becomes activated as a protein kinase at the time of entry into S phase [Rosenblatt et al., 1992]. Inactivation of cdk2 by microinjection of neutralizing antibodies to cyclin A [Pagano et al., 1992] or cdk2 [Pagano et al., 1993], treatment with antisense cyclin A nucleotides [Girard et al., 1991], or transfection of a dominant negative mutant of cdk2 [van den Heuvel and Harlow, 1993] prevents cells from entering S phase. Cdk2-cyclin A seems to affect DNA replication directly. Cdk2-cyclin A complexes are localized at foci of nuclear replication in terminally differentiated myotubes induced to re-enter S phase by transfection with T antigen [Cardoso et al., 1993], and phosphorylation of the p34 subunit of RPA is required for initiation and maintenance of DNA replication [Fang and Newport, 1993; Waga and Stillman, 1994]. Immunodepletion of cdk2 prevents DNA synthesis in a *Xenopus* cell free system [Fang and Newport, 1991] by preventing phosphorylation of the p34 subunit of RPA [Fang and Newport, 1993]. Like *cdc2*, cdk2 is subject to inactivating phosphorylations at the active site [Gu et al., 1992; Sebastian et al., 1993]. Cdk2 is dephosphorylated and activated by CDC25A: transcription of CDC25A begins in early G1 and peaks in late G1, and microinjection of anti-CDC25A antibodies blocks entry into S phase [Jinno et al., 1994].

Activation of cdk2-cyclin A appears to be the necessary, final step required for exit from G1 and commencement of DNA replication. Cyclin A is unique among the cyclins, being required at two points in the cell cycle: both for mitosis and for DNA replication [Girard et al., 1991; Pagano et al., 1992; Zindy et al., 1992]. In this light, it is interesting to note that apoptosis seems to depend on cell cycle processes acting at or about the time of the G1-S transition and comprises some of the same morphological changes found

in mitosis [Lazebnik et al., 1993; Rubin et al., 1993; Sen and D'Incalci, 1992; Ucker, 1991], the most prominent being chromatin condensation. Could activation of cyclin A-dependent protein kinases be the molecular link between apoptosis, the cell cycle, G1-S, and mitosis? Circumstantial evidence certainly seems to point in this direction. Arrest in the cell cycle prior to the appearance of cyclin A tends to protect cells from apoptosis [Belizario and Dinarello, 1991; Boehme and Leonardo, 1993; Colotta et al., 1992; Howard et al., 1993; Meikrantz et al., 1994; Nishioka and Welsh, 1994; Nuñez et al., 1990; Ryan et al., 1993; Yonish-Rouach et al., 1993]. This is not merely reflective of a requirement for p53 synthesis, since protection extends to p53-independent apoptosis-inducing agents as well (e.g., TNF- α , TCR ligation). Two oncogenes which induce apoptosis under restrictive growth conditions (i.e., in the absence of serum or without expression of a cooperating oncogene), *c-myc* and adenovirus E1A, are both transcriptional activators of the cyclin A gene [Buchou et al., 1993; Jansen-Dürr et al., 1993; Shibuya et al., 1992]. In the rat prostate, cyclin A follows *c-myc* as one of the proliferation-associated genes whose transcription is enhanced during castration-induced apoptosis (R. Buttyan, personal communication). WEHI-231 cells accumulate increased levels of cyclin A protein following anti-IgM antibody treatment (Meikrantz, Boothby, and Schlegel, unpublished observations). Apoptosis due to prolonged S phase arrest (22 h) of the early CD4⁺ T cell line AGF in high concentrations of thymidine is associated with translocation of cyclin A and *cdc2* to the nucleus [Gazitt and Erdos, 1994]. Phosphorylation of the p34 subunit of RPA, which depends on *cdk2* [Fang and Newport, 1993], is stimulated by exposure of cells to ionizing radiation [Liu and Weaver, 1993]. This suggests that *cdk2*-cyclin A activity may actually be stimulated by apoptosis-inducing agents. Finally, staurosporine, a known activator of CDKs, including *cdk2*-cyclin A [Meikrantz et al., 1994; Tam and Schlegel, 1992], is an acute activator of apoptosis in human cells [Jacobson et al., 1993; Meikrantz et al., 1994].

In S phase-arrested HeLa cells, apoptosis was induced by agents known to activate CDKs (caffeine, 6-dimethylaminopurine, staurosporine, and okadaic acid), with activation of cyclin A-dependent kinases occurring within 2 h after exposure [Meikrantz et al., 1994]. The extent of cyclin A-dependent kinase induction (two- to sevenfold) correlated well with the extent of

apoptosis induction (15–80% of cells with apoptotic nuclei at 8 h following treatment). Where examined, both *cdc2*-cyclin A and *cdk2*-cyclin A were activated. These results raised the following question: since CDK-activating agents induced apoptosis, would other apoptosis-inducing agents activate CDKs? This was found to be the case with TNF- α , which has previously been reported to act synergistically with chemical CDK activators such as caffeine [Belizario et al., 1993] and staurosporine [Beyaert et al., 1993]. HeLa cells are normally resistant to TNF- α , unless it is given simultaneously with cycloheximide [Wallach, 1984] or to S phase-arrested cells [Meikrantz et al., 1994]. In hydroxyurea-arrested cells, TNF- α stimulated CDK-cyclin A activity about threefold. Cotreatment with TNF- α and cycloheximide increased kinase activity about sevenfold. Interestingly, cycloheximide alone induced a fivefold increase in CDK-cyclin A activity, suggesting a mechanism for its potentiation of TNF- α killing. HeLa cells in this model system had to be *arrested* in S phase in order to undergo apoptosis: cells passing synchronously through S phase were not sensitized to killing. We suggest that this system is a "chemical" model for apoptosis: S phase arrest by hydroxyurea chemically imposes the cell cycle arrest component, while TNF- α or chemical treatment provides the signal(s) for CDK activation.

Induced transcription of cyclin A followed by CDK-cyclin A activation also takes place in serum-starved Rat 1a fibroblasts during *myc*-induced apoptosis [Hoang et al., 1994]. Two types of analysis were carried out to study cyclin expression in this system. First, cell lines were created in which *myc* was constitutively overexpressed. In these cells, serum withdrawal led to extensive apoptosis, whereas serum withdrawal from cultures of parental cells had little effect. Analysis of cyclin transcription in the *myc*-overexpressing cells showed that cyclin A transcription was elevated compared with parental Rat 1a cells, while transcription of cyclin B, cyclin C, cyclin D1, and cyclin E were unaffected. A second group of cell lines was then created, expressing cyclin A under control of the Zn²⁺-inducible metallothionein promoter. Serum withdrawal plus exposure to 50 μ M Zn²⁺ led to apoptosis. Not all of the cells expressed cyclin A to the same level following Zn²⁺ induction, and the extent of apoptosis observed in low serum plus Zn²⁺ conditions indicated a dose-dependent induction of apoptosis by cyclin A. Thus, in the

presence of growth arrest by serum starvation, expression of cyclin A alone was sufficient to induce apoptosis.

Is CDK-cyclin A activation required for apoptosis? The ability of a wide variety of apoptosis-inducing agents to induce CDK-cyclin A activity in S phase-arrested HeLa cells is suggestive of a common requirement for this enzyme, and the sufficiency of cyclin A induction alone to promote apoptosis in serum-starved fibroblasts is further evidence for such a mechanism. However, the most direct evidence for a requirement for CDK activity in apoptosis comes from studies of cytolysis induced by fragmentin-2/granzyme B, a protease component of the lytic granules released by cytotoxic T cells or natural killer cells during destruction of their targets [reviewed in Greenberg and Litchfield, in press]. Fragmentin-2/granzyme B rapidly (within a few hours) induced apoptosis when added to detergent-permeabilized cells. Premature activation of *cdc2* was found to be required for fragmentin-2/granzyme B-induced apoptosis [Shi et al., 1994]. *Cdc2* was rapidly activated at the initiation of apoptosis, and DNA fragmentation and nuclear condensation could be prevented by an excess of *cdc2* peptide substrate. At least part of the induced kinase activity has subsequently been found to be associated with cyclin A (A. Greenberg, personal communication). Most importantly, fragmentin-2/granzyme B-induced apoptosis was temperature dependent in FT210 cells, which are homozygous for a temperature sensitive *cdc2* allele [Th'ng et al., 1990]. At the restrictive temperature, where mutant *cdc2* is rapidly degraded, fragmentin-2/granzyme B-induced apoptosis was inhibited by as much as 50%. This strongly implicates CDK-cyclin A activation as a necessary step in apoptosis induction. Although the maximum inhibition seen was 50%, we note that FT210 cells contain a significant amount of *cdk2*-cyclin A that is unaffected by shift to the *cdc2*-restrictive temperature [Hamaguchi et al., 1992]. Since both *cdc2*-cyclin A and *cdk2*-cyclin A are activated in HeLa cells undergoing apoptosis, it is possible that elimination of all cyclin A-dependent kinase activity might inhibit apoptosis to an even greater degree. The relative predominance of *cdc2*-cyclin A complexes versus *cdk2*-cyclin A complexes early in the cell cycle seems to vary from cell type to cell type and as a function of growth conditions. For example, significant differences between monolayer and suspension HeLa cultures have been reported [Elledge et al., 1992].

Recently, we have found that transient expression of dominant negative mutants of *cdc2* and *cdk2* both suppress TNF- α -induced apoptosis in HeLa cells, although *cdk2*-cyclin A seems to be the crucial complex mediating apoptosis in these monolayer cells [Meikrantz and Schlegel, manuscript in preparation].

Much has been made in the literature of the morphological similarities between mitotic catastrophe in yeast, premature mitosis in mammalian cells, and apoptosis. It has been suggested that apoptosis is a premature, abortive mitosis. At the beginning of this review, we presented the conceptual similarity between apoptosis and mitotic catastrophe in yeast, relating the two processes to checkpoint controls that impinge on CDKs. However, some clear biochemical distinctions can be made between the two processes. First and most importantly, *cdc2*-cyclin B, whose activation is absolutely required for mitosis [Nurse, 1990] and premature mitosis [Steinmann et al., 1991] is not activated during apoptosis in HeLa cells [Meikrantz et al., 1994]. Instead, activation of CDK-cyclin A complexes is observed. We note that cyclin A has no clear counterpart among the yeast cyclins. The chromatin morphology induced by premature activation of *cdc2*-cyclin B is different from that induced in apoptosis: chromosome spreads prepared from apoptotic cells stain homogeneously, typically containing one or more nonstaining "holes" [Arends et al., 1990; Meikrantz et al., 1994], and do not present the characteristic "pulverized" appearance of premature chromatin condensation [Rao and Johnson, 1970]. In a cell free system that mimicks many of the morphological characteristics of apoptosis [Lazebnik et al., 1993], depletion of *cdc2*-cyclin B did not suppress the response, mitosis-specific phosphorylation of the MPM-2 proteins was not detected, and the nuclear lamins were not phosphorylated. In fact, the nuclear lamina disassembly that was seen seems to have been due to proteolysis rather than phosphorylation of the lamin proteins.

Although we have suggested that apoptosis correspond may to the cyclin A-dependent half of mitosis [Meikrantz et al., 1994], it could equally be viewed as a premature and aberrant entry into S phase, resulting in condensation and cleavage of chromatin rather than DNA replication (see Fig. 2). Thus, while agents like staurosporine induce premature mitosis in G2-arrested cells, they induce apoptosis in G1/S-arrested cells. In this sense, apoptosis resembles

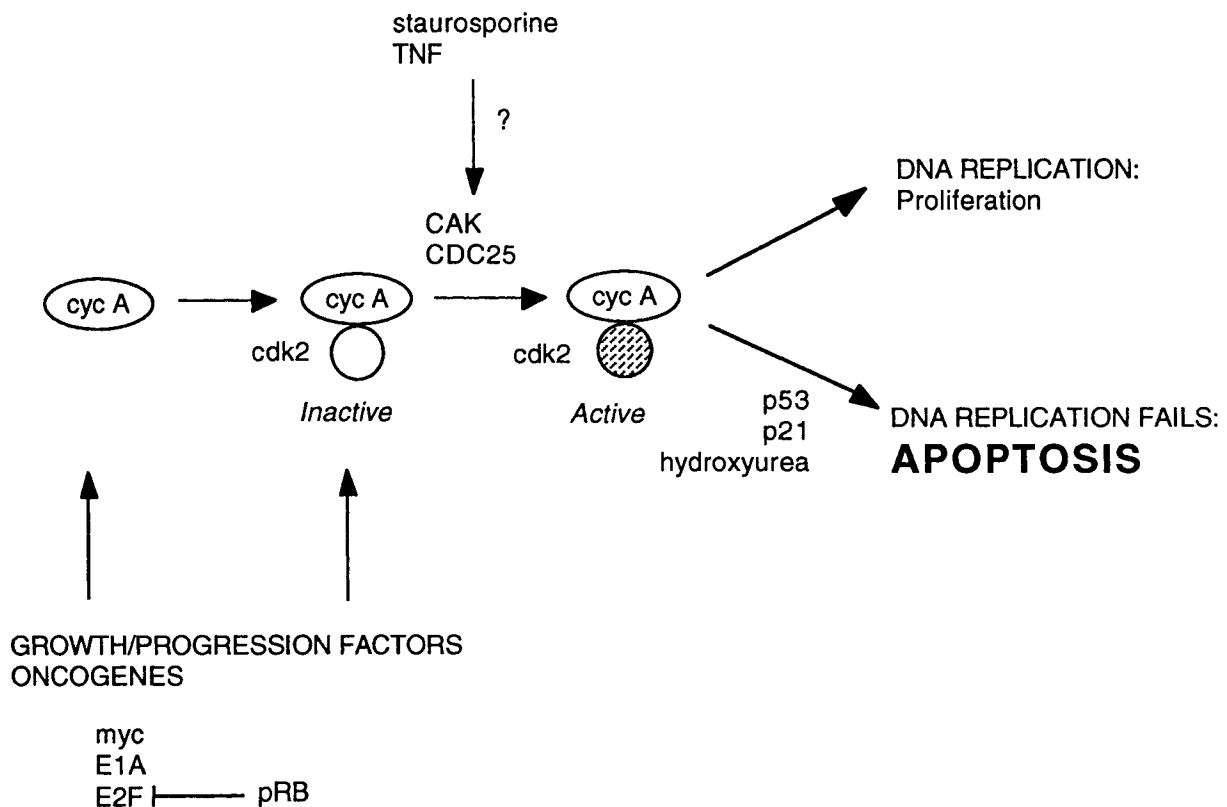


Fig. 2. Relationship of CDK-cyclin A activation to DNA replication and apoptosis.

mitotic catastrophe: both events are characterized by improper timing of CDK activation and by uncoupling the normal interdependence of cell cycle events.

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