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p53, Apaf-1, caspase-3, and -9 are dispensable for Cdk5 activation during cell death

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Abstract

Cyclin-dependent kinase 5 (Cdk5) is a member of the cyclindependent kinase family that is mostly seen in neurons, does not vary with cell cycle, and is activated in many neurodegenerative disorders and other non-neuronal pathologies, but its relationship to non-neuronal apoptosis is not understood, nor is the control of the activation of Cdk5 by its activators. The most widely studied activator of Cdk5, p35, is cleaved to p25 by calpain, an event that has been linked with activation of Cdk5 and neuronal death. Here we report that calpain-mediated Cdk5/p25 activation accompanies nonneuronal as well as neuronal cell death, suggesting that the p35/calpain/p25/Cdk5 activation sequence is a general feature of cell death. We further demonstrate that Cdk5 can be activated in the absence of p53, Apaf-1, caspase-9, and -3 during cell death, indicating that its activation relates more to cell death than to a specific pathway of apoptosis.

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Abbreviations: Cdk, cyclin-dependent kinase; Cdk5, cyclindependent kinase 5; CHX, cycloheximide; CPT, camptothecin; CS, calpastatin; EtOH, ethanol; PARP, poly (ADP ribose) polymerase; PD, PD150606; SBDPs, spectrin breakdown products; TUNEL, terminal deoxynucleotidyl transferase-mediated dUTP nick end labeling

Introduction

Here we document that activation of cyclin-dependent kinase 5 (Cdk5) occurs during cell death in mesodermal cells in the same situations as has been reported for neurons and other cells, thus demonstrating that activation of Cdk5 occurs generally in most or all dying cells. We then address the mechanism of its activation during cell death.

Cdk5 is a member of the cyclin-dependent kinase (Cdk) family originally identified from bovine brain by its sequence

homology to human Cdc2.¹ Cdk5 is expressed at both mRNA and protein levels in a broad spectrum of tissues and cell lines. with the highest levels detected in neurons.²⁻⁶ Unlike other cell cycle-promoting members of Cdk family, Cdk5 kinase primarily modulates neuroskeletal dynamics in postmitotic neurons.^{3,7,8} Cdk5 plays an essential role in both developing and adult neurons, including neuronal migration,^{9,10} axon guidance,¹¹ neurite outgrowth,⁷ dynamics of synaptic structure,^{12–14} neurotransmission,^{15,16} and neuronal secretion.¹⁷ Cdk5 null mice suffer perinatal mortality, with severe defects in neuronal migration and laminar configuration in the cerebral cortex.¹⁸ However, Cdk5 is not restricted to neurons. It is also prominent in the testis, where its level is high but its function is unknown (our unpublished data), developing muscle, 19,20 bFGF-stimulated bovine aortic endothelial cell proliferation,²¹ and differentiating monocytes.22,23

Cdk5 is activated in dying cells, including both developing tissues (interdigital areas of mouse limbs,²⁴ differentiating rat lens,⁴ and mouse brain⁵) and adult tissues (regressing mouse prostate following androgen withdrawal and atrophic mouse ovaries⁵). The association between Cdk5 activation and cell death is seen in pathology: rat lungs infected with *Strepto-coccus pneumoniae*,²⁵ mouse limbs infected with *Leishmania major*,²⁶ preterm lamb lungs after ventilation,²⁷ cyclophosphamide-treated mouse embryos,²⁸ and several neuro-disorders and neuronal injuries.^{29–37} The regulatory signals and mechanisms by which Cdk5 is activated remain undefined.

As with other Cdks, Cdk5 is activated by association with an activator. p35 is the most important regulatory subunit of Cdk5 during neurodevelopment,^{38–40} and we confirmed that it interacts with Cdk5 during cell death.²⁸ Cleavage of p35 to p25 by calpain has been linked to activation of Cdk5. Inhibition of calpain activation, which blocks p25 production and Cdk5 activation, prevents neuronal cell death.^{29,31,32,35,41,42} Here we document the generation of p25 during cell death, as we had previously suggested.²⁸ We also establish that calpain-mediated Cdk5/p25 activation coincides with cell death in both non-neuronal and neuronal systems, suggesting the generality of the role of Cdk5 in cell death.

The signals that activate calpain and Cdk5 are unclear, as is the question of how activation of Cdk5 relates to cell death. To address this latter issue, we investigate the relationship of Cdk5 to major cell death pathways. We first examine the relationship of Cdk5 to the p53-mediated signaling pathway.^{43–45} p53 may mediate mitochondrial cell death signaling through the elevation of the level of reactive oxygen species,⁴⁶ while oxidative stress can stimulate Cdk5 activity by inducing an upregulation of p35 in human neuroblastoma IMR-32 cells.⁴⁷ Reactive oxygen species (ROS) are instrumental in a number of neurodegenerative disorders where Cdk5 is activated.^{48,49} Furthermore, *in vitro* p53 is a substrate of Cdk5.⁵⁰ We therefore investigate crosstalk between p53 and Cdk5.

Activation of the apoptosome and the subsequent activation of caspase-9 and -3 follow mitochondrial dysfunction⁵¹ in the intrinsic pathway. We therefore also ask how Cdk5 interacts with these caspases. Finally, we demonstrate that Cdk5 can be activated during cell death in the absence of p53, Apaf-1, caspase-9, and -3.

Nevertheless, activation of Cdk5 is a consistent feature of many or most deaths of ectodermal and mesodermal cells. Since Cdk5 activation is coincident but independent of these pathways, the mechanism by which it is activated and the function of its activation remain to be elucidated.

Results

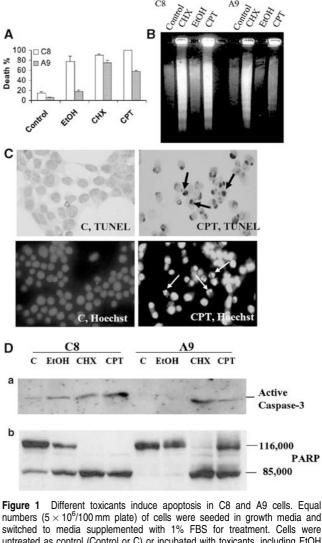
Apoptotic cell death is induced by different stimuli in C8 and A9 cells

To examine the involvement of p53 in the activation of Cdk5, we used C8 and A9 cells, which are mouse embryonic fibroblast cells oncogenically transformed with both *E1A* and *ras*, and containing wild-type p53 or p53–/–, respectively.⁴³ Toxins such as ethanol (EtOH), cycloheximide (CHX), and camptothecin (CPT) kill these cells. EtOH at high concentrations damages the cell membrane.⁵² CHX inhibits protein synthesis and is otherwise toxic at the pharmacological levels normally used to induce apoptosis.⁵³ CPT inhibits DNA topoisomerase I.⁵⁴ All lead to apoptosis in a variety of cells.^{55–58}

Death induced in C8 and A9 cells after exposure to EtOH (2.5%), CHX ($25 \mu g/ml$), or CPT ($20 \mu M$) for 18h was determined by trypan blue (TB) exclusion assay. Owing to their sensitivity to low serum (1% FBS), untreated control C8 cells manifested of the order of 15% cell death (Figure 1A). All three toxins induced substantial cell death in C8 – up to 77, 90, and 100%, respectively (Figure 1A). In p53–/– A9 cells, after 18h CHX and CPT induced 75 and 58% cell death, respectively (Figure 1A), indicating that these compounds could induce p53-independent cell death. EtOH, however, caused 77% cell death in C8 cells, but only 18% cell death in A9 cells (Figure 1A), indicating that EtOH induced p53-dependent death in C8.

EtOH, CHX, and CPT can induce apoptosis or necrosis, depending on cell line or conditions.^{54,59,60} To evaluate the cell death induced under our conditions, we employed several markers to characterize apoptotic cell death, including examination of DNA fragmentation both by gel electrophoresis and *in situ* detection of fragmented DNA by terminal deoxynucleotidyl transferase-mediated dUTP nick end labeling (TUNEL), measurement of nuclear fragmentation by Hoechst staining, and activation of caspase-3 and cleavage of poly (ADP ribose) polymerase (PARP).

As shown in Figure 1B, in C8, all three chemicals induced DNA ladders above background, indicating fragmented chromatin and internucleosomal DNA fragmentation. CHXand CPT-treated A9 cells also showed typical DNA ladders, but ladders were barely detectable in EtOH-treated cells, correlating with the low cell death. Control A9 cells were negative. In C8, DNA fragmentation was further confirmed by TUNEL, and nuclear fragmentation and condensation were detected by Hoechst staining in all treated samples, as exemplified in Figure 1C. CHX- and CPT-treated A9 cells



numbers (5 × 10⁶/100 mm plate) of cells were seeded in growth media and switched to media supplemented with 1% FBS for treatment. Cells were untreated as control (Control or C) or incubated with toxicants, including EtOH (2.5%), CHX (25 μ g/ml), and CPT (20 μ M), for 18 h. (**A**) Levels of cell death induced in C8 and A9 were determined by trypan blue exclusion. (**B**) DNA ladder was examined on a 2% agarose gel. (**C**) Untreated and CPT-treated C8 cells were fixed with 3% paraformaldehyde, and DNA fragmentation was determined by TUNEL assay (upper panels). C, TUNEL control, no signal; CPT, TUNELpositive signal is the brown dark staining in dying/dead cells as indicated by arrows. Nuclear morphology was examined using Hoechst fluorescent dye (lower panels). Arrow indicates fragmented nuclei. (**D**) Cells were lysed with RIPA buffer, and equal amounts of lysates were resolved by SDS-gel electrophoresis, followed by immunoblotting with active caspase-3 antibody (a) or PARP antibody (b), as described in Materials and Methods

were also TUNEL positive and showed nuclear fragmentation (data not shown).

Activation of caspase-3 was demonstrated by Western blot (WB) using an antibody that detects active 17 kDa caspase-3. In C8 cells treated with EtOH, CHX, or CPT for 18 h, caspase-3 was active (Figure 1D, a), and its activation corresponded to the level of cell death (Figure 1A). Activation was further confirmed by PARP cleavage by WB. When C8 cells were exposed to the three toxins, the 85 kDa band corresponding to caspase-3-specific PARP cleavage was substantially above the basal level corresponding to the 15% background cell death shown in Figure 1A (Figure 1D, a and b). In A9 cells, as in C8 cells, we found caspase-3 activation (Figure 1D, a) and PARP cleavage (Figure 1D, b) during cell death by CPT and CHX. However, in EtOH-treated A9 cells, there was neither caspase-3 activation nor PARP cleavage (Figure 1D, a and b). Thus, the death of C8 cells induced by EtOH, CHX, or CPT was apoptotic. CHX and CPT can induce apoptotic cell death independent of p53 in A9, while induction of cell death by EtOH depends on p53.

We further evaluated the kinetics of CPT-induced cell death, activation of caspase-3, and activation of Cdk5 in C8 and A9. By 6 h, C8 cell death was statistically greater than that in A9 (49% positive C8 cells by TB, compared with 4% in A9; Figure 2A) and continued in this trajectory for the next 2 h. However, by 12 h, A9 cells started to die (20%), increasing to 40% by 15 h (Figure 2A). Activation of caspase-3 started at 6 h in C8 (Figure 2B, a), correlating with strong induction of p25, indicative of Cdk5 activation (Figure 2B, b). Induction of p25 is a reliable indicator and measure of Cdk5 activation.^{28,29,31} In

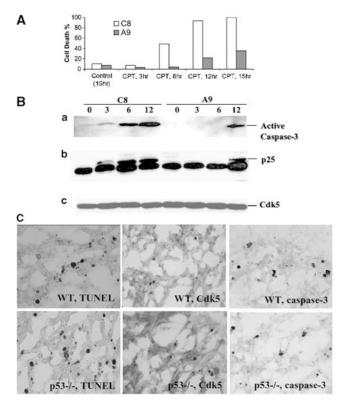


Figure 2 Lack of p53 causes a delayed induction of cell death and activation of Cdk5, but does not affect cell death or Cdk5 protein expression in developing mouse embryos. (A) There is a delayed induction of cell death in p53-deficient A9 cells. C8 and A9 cells were exposed to CPT, and cell death was determined by trypan blue exclusion. Averages of four independent experiments are shown, with S.D. less than 5%. (B) Immunoblots of protein lysates from C8 and A9 probed with antibody against active caspase-3 (a), p35 (b), or Cdk5 (c) cells indicate a delayed but eventual clear activation of Cdk5 and caspase-3 in CPT-induced cell death in A9 cells. (C) Both p53 wild-type and mutant mouse embryos (ED 13.5) were fixed and sectioned as described in Materials and Methods. TUNEL assay and immunohistochemistry study were performed on near-serial sections for comparison. There is no difference in the level or pattern of cell death and Cdk5 protein expression between the brains of wild-type and p53-/- embryos. Caspase-3 activation further indicates a caspase-3-involved cell death in this region. Magnification, $\times 40$

A9, it took 12 h to activate caspase-3 (Figure 2B, a) and induce p25 (Cdk5 activation – Figure 2B, b). As expected, the level of Cdk5 protein was constant for the 18 h in both C8 and A9 (Figure 2B, c, data for 8, 15, and 18 h not shown). Thus, although p53 was not required for p53-independent apoptosis, such as CHX- and CPT-induced cell death, lack of p53 resulted in a delay of both apoptosis and Cdk5 activation. Cdk5 activation correlates with apoptosis, but not p53.

The influence of p53 on the activation of Cdk5 was further examined in p53 knockout mouse embryos. Developing mouse embryos manifest apoptotic cell death in many organs. We first examined apoptotic cell death by TUNEL in developing p53-/- mouse embryos at ED13.5, 14.5, and 15.5 (data for 14.5 and 15.5 not shown). TUNEL assay on sections of wild-type and p53-/- embryos indicated substantial levels of cell death in the trigeminal region of both embryos (Figure 2C); this death was therefore p53-independent. On an adjacent serial slide, the strong caspase-3 immunoreactivity indicated activation of caspase-3 in both wild-type and p53-/- embryos (Figure 2C). Similarly, Cdk5 immunoreactivity, indicative of its activation,²⁸ was detected in both wild-type and p53 knockout mouse embryos. We found no difference in the level of cell death or activation of Cdk5 between wild-type and p53-/- (Figure 2C), again correlating Cdk5 with cell death, but not p53.

Cdk5 is activated in the absence of functional apoptosome (Apaf-1) and subsequent caspase-9 activation

To address whether an intact intrinsic cell death pathway is required for Cdk5 activation, we first examined Apaf-1. We used Apaf-1 null mouse embryos at stages ED12.5, 13.5, and 15.5. In all the three stages in both wild-type and null mice, cell death correlated with Cdk5 expression. In ED13.5 Apaf-1 knockout mouse embryos, many cells in the periphery of the liver were positive for both DNA fragmentation and Cdk5 (Figure 3A, a, black arrows). Therefore, Apaf-1 is not required for Cdk5 activation during cell death.

As confirmation, primary mouse embryonic fibroblasts lacking Apaf-1 as well as their wild-type counterparts were exposed to CHX (25 μ g/ml) or CPT (20 μ M). Cells began to show apoptotic morphology at 36 h. By this time, 40% of CHX-exposed Apaf-1+/+ and 17% of Apaf-1-/- cells were TB-positive, while 45% of CPT-exposed Apaf-1+/+ and only 20% of Apaf-1-/- cells were dead (Figure 3A, b). Apaf-1-deficient cells are more resistant to cell death, in accordance with the argument that Apaf-1 is apoptogenic.

Cdk5 protein did not change (Figure 3A, c). However, Cdk5 was activated, as indicated by generation of p25 in all dying cells (Figure 3A, d). Thus, again, Cdk5 activation correlates with cell death. In these experiments, activation can occur in the absence of Apaf-1.

Caspase-9 is the direct downstream target of the apoptosome. We therefore investigated the relationship between caspase-9 and Cdk5 activation. Both caspase-9–/– primary mouse embryonic fibroblast cells and their counterparts were exposed to CHX (25 μ g/ml) or CPT (20 μ M) and cell death was determined by TB. At 24 h after exposure, CHX induced 88

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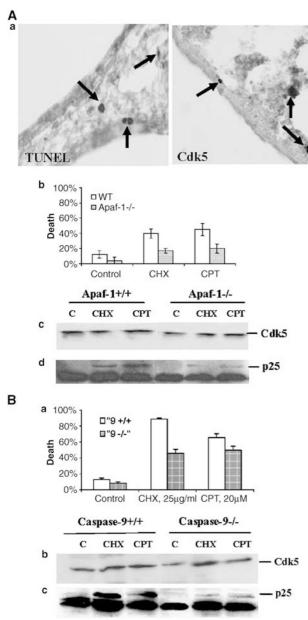


Figure 3 Cdk5/p25 activation during cell death does not require a functional apoptosome. (A) Cdk5 activation during cell death is Apaf-1-independent. (a) Apaf-1 mutant mouse embryos (ED 13.5) were fixed and sectioned as described in Materials and Methods. TUNEL assay and immunohistochemistry study were performed on near-serial sections for comparison. TUNEL assay demonstrates cells with fragmented DNA in the periphery of liver in Apaf-1 knockout embryo and immunohistochemistry using Cdk5 antibody shows cells expressing Cdk5 protein in the same region. Magnification, \times 100. (b-d) Cdk5 is activated during cell death in both wild-type and Apaf-1-/- MEFs. Wild-type and Apaf-1-/-MEFs were exposed to CHX (25 µg/ml) or CPT (20 µM) for 36 h. Cell death was measured by trypan blue (b), and Cdk5 protein (c) and p25 production (d) was examined by immunoblotting of proteins from treated cells using Cdk5 or p35 antibody. (B) Cdk5 is activated during cell death in both wild-type and caspase-9-/- MEFs. Wild-type and caspase-9-/- MEFs were exposed to CHX (25 μ g/ ml) or CPT (20 μ M) for 24 h. Cell death was measured by trypan blue (a), and Cdk5 protein (b) and p25 production (c) was examined by immunoblotting of proteins from treated cells using Cdk5 or p35 antibody

and 46% cell death in wild-type and caspase-9–/– cells, respectively. CPT killed 65% wild type and 50% caspase-9–/– cells (Figure 3B, a). Cdk5 expression did not differ between wild-type and caspase-9-deficient cells (Figure 3B, b). p25 was generated where cell death was induced in both cells, and the level of the p25 induction (Figure 3B, c) corresponded with that of cell death measured by TB exclusion (Figure 3B, a). Thus, Cdk5 is activated whether or not caspase-9 is present.

Therefore, both Apaf-1 and caspase-9 are dispensable for Cdk5 activation during cell death, and Cdk5 can be activated in apoptosome-independent cell death.

Cdk5 can be activated in the absence of caspase-3

Caspase-3 is the primary effector caspase. We therefore investigated the relationship of Cdk5 activation to activation of caspase-3. TUNEL assay and immunohistochemistry were

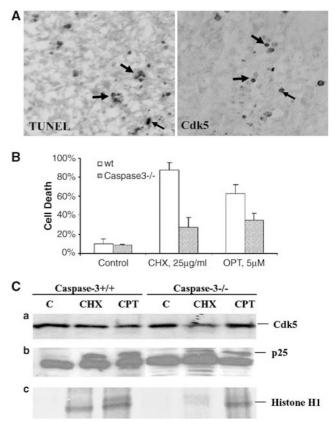


Figure 4 Cdk5/p25 activation during cell death does not require caspase-3 activity. (A) Caspase-3 knockout mouse embryos (ED 13.5) were fixed and sectioned as described in Materials and Methods. Cell death and Cdk5 expression in the brain of caspase-3 knockout mouse embryos were examined by TUNEL and immunohistochemistry, respectively. TUNEL assay demonstrates cells with fragmented DNA (left panel, black arrows) in the brain of caspase-3 knockout embryo, and Cdk5 antibody reveals cells expressing Cdk5 protein (right panel, black arrows) in the same brain region. Magnification, \times 40. (B, C) Wild type and caspase-3 –/– MEFs were exposed to CHX or CPT for 24 h. Cell death was reduced in the knockouts, when it occurred, Cdk5 amounts were not changed (C, a) but p25 was produced (C, b) and the kinase was activated (C, c), indicating that activation does not require caspase-3

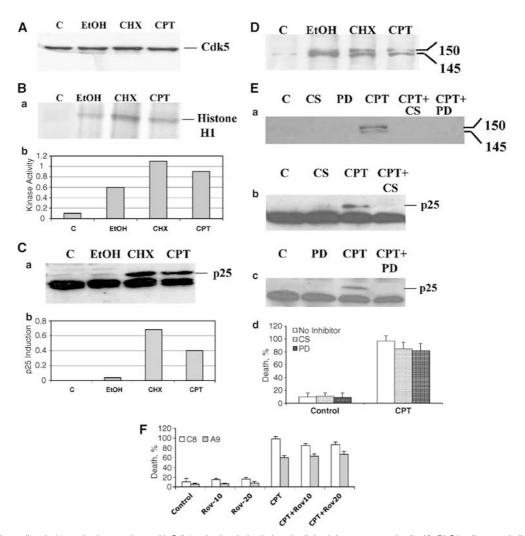


Figure 5 Calpain-mediated p25 production correlates with Cdk5 activation during induced cell death in non-neuronal cells. (**A–D**) C8 cells were challenged with EtOH, CHX or CPT for 18 h. Cells were collected and lysed with RIPA buffer. (**A**) Immunoblot probed with Cdk5 antibody of equal protein samples as noted above. There is no significant difference in the amount of Cdk5 protein expressed. (**B**) Kinase activity of Cdk5 immunoprecipitate was determined using histone H1 as an *in vitro* substrate in the presence of γ [³²P]ATP. The incorporated γ [³²P] in phosphorylated histone H1 was resolved on a 12.5% acrylamide gel and identified by autoradiography (a) and quantified by densitometry (b). Kinase activity is increased by the toxins. (**C**) Immunoblot probed with p35 antibody (a) to identify p25 as quantified by examining the production of specific SBDPs at 150 and 145 kDa, resulting from cleavage of nonerythroid α -spectrin by calpain. Calpain is activated in all three experimental situations. (**E**) C8 cells were treated with CPT (20 μ M) for 18 h, in the presence or absence of either of the two calpain inhibitors, CS (10 μ M) or PD (10 μ M). Cells were treated with RIPA buffer. Immunobloting using spectrin antibody (a) confirms that CS or PD inhibits calpain activation, and immunoblating using spectrin antibody (a) confirms that CS or PD inhibits calpain activation, and immunoblating using spectrin antibody (a) shows that neither CS nor PD protected C8 cells from CPT-triggered cell death. (**F**) Roscovitine does not protect cells with either wild-type (C8) or p53 knockout (A9) from CPT-induced cell death. C8 and A9 cells were treated with CPT (20 μ M) in the presence of roscovitine (10 or 20 μ M) for 18 h. Cell death was measured by trypan blue staining

conducted on near-serial sections of ED13.5 caspase-3 null mouse embryos. TUNEL assay showed many dead cells containing fragmented DNA in the trigeminal region (Figure 4A, left panel, black arrows). This suggested that some cells died in a caspase-3-independent manner, but were still TUNEL positive. Cdk5 protein (active Cdk5) was also expressed in many cells in the same area on an adjacent section (Figure 4A, right panel, black arrows), indicating caspase-3-independent Cdk5 expression and activation in dying cells. In addition to the dying trigeminal cells, we also found cell death in the liver, dorsal root ganglia, and other regions (data not shown). To further confirm the *in vivo* data, we used an *in vitro* system, in which both wild-type and caspase-3–/– primary mouse embryonic fibroblast cells were exposed to CHX ($25 \,\mu$ g/ml) or CPT ($20 \,\mu$ M) for 24 h. Cells lacking caspase-3 were more resistant to induction of cell death by CHX or CPT. In all, 27 and 35% of caspase-3–/– cells were killed by CHX and CPT, respectively, in contrast to 87% cells killed by CHX and 63% by CPT in wild-type cells (Figure 4B). The level of Cdk5 expression remained constant in both types of cells (Figure 4C, a). However, p25 was generated only when cell death was induced, and its level corresponded to the level of cell death (Figure 4C, b).

To further verify whether Cdk5 was activated during cell death induced in caspase 3–/– cells, we performed Cdk5 kinase assay using histone H1 as an *in vitro* substrate. The phosphorylated histone H1 band at 33 kDa indicated activation of Cdk5 in both wild-type and caspase-3–/– cells induced to die by CHX and CPT. The intensity of the band corresponded to that of the p25 band and to the level of induced cell death (Figure 4C, c).

Thus, caspase-3 is not required for p25-mediated Cdk5 activation during cell death.

Cdk5 is not necessary for cell death in non-neuronal C8 cells

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Having established that apoptotic cell death could be induced in the MEF C8 cells, we used these cells to examine the activation of Cdk5 activity and its regulation. The level of Cdk5 protein was equal in both control and cells exposed to the three toxins (Figure 5A), as we had seen in dying cells of developing mouse embryos.^{24,28} However, we found little, if any, activity of Cdk5 kinase in the control, but substantial activity in all exposed samples (Figure 5B, a). The level of Cdk5 kinase activity quantified by densitometry (Figure 5B, b) approximated the level of cell death (Figure 5A).

p25, presumptively generated from p35, is an activator of Cdk5, is involved in neuronal cell death, and is the only Cdk5 interacting protein identified in dying cells.²⁸ Here we examined the generation of p25 in non-neuronal cells in relation to cell death induced by the different reagents. By WB, p25 induction was seen for all the three treatments (Figure 5C, a) and the level, as quantified by densitometry (Figure 5C, b), approximated that of cell death (Figure 1A) and correlated with Cdk5 activity, as measured by the histone 1 kinase assay (Figure 5B).

Calpain is the only protease identified as cleaving p35 to form p25. Calpain activation can be determined by the cleavage of its endogenous substrate spectrin to form specific spectrin breakdown products (SBDPs) at 150 and 145 kDa.⁶¹ In exposed C8 cells, anti-spectrin antibody detected a dual band at 150 and 145 kDa, indicating calpain activation (Figure 5D).

The level of calpain activation (Figure 5D) corresponded to the level of p25 generated (Figure 5C). The relationship between calpain activation and generation of p25 was further demonstrated by the use of two specific calpain inhibitors. calpastatin (CS) peptide (an endogenous inhibitor of calpain) and PD150606 (PD, a synthetic inhibitor). In C8 cells exposed to CPT in the presence of either CS or PD, SBDP bands representing calpain activation were not present as expected (Figure 5E, a), and the p25 band failed to appear as well (Figure 5E, b and c). Similar results were observed in C8 cells treated with EtOH and CHX (data not shown). Since we and many others have demonstrated that p35/p25 acts directly upstream of Cdk5 activation and p25 production is a reasonable measure of Cdk5 activation due to its more accessible measurement, these results document that Cdk5 activation is induced by calpainmediated p25 production during cell death in cells other than neurons.

However, although inhibition of calpain activity by either CS or PD abolished Cdk5 activation, it failed to protect cells from death trigged by CPT. In C8, either CS or PD alone did not induce cell death above the basal level in untreated control cells, and neither inhibitor significantly protected C8 from CPT-induced apoptosis (Figure 5D, d). To more directly link Cdk5 activation and apoptosis, we further exposed both C8 and A9 (p53–/–) cells to CPT in the presence or absence of roscovitine, a widely used Cdk5 inhibitor. At low concentrations as we used in our experiment, roscovitine specifically inhibits Cdk5, while sparing other Cdks.⁶² In both C8 and A9, 5–20 μ M rovcovitine did not affect the level of cell death induced by CPT after 18 h (Figure 5F). Therefore, Cdk5 activation does not cause cell death and inhibition of Cdk5 activity will not protect cells.

Discussion

In this paper, we demonstrate that the finding that p25 is generated when Cdk5 is activated in neuronal cells applies also to cells other than neurons, and we extend the finding by dissociating this activation from several well-known pathways of apoptosis. As expected, p53 protects cells at low doses of CHX and CPT, but high levels of these very toxic drugs ultimately lead to cell death although with different kinetics, and this death occurs with Cdk5 activation and p25 generation. These findings corroborate those of Neet et al., who observed that delayed apoptosis occurs even in p53-defective NGF-deprived PC-12 cells.⁶³ In our experiments, the fact that EtOH kills far fewer A9 cells (12% after 18 h compared to 54% of C8) indicates that EtOH lethality in these fibroblasts is dependent on p53. However, whether or not cell death requires a functional p53 pathway, the Cdk5/p25 activation correlates with the induction of cell death.

Cdk5 is activated even when the intrinsic pathway is not. The formation of an active apoptosome (cytochrome c/Apaf-1) and consequent caspase-9 and 3 activation define the intrinsic cell death pathway.⁶⁴ Here we demonstrate that Cdk5 is activated when cells die in the absence of Apaf-1, caspase-9, or -3. Cdk5 is also activated in CPT-induced cell death in C8 (wild-type) fibroblasts in the presence of the pan-caspase inhibitor zVAD-FMK (50 μ M), further documenting a general correlation between Cdk5 activation and cell death, regardless of caspase activation.

Since caspase activity is necessary for the classical characteristics of apoptosis, such as internucleosomal DNA fragmentation,⁶⁵ the deaths in the Apaf-1, caspase-9, -3 knockouts, and in the presence of zVAD-FMK are presumptively nonapoptotic or represent a variation of apoptosis. If so, we have demonstrated a correlation between Cdk5 activation and both classical apoptotic and nonapoptotic or nonclassical apoptotic cell death. Caspase-3 activation, when present, coincides with the activation of Cdk5, but lack of caspase-3 does not hinder the cell death-related activation of Cdk5, indicating that Cdk5 activity is independent of caspase-3 during cell death. Although we did not directly test the correlation with caspase-3, inhibition of Cdk5 does not delay cell death. Sandal *et al.*⁶⁶ argued that Cdk5 activation is located upstream of caspase-3 activation and that Cdk5

activity is needed for cleavage of pro-enzyme caspase-3 to its active (17 kDa) form, in cAMP-induced apoptosis in rat leukemia cells. However, Li *et al.*⁴¹ demonstrated caspase-3 activation in the brain of Cdk5 null mouse embryos during cell death, indicating Cdk5-independent caspase-3 activation in neuronal cell death. Our results are more consistent with those of Li *et al.* The discrepancy may perhaps be ascribed to the different systems utilized.

The activation and generation of p25 are coincident with the activation of calpain. However, although blockage of calpain using either CS or PD prevents activation of Cdk5 (Figure 5E, b and c), it does not reduce the level of cell death (Figure 5E, d). Furthermore, we have conducted studies where inhibition of Cdk5 activity using either pharmacological inhibitors, such as roscovitine, or dominant-negative mutants cannot protect cells from cell death in which Cdk5 is usually activated (Figure 5F; DN, data not shown). Therefore, our data suggest that Cdk5/p25 activation is a consequence rather than a cause of cell death in neuronal and non-neuronal cells. That neurons from Cdk5 knockout mice can die via apoptosis supports our finding.⁴¹

How Cdk5 is activated seems to vary according to the experimental paradigm. It has been shown that Cdk7, purified from bovine brain, is a Cdk5-activating kinase that phosphorylates Cdk5 at Ser159 and enhances Cdk5/p25 activity.⁶⁷ Cdk5 can also be activated without p25 in rat cerebellar granule neurons following glutathione depletion.⁴² Our data indicate activation by p25 in these mesodermal cells. The fact that Cdk5 protein does not change in either treated or untreated C8 and A9 cells or other cells with specific knockouts supports the argument that Cdk5 is a relatively stable protein and that its activation during cell death is not regulated by synthesis or degradation of protein.²⁸

Cdk5 may mediate neuronal apoptosis through two distinct mechanisms, a cytoplasmic mechanism that leads to cytoskeleton disruption⁶ and a nuclear mechanism that interferes with the transcription of prosurvival genes.⁶⁸ Since the presence or absence of active Cdk5 does not appear to affect apoptosis or nonapoptotic cell death, the significance of these findings is unclear.

We conclude that Cdk5 protein is present in many, if not all, types of cells *in vivo* and *in vitro*, and that this protein is activated upon induction of cell death. Activation of Cdk5 is dependent on generation of p25 from p35 by means of activated calpain, and is independent of p53, apoptosome, caspase-3, and -9. However, a cell can also die without activation of Cdk5 by p25 generation, thus producing a death either independent of Cdk5 or relying on activation of Cdk5 by unidentified factors. We are examining the latter possibility, as we continue to explore the significance of Cdk5 activation in cell death.

Materials and Methods

Antibodies and reagents

Chemical reagents were ordered from Sigma (St Louis, MO, USA), unless otherwise mentioned. The calpain inhibitors, CS peptide (Cat# 208902) and PD (Cat# 513022), were obtained from CalBiochem (La Jolla, CA, USA). The antibodies used in this paper include anti-Cdk5 (C-8), anti-p35

(C-19), and anti-PARP (H-250) antibodies from Santa Cruz Biotechnology (Santa Cruz, CA, USA), anti-active caspase-3 antibody from BD Biosciences (San Diego, CA, USA), and anti-spectrin antibody from Chemicon International (Temecula, CA, USA).

Cell culture and treatment

C8 and A9 cells are mouse embryonic fibroblast cell lines transformed with E1A and ras, with C8 p53 + /+ and A9 p53-/- (gift from Dr. Scott Lowe, CSHL). Cells were maintained in Dulbecco's modified Eagle medium (DMEM, Life Technologies, Rockville, MD, USA) supplemented with 10% fetal bovine serum, 50 U/ml penicillin, and 100 μ g/ml streptomycin, at 37°C, in a humidified atmosphere of 7.5% CO₂. At 80% confluence, cells were exposed in DMEM containing 1% FBS to 2.5% EtOH, 25 μ g/ml cycloheximide (CHX, dissolved in EtOH), or 20 μ M CPT (dissolved in DMSO), for 18 h. For the kinetic study of cells exposed to 20 μ M CPT, cells were collected at 0, 3, 6, 8, 12, 15, and 18 h. To block calpain activation induced by CPT in both C8 and A9 cells, two calpain inhibitors, 10 μ M CS or 10 μ M PD, were coadministered with CPT and cells were incubated for 18 h. To inhibit Cdk5 activation, cells were treated with roscovitine alone or in combination with CPT for 18 h.

Cell death by trypan blue

Dead and dying cells lose membrane integrity, allowing the preferential uptake of vital dye trypan blue.⁶⁹ Following exposure to toxins, cells were pelleted and washed with PBS. In all, 100 μ l of well-suspended cells was mixed with 100 μ l of 0.4% trypan blue solution in phosphate-buffered saline (PBS, 145.4 mM NaCl, 2.68 mM KCl, 10 mM Na₂HPO₄, 1.8 mM KH₂PO₄, pH 7.4), and left at room temperature (RT) for 5 min. Cells were then viewed under a light microscope; blue cells were considered nonviable.

Detection of DNA fragmentation by agarose gel electrophoresis

Cells were lysed in lysis buffer (0.2% Triton X-100, 10 mM Tris/HCl 10 mM EDTA, pH 7.5). The cell lysate was incubated on ice for 15 min, followed by centrifuging at 4°C at 13 000 rpm for 30 min. The supernatant containing the low-molecular-mass DNA was incubated with RNase (100 μ g/ml) for 1 h and then extracted twice with equal volumes of phenol/ chloroform/isoamyl alcohol (24 : 24 : 1) and once with chloroform/isoamyl alcohol (24 : 24 : 1) and once with chloroform/isoamyl alcohol (24 : 300 mM NaCl and 2.5 volume of EtOH at -20° C overnight and spun at 13 000 rpm for 30 min at 4°C. The pellet was washed with 70% EtOH once, air dried, and suspended in 10 mM Tris/HCl, 1 mM EDTA, pH 8.0. Low-molecular-mass DNA from equal numbers of cells was run on 2% agarose gel. Separated DNA bands were visualized by ethidium bromide staining of the gel.

Animals, tissue, and slide preparation

Mice (p53 heterozygous mice C57BL/6J-Trp53^{tmlTy} (Jackson Laboratories, Bar Harbor, ME, USA) and caspase-3 heterozygous mice (gift from Dr. Flavell, Yale University)) were mated overnight, and early next morning females were checked for vaginal plugs. Positive females were designated as gestational day 0.5. On gestational day 13.5, females were killed by cervical dislocation and the embryos were removed. Freshly dissected tissue samples were fixed in 4% paraformaldehyde by shaking at 4°C

overnight, then washed with PBS and partially dehydrated by immersing in 20% sucrose at 4°C by shaking for 24 h. Embryos were then embedded in OTC compound (Miles, Elkhart, IN, USA) for frozen blocks, which can be stored at -80° C and sections were made on Vectabond[®] slides.

PCR genotyping of p53 knockout mice

Genotyping of embryos was performed according to the protocol recommended by Jackson Laboratories (http://www.jax.org/resources/ documents/imr/supp proto.html). Two sets of primers were used for p53: 5'-CTT GGG TGG AGA GGC TAT TC-3' and 5'-AGG TGA GAT GAC AGG AGA TC-3': 5'-ATA GGT CGG CGG TTC AT-3' and 5'-CCC GAG TAT CTG GAA GAC AG-3'. Three primers were used for caspase-3: 5'-CTT GGG TGG AGA GGC TAT TC-3' and 5'-AGG TGA GAT GAC AGG AGA TC-3'; 5'-ATA GGT CGG CGG TTC AT-3' and 5'-CCC GAG TAT CTG GAA GAC AG-3'. Briefly, the hind limb was lysed in 170 μ l of lysis solution (50 mM Tris, pH 8.3, 100 mM NaCl, 5 mM EDTA, 0.8% SDS) and 30 µl of Proteinase K solution (10 mg/ml) at 55°C for 5 h or overnight. After centrifuging at 13 000 rpm for 15 min, the supernatant containing DNA was precipitated using the same volume of 100% isopropyl alcohol. The DNA pellet obtained by centrifuging at 13 000 rpm for 15 min was dissolved in autoclaved ddH₂O and stored at -20°C. In all, 10-200 ng of the extracted DNA was used for PCR. This PCR assay applied the 'touching down cycling' technique. For the first 12 cycles, the annealing temperature, which started at 64°C, was reduced by 0.5° per cycle. For the remaining 25 cycles, the annealing temperature stayed at 58°C. The PCR products were separated by gel electrophoresis on a 1.5% agarose gel. P53 or caspase-3 wild type gave a band at 600 or 320 bp, and their knockout counterparts showed a band at 280 or 300 bp, respectively.

DNA fragmentation assay by TUNEL

DNA fragmentation was detected by using TUNEL peroxidase (TUNEL POD) combined with nonisotopic digoxigenin-11-dUTP and terminal transferase according to our previously published method²⁴ and the manufacturer's instruction (Roche Molecular Biochemicals, Germany). Briefly, slides were incubated with permeabilization solution (0.1% Triton X-100 in 0.1% sodium citrate) on ice for 2 min, and endogenous peroxidase activity was guenched with 0.3% hydrogen peroxide in methanol at RT for 30 min. After two rinses in PBS, TUNEL reaction mixture (9 volume of TUNEL label and 1 volume of TUNEL enzyme) was applied to slides, which were incubated for 30 min at 37°C followed by three washes with PBS. TUNEL POD was then applied to the slides to bind to the FITC-dUTP enzymatically added to the free end of the oligonucleotide and visualized with DAB (diaminobenzidine, Research Genetics. Huntsville, AL, USA). The sections were counterstained with methylene blue and mounted with Permount[®] (Fisher Scientific, Burr Ridge, IL, USA).

Immunohistochemistry

In situ levels of specific proteins were estimated using the Avidin-Biotinylated-peroxidase Complex (ABC) kit (Vectastain ABC kit, Vector Laboratory, Burlingame, CA, USA). Sections were quenched with 0.3% hydrogen peroxide in methanol at RT for 20 min to abolish endogenous peroxidase activity. After three washes with 1 × PBST (0.1% Tween 20 in PBS), sections were incubated in blocking solution at RT for 1 h, and treated with primary antibody (0.5 μ g/ml for Cdk5 and p35, 0.1 μ g/ml for caspase-3) (Santa Cruz Biotechnology, Santa Cruz, CA, USA) at 4°C in a humidified chamber overnight. Following three washes with PBST, secondary biotinylated antibody was applied to the sections for 1 h at RT. Sections were then washed with PBST for three times and incubated with ABC reagent for 2 h at RT. Sections were washed with PBST for three times before being visualized with DAB, counterstained with methylene blue, and mounted with Permount.

Western blot

Cells or tissues were lysed in RIPA buffer (50 mM Tris, pH 7.5, 150 mM NaCl, 1% Triton X-100, 0.1% SDS, 0.5% sodium deoxycholate, 2 mM EDTA) supplemented fresh with protease inhibitor cocktail tablets (Boehringer Mannheim, Germany), and cell/tissue debris were removed by centrifugation at 13 000 rpm for 30 min at 4°C. Protein concentration was determined using the Bio-Rad microassay (Bio-Rad laboratories, Hercules, CA, USA). Equal amounts of protein were run on 8, 12.5 or 15% SDS-polyacrylamide gels after addition of equal volumes of 2 × Laemmli loading buffer (100 mM Tris, pH 7.5, 4% SDS, 20% glycerol, 0.002% bromophenol blue).⁷⁰ After protein transfer, nitrocellulose membrane blots were blocked with 5% non-fat dry milk in PBST for 30 min at RT and incubated with primary antibody (0.5 μ g/ml for Cdk5 or p35 or caspase-3, 0.1 μ g/ml for spectrin and PARP) on the shaker overnight at 4°C. After three washes with PBST, blots were incubated with horseradish peroxidase (HRP)-conjugated goat anti-rabbit antibody (Jackson Immuno Research Laboratory, West Grove, PA, USA) for 1 h at RT, followed by three washes. The immune complexes were detected by chemiluminescence (ECL Plus kit, Amersham, Chicago, IL, USA) and exposed to an autoradiographic film.

In vitro histone H1 kinase assays

Cdk5 kinase activity was assayed as described by Zhang et al.⁵ Equal amounts of lysates from different cell samples were incubated with 1.5 μ g/ ml Cdk5 antibody for 1 h at 4°C, and then purified by the addition of protein A-Sepharose (Boehringer Mannheim, Germany). The precipitated beads were equilibrated in kinase buffer (50 mM Tris, pH 7.5, 10 mM MgCl₂, 1 mM DTT, 20 mM EGTA, 0.1 mM sodium vanadate, 80 mM ßglycerophosphate) and collected by centrifugation. Histone H1 kinase assays were performed in kinase buffer supplemented with 0.1 μ Ci/ml [y-32P]ATP (6000 Ci/mM) (Amersham Pharmacia Biotech., Piscataway, NJ, USA), 10 mM ATP, 50 µg/ml calf thymus histone H1 (Boehringer Mannheim, Germany), and 5 µM cAMP-dependent protein kinase inhibitor (Sigma, St Louis, MO, USA) at 30°C for 30 min. An equal volume of 2 imesLaemmli buffer was added to each sample before they were denatured at 100°C for 2 min. Following electrophoresis, gels were fixed, dried, and exposed to film. The intensity of the bands was quantified using a Storm Imaging System (Molecular Dynamics Inc., Sunnyvale, CA, USA).

Condensed nuclei stained by bis-benzimide

The DNA fluorochrome bis-benzimide (Hoechst 33258, Sigma, St Louis, MO, USA) was used to stain the fragmented nuclei of apoptotic cells. Briefly, cells were trypsinized, pelleted, and washed once with ice-cold PBS. The cells were then resuspended and fixed in 3% paraformaldehyde solution (in PBS) for 30 min, washed with ice-cold PBS, and incubated with 16 μ g/ml bis-benzimide (in PBS) for 25 min at RT. Finally, the cells were washed with ice-cold PBS once and resuspended in PBS. A volume of 20 μ l of the cell suspension was evenly distributed on a slide and coverslipped to be viewed under an Eclipse TE300 microscope (Nikon Inc., Melville, NY, USA).

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