

Involvement of Glyceraldehyde-3-Phosphate Dehydrogenase (GAPDH) and p53 in Neuronal Apoptosis: Evidence That GAPDH Is Upregulated by p53

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We recently reported that cytosine arabinoside (AraC)-induced apoptosis of cerebellar neurons involves the overexpression of glyceraldehyde-3-phosphate dehydrogenase (GAPDH). The present study was undertaken to investigate whether p53 and/or Bax overexpression participates in the AraC-induced apoptosis of cerebellar granule cells and, if so, the relationship between p53 induction and GAPDH overexpression in these cells.

AraC-induced apoptosis of cerebellar granule cells was preceded by an increase in levels of p53 mRNA and protein detected between 1 and 8 hr after treatment. The mRNA level for a p53 target gene, Bax, was also increased. The increase in GAPDH mRNA lasted longer than that of either p53 or Bax, and the level of GAPDH protein in the particulate fraction increased after induction of GAPDH mRNA. The antisense oligonucleotide to p53 protected granule cells from AraC-induced chromatin condensation, internucleosomal cleavage, and apoptotic death. The inhibition of p53 expression by the p53 antisense

oligonucleotide not only blocked the expression of Bax but also partially suppressed the increased GAPDH mRNA and protein levels. Conversely, the suppression of GAPDH expression and subsequent attenuation of apoptosis of granule cells by GAPDH antisense oligonucleotide did not influence the expression of p53 or Bax. Cerebellar granule cells prepared from p53 knock-out mice were resistant to AraC toxicity, and the p53 gene knock-out suppressed AraC-upregulated GAPDH expression. Moreover, infection of PC12 cells with an adenoviral vector containing p53 gene dramatically increased GAPDH expression and triggered cell apoptosis. These results suggest that AraC-induced apoptosis of cerebellar granule cells involves the expression of both GAPDH and p53 and that, similar to Bax, GAPDH is upregulated by p53 after exposure to the apoptotic insult.

Key words: p53; GAPDH; cerebellar granule cell; PC12; cytosine arabinoside; apoptosis; adenovirus

Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) is an essential glycolytic enzyme that is expressed in all prokaryotic and eukaryotic organisms. Its role in cellular metabolism is to convert glyceraldehyde-3-phosphate to 1,3-diphosphoglycerate. Recent studies suggest that GAPDH is a multifunctional protein endowed with a number of diverse activities seemingly independent of its role in glycolysis. These activities include phosphorylating transverse-tubule proteins (Kawamoto et al., 1986), stimulating RNA transcription (Morgenegg et al., 1986), interacting with microtubules (Huitorel and Pantaloni, 1985), influencing RNA catalysis by binding to hammerhead ribozyme (Sioud and Jespersen, 1996), and acting as a diadenosine tetraphosphate binding protein to influence DNA replication and DNA repair (Baxi and Vishwanatha, 1995).

Cytosine arabinoside (1- β -D-arabinofuranosyl cytosine; AraC) is usually thought to kill proliferating cells through incorporation into DNA during replication, followed by chain termination (Hedley and McCulloch, 1996). However, we and others have

reported that AraC induces apoptosis of postmitotic cerebellar granular cells cultured from postnatal rats (Dessi et al., 1995; Ishitani and Chuang, 1996; Saunders et al., 1997, 1999). AraC has been shown to induce apoptosis cerebellar granule cells *in vitro* in a concentration-dependent manner only when added within 24 hr after cell plating (Dessi et al., 1995). Previously, we found that AraC-induced apoptosis of granule cells can be blocked by inhibitors of transcription, translation, and deoxyribonuclease (Ishitani and Chuang, 1996). The AraC-induced apoptosis of granule cells is preceded by an increase in the levels of GAPDH mRNA and protein (Ishitani and Chuang, 1996), which is followed by nuclear translocation of the GAPDH protein (Saunders et al., 1997; Ishitani et al., 1998). Moreover, an antisense oligonucleotide against GAPDH attenuates GAPDH overexpression and nuclear translocation and robustly protects granule cells from apoptosis (Ishitani and Chuang, 1996; Saunders et al., 1997, 1999; Ishitani et al., 1998), suggesting that GAPDH plays a prominent role in AraC-induced apoptotic death. It has also been shown that GAPDH nuclear translocation is involved in apoptosis of non-neuronal and neuronal cells (Sawa et al., 1997). The role of GAPDH in the pathway and the events leading to its induction, however, are unknown.

Accumulating evidence supports the view that p53, a nuclear phosphoprotein, may be necessary for neuronal apoptosis. For example, cerebellar granule cells cultured from p53 null mice are resistant to treatments with excitotoxins (Xiang et al., 1996) or

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DNA-damaging agents, such as irradiation or AraC (Enokido et al., 1996a,b). Moreover, whole body γ -irradiation induces apoptosis of granule cells in wild-type, but not p53-deficient, neonatal mice (Wood and Youle, 1995). Other *in vivo* studies have shown that knocking out the p53 gene protects hippocampal neurons from seizure-induced cell death (Morrison et al., 1996), dopamine neurons from *N*-methyl-4-phenyl-1,2,3,6-tetrahydropyridine-induced toxicity (Trimmer et al., 1996) and the brain from ischemia-induced infarction (Crumrine et al., 1994). Conversely, adenovirus-mediated transfection of the p53 gene in hippocampal or cortical neurons results in the apoptotic death of these neurons (Xiang et al., 1996; Jordan et al., 1997). In view of the emerging evidence suggesting a prominent role for p53 in neuronal apoptosis, this study examined whether p53 and its responsive gene, Bax, are overexpressed during AraC-induced apoptosis of granule cells. In addition, we explored whether GAPDH overexpression is regulated by p53 in granule cells treated with AraC and in other neurally related cell types.

MATERIALS AND METHODS

Primary cultures of cerebellar granule cells and treatment conditions. Cerebellar granule cells were prepared from 8-d-old Sprague Dawley rat pups or 7-d-old wild-type and p53 knock-out mouse pups (Donehower et al., 1992) supplied by Taconic Farms (Germantown, NY), as described previously (Nonaka et al., 1998). Briefly, cerebella were chopped into 400 μ m cubes, and the cells were dissociated by trypsinization, followed by DNase treatment. The dissociated cells were resuspended in a basal modified Eagle's medium containing 10% fetal calf serum, 2 mM glutamate, 50 μ g/ml gentamicin, and 25 mM KCl. Granule cells from rat were seeded at a density of 2.5×10^5 cells/cm² onto 24-well plates or 100 mm culture dishes and granule cells from mouse onto 96-well plates or 60 mm dishes precoated with poly-L-lysine. To induce cell apoptosis, AraC was added to culture medium to a final concentration of 300 μ M 12 hr after cell plating. When used, oligonucleotides were added to the cultures 2 hr before the addition of AraC. After 24 hr of AraC treatment, viability of granule cells grown on plates was measured by MTT assays as described previously (Nonaka et al., 1998). The tetrazolium ring of MTT is reduced by active dehydrogenase in viable mitochondria, forming a blue-colored precipitate that was then dissolved in dimethyl sulfoxide and quantified spectrophotometrically at 540 nm.

Isolation of fragmented DNA. For DNA fragmentation analysis, cells grown on 100 mm dishes were also harvested after treatment essentially as described by Nonaka et al. (1998). Briefly, cells were washed with ice-cold PBS and then lysed in a buffer containing 10 mM Tris-HCl, pH 7.4, 10 mM EDTA, and 0.5% Triton X-100. Lysates were centrifuged at $10,000 \times g$ for 10 min at room temperature. Fragmented DNA in the supernatant was precipitated by the addition of 400 μ l of NaI (7.4 M) and 700 μ l of isopropanol and then isolated by centrifugation at $12,000 \times g$ for 15 min. The DNA pellet was dissolved in 10 mM Tris-HCl, pH 7.4, containing 1 mM EDTA [Tris-EDTA (TE)]. The DNA samples were subjected to electrophoresis on 1.2% agarose gel, and the DNA bands were visualized by ethidium bromide staining.

Northern blotting. Cerebellar granule cells were harvested after different time lengths of AraC treatment. Total RNA was extracted from cells with guanidine thiocyanate and isolated by cesium chloride gradient centrifugation as described previously (Fukamauchi et al., 1993), with modifications. After electrophoresis on 1% agarose gel containing formaldehyde, RNA was transferred to a Duralose membrane (Stratagene, La Jolla, CA) and hybridized with cDNA probes for Bax, Bcl-2, p53, and GAPDH, separately. The cDNA probes were labeled with [α -³²P]dCTP by the random priming method. Hybridizations were performed at 42°C for 16 hr, followed by two washes at room temperature with $2 \times$ SSC-0.1% SDS and then two additional washes at 50°C in $0.1 \times$ SSC-0.1% SDS. Northern blots were quantified by Betascope Model 603 blot analyzer (Betagen, Waltham, MA).

Western blotting. To measure p53 and Bax protein levels, cells cultured in 60 or 100 mm dishes at indicated times were detached by scraping and sonicated 30 sec in Laemmli lysis buffer containing 6.25 mM Tris-HCl, pH 6.8, 2 mM EDTA, 15% sucrose, 10% glycerol, 3% SDS, and 0.7 M β -mercaptoethanol. For measurement of GAPDH immunoreactive protein, the samples were prepared as reported previously (Ishitani and

Chuang, 1996), with slight modifications. Briefly, scraped cells were ruptured by sonication in 0.32 M sucrose, and the homogenates were centrifuged at $2 \times 10^5 g$ for 30 min. The particulate fraction (pellet) was resuspended in a small volume of lysis buffer. Protein concentration was determined by using the BCA protein assay kit (Pierce, Rockford, IL), and samples containing 10–30 μ g of protein were loaded and separated by 12% SDS-PAGE. Proteins were subsequently transferred to an Immobilon-P membrane (Millipore, Bedford, MA). After blocking for 1 hr in PBST ($1 \times$ PBS and 0.1% Tween 20) containing 5% nonfat dry milk, blots were incubated for 1 hr at room temperature with antibodies against GAPDH (Advanced Immunochemical, Long Beach, CA), rat p53 (Pab240), human p53 (DO-1) (Santa Cruz Biotechnology, Santa Cruz, CA), or Bax (Ab-1; Oncogene Research, Cambridge, MA) in PBST containing 3% nonfat milk. Blots were washed four times in PBST during a 40 min period and were then incubated 1 hr with horseradish peroxidase-conjugated second antibodies in PBST containing 3% nonfat dry milk. Immunoreactivities of the protein bands were detected by enhanced chemiluminescent autoradiography (ECL kit; Amersham Pharmacia Biotech, Arlington Heights, IL) as instructed by the manufacturer. Western blots were quantitatively analyzed by capturing images on films using a CCD camera (Sierra Scientific, Sunnyvale, CA) and the Macintosh NIH Image 1.5 software (Wayne Rasband, National Institute of Mental Health, Bethesda, MD).

Morphological evaluation. Changes in nuclear morphology were examined by staining cerebellar granule cells with Hoechst 33258 (Nonaka et al., 1998). Cells growing on 35 mm dishes were fixed with 4% formaldehyde. After gentle washing with cold PBS, cells were stained with

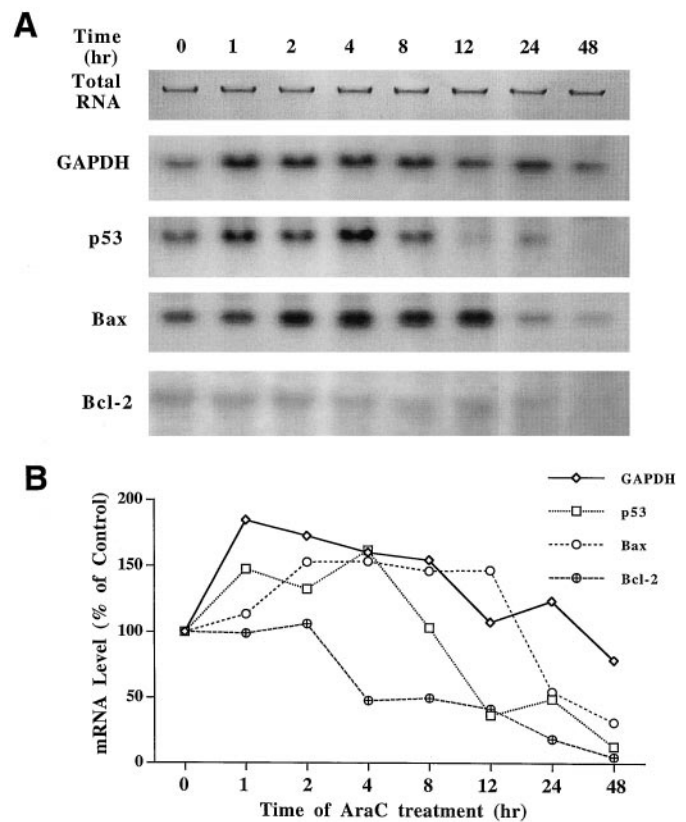


Figure 1. Time course of AraC-induced changes in the mRNA levels of GAPDH, p53, Bax, and Bcl-2 in cerebellar granule cells. AraC was added to culture medium to a final concentration of 300 μ M 12 hr after plating. At the indicated times after AraC exposure, cells were harvested for RNA isolation. An aliquot of $1/10$ of each sample was used for quantification of total cellular RNA. The RNA was allowed to migrate ~ 5 mm from the sample well into 1% agarose gel, and the total RNA was quantified by image analysis (Fukamauchi et al., 1993). The same amount (10 μ g) of total RNA from each sample was applied to each well of the agarose gel for Northern blot analysis. *A*, Northern blots; *B*, quantified results of blots shown in *A*.

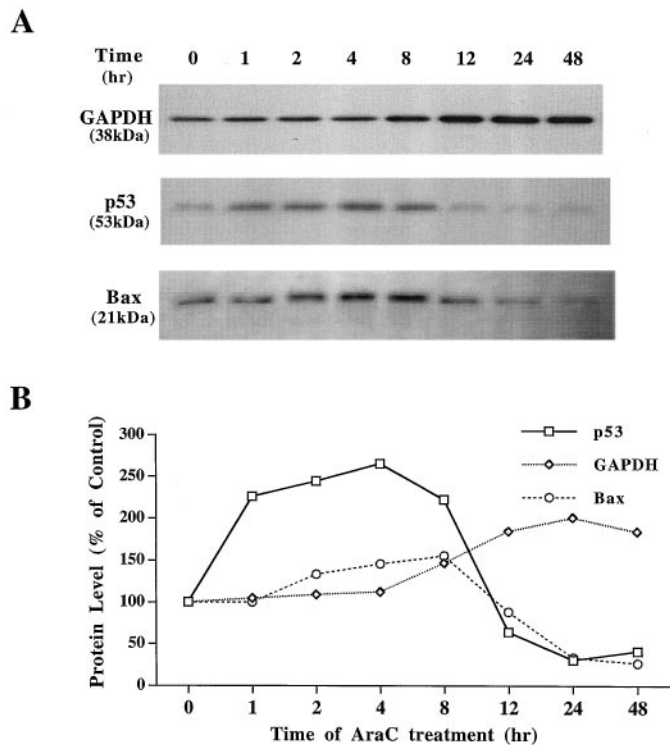


Figure 2. Time course of AraC-induced increases in GAPDH, p53, and Bax protein levels in cerebellar granule cells. Freshly cultured cells were treated with AraC as described in Figure 1. At different time points, cells were collected in lysis buffer and sonicated for 30 sec. For measurement of GAPDH protein, an aliquot containing 10 μ g of protein from the particulate fraction of the cells was loaded and separated by SDS-PAGE. For examination of p53 and Bax protein expression, an aliquot containing 30 μ g of total protein was used for SDS-PAGE. *A*, Western blots; *B*, quantified results of blots shown in *A*. Note that protein levels of GAPDH, p53, and Bax were time dependently changed after AraC treatment.

Hoechst 33258 (5 μ g/ml in PBS) for 5 min at 4°C. Photomicrographs were taken using an inverted UV illumination microscope.

Oligonucleotide preparations. Antisense and sense oligonucleotides to GAPDH and p53 were synthesized by Cruachem (Dulles, VA). The first and last bases of each oligonucleotide were phosphorothioated. The GAPDH oligonucleotide sequence (GAPDH/AS) was 5'-GACCTTCAC-CATCTTGICTA-3', corresponding to a sequence of the rat GAPDH gene flanking the mRNA translation initiation region (GenBank GAPDH cDNA clone M17701). The p53 antisense oligonucleotide (p53/AS) sequence was 5'-GACCTCAGGTGGCTCATACGG-3', corresponding to residues between 390 and 410 of rat p53 coding sequence (GenBank p53 cDNA clone X13058). The sequences of the GAPDH sense (GAPDH/S) and p53 sense (p53/S) oligonucleotides were the exact inverse of those of their corresponding antisense oligonucleotides. The stock concentration of oligonucleotides was 1 mM in TE buffer, pH 7.4.

Construction of adenoviral vectors and transfection into PC12 cells. The construction of adenovirus expressing human wild-type p53 cDNA (Ad-WTp53), β -galactosidase gene (Ad β -gal), and adenovirus containing no transgene (AdControl) have been described previously (Katayose et al., 1995). Briefly, the human wild-type p53 gene was cloned to the site downstream of cytomegalovirus promoter and upstream of SV40 polyadenylation sequence in the replication-defective recombinant adenovirus. Recombinant adenoviruses were propagated in human embryonic kidney 293 cells, purified by double cesium chloride gradient centrifugations, titrated, and stored at -70°C.

PC12 cells were maintained in 100 mm dishes containing RPMI medium 1640 plus 12% fetal bovine serum and 6% horse serum. Cells were infected with 20 pfu/cell of recombinant adenoviruses at 24 hr after plating. The infection efficiency of PC12 cells was determined by counting the β -galactosidase-positive cells under a light microscope at 24 hr after infection with Ad β -gal (Li et al., 1998).

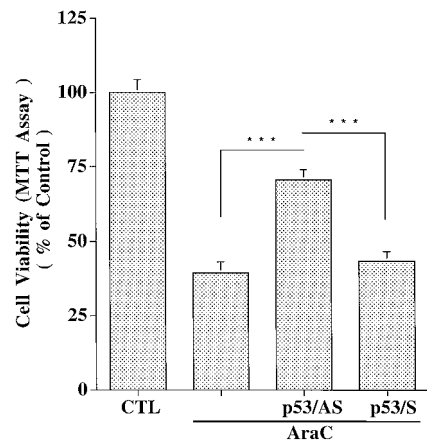


Figure 3. Antisense oligonucleotide to p53 protects against AraC-induced neuronal death. Cerebellar granule cells were pretreated with p53 oligonucleotides (3 μ M) for 2 hr before the addition of AraC (300 μ M). After 24 hr exposure to AraC, neuronal viability was measured using the MTT colorimetric assay. The results are expressed as percentage of mean \pm SEM of control group from three independent experiments. CTL, Control. *** p < 0.001 using one-way ANOVA with Fisher's PLSD test.

RESULTS

Levels of GAPDH, p53, and Bax expression are increased during the AraC-induced apoptosis of cerebellar granule cells

Northern blot hybridization showed that the expression of GAPDH mRNA in freshly cultured granule cells increased from 1 to 24 hr after exposure to AraC (Fig. 1). p53 mRNA levels

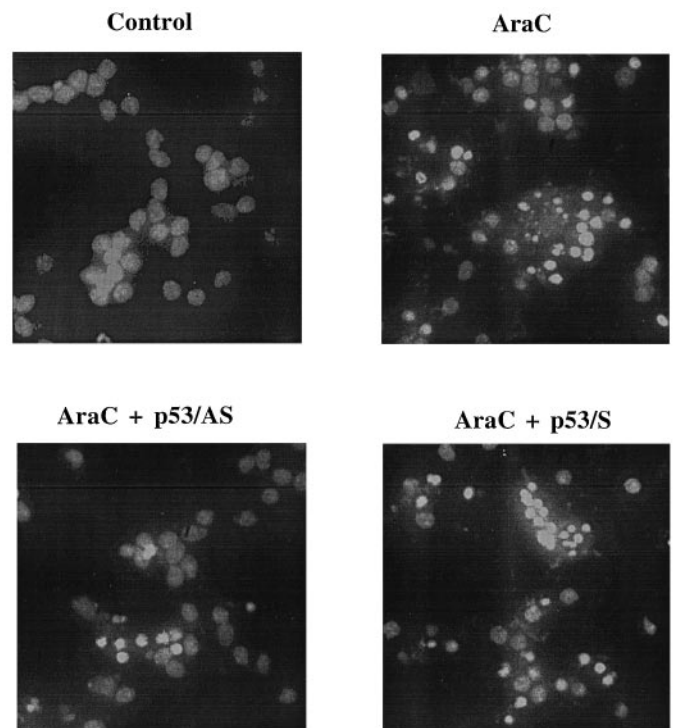


Figure 4. Antisense but not sense oligonucleotide to p53 protects against AraC-induced cell death: morphological studies. Cerebellar granule cells were treated as described in Figure 3. After exposure to AraC for 24 hr, cells were fixed with 4% formaldehyde and then stained with Hoechst dye 33258 (5 μ g/ml in PBS) for 5 min at 4°C. Photomicrographs were taken using an inverted UV illumination microscope.

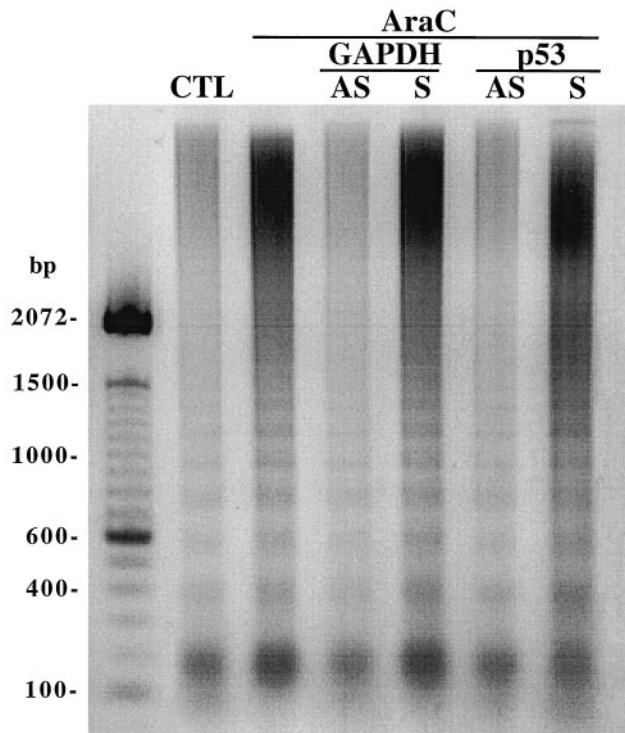


Figure 5. Antisense oligonucleotides to GAPDH and p53 reduce AraC-induced DNA fragmentation. Cerebellar granule cells were treated with AraC as described in Figure 3 and pretreated with GAPDH or p53 oligonucleotides ($3 \mu\text{M}$) for 2 hr. After exposure to AraC for 24 hr, soluble DNA was extracted from cells and subjected to electrophoresis as described in Materials and Methods. The DNA size markers are shown to the left. AS, Antisense oligonucleotide; S, sense oligonucleotide; CTL, control.

increased from 1 to 4 hr and then declined, whereas the Bax mRNA levels were elevated between 2 and 12 hr. Thus, the upregulation of GAPDH mRNA lasted longer than that of either p53 or Bax. The expression of Bcl-2 mRNA was very low and appeared to be decreased after 4 hr of AraC treatment.

Western blots and their quantified results showed that endogenous cellular p53 protein levels were increased at 1 hr, reached their peak (265% of the control) at 4 hr, and then decreased to less than the baseline value by 12 hr (Fig. 2*A,B*). Levels of cellular Bax protein were increased at 2 hr and peaked at ~ 8 hr (155% of the control). Compared with p53 and Bax protein, the GAPDH protein levels in the particulate fraction of AraC-treated granule cells increased rather late; the increase was detected at 8 hr, and reached its maximum (200% of the control) ~ 24 hr after exposure.

p53 and GAPDH antisense oligonucleotides protect cerebellar granule cells against AraC-induced apoptosis

Cell viability, as measured by MTT assay at 24 hr after treatment with AraC, was reduced to $40.4 \pm 3.2\%$ ($n = 4$) (Fig. 3). Photomicrographs showed that AraC caused condensed and/or fragmented cell nuclei, which are characteristic of apoptosis (Fig. 4). Exposure to AraC for 24 hr also caused internucleosomal cleavage of the DNA (Fig. 5). To identify whether the increase in p53 expression was directly related to the cell death, granule cells were pretreated with $3 \mu\text{M}$ p53/AS before exposure to AraC. The cell viability for the p53/AS-pretreated group was $72 \pm 4.0\%$

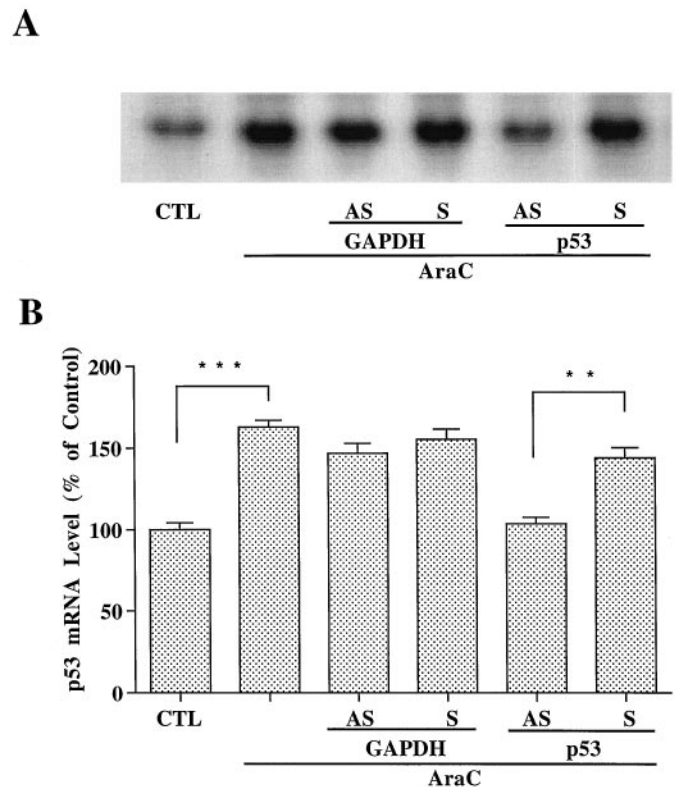


Figure 6. Antisense oligonucleotide to p53 but not GAPDH suppresses AraC-induced increases in p53 mRNA. Cerebellar granule cells were exposed to $300 \mu\text{M}$ AraC for 12 hr after plating. Cells were pretreated with GAPDH or p53 oligonucleotides for 2 hr before the addition of AraC. RNA was extracted from cells after exposure to AraC for 4 hr. *A*, Northern blot results of p53 mRNA levels from a typical experiment. *B*, Levels of p53 mRNA expressed as percent of the control quantitated with a Betascope Model 603 blot analyzer. Each column represents the mean \pm SEM from three independent experiments. AS, Antisense oligonucleotide; S, sense oligonucleotide; CTL, control. $**p < 0.01$; $***p < 0.001$ using one-way ANOVA with Fisher's PLSD test.

the control ($n = 4$), which was significantly greater than that of AraC alone ($p < 0.001$) (Fig. 3). Pretreatment with the corresponding p53/S was ineffective; cell viability was $44.6 \pm 3.3\%$ ($n = 4$; $p > 0.05\%$). p53/AS, but not p53/S, markedly reduced chromatin condensation and nuclear fragmentation (Fig. 4) and essentially blocked internucleosomal DNA cleavage in AraC-treated cells (Fig. 5). These results suggest that AraC-induced apoptosis of granule cells is not only dependent on GAPDH, but on p53 as well. Moreover, similar to the GAPDH/AS (Ishitani and Chuang, 1996; Saunders et al., 1997), the p53/AS was effective in protecting granule cells from AraC-induced apoptosis.

Abrogation of p53 expression suppresses GAPDH mRNA and protein levels

Northern blot results showed that p53/AS, but not p53/S, blocked AraC-induced upregulation of p53 and Bax mRNA, whereas neither GAPDH/AS nor GAPDH/S was effective (Figs. 6, 7). AraC-induced GAPDH mRNA upregulation was blocked by GAPDH/AS and partially suppressed by p53/AS (Fig. 8). p53/AS also concentration dependently suppressed AraC-increased levels of GAPDH protein in the particulate fraction of granule cells (Fig. 9*A,B*). At concentrations of 1 and $3 \mu\text{M}$, p53/AS suppressed upregulated GAPDH protein by 37.6 and 74.7%, respectively, whereas p53/S was ineffective at these two concentrations. At 5

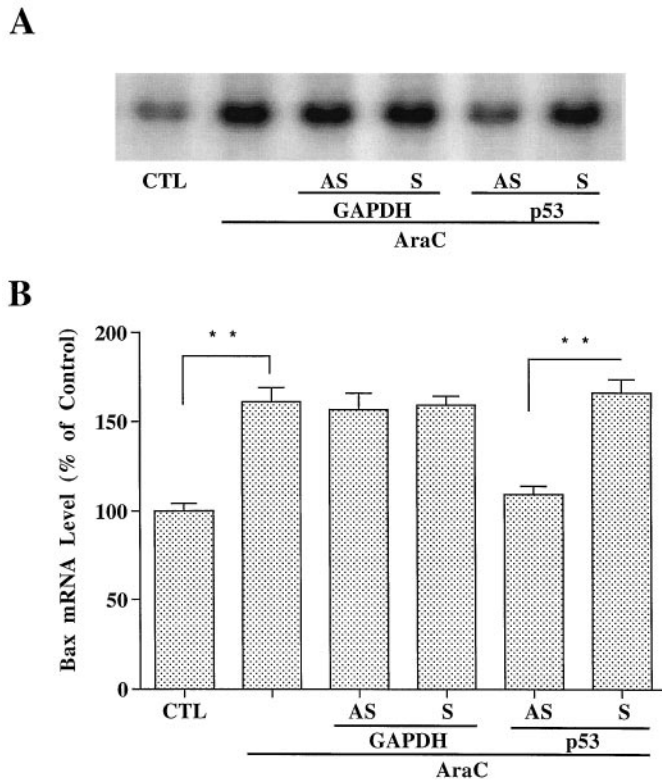


Figure 7. Antisense oligonucleotide to p53 inhibits AraC-induced Bax mRNA upregulation. Cerebellar granule cells were treated with AraC, and RNA was prepared as described in Figure 6. *A*, Northern blot results of Bax mRNA levels from a typical experiment. *B*, Levels of Bax mRNA expressed as percent of the control quantitated with a Betascope Model 603 blot analyzer. Each column represents the mean \pm SEM from three independent experiments. *AS*, Antisense oligonucleotide; *S*, sense oligonucleotide; *CTL*, control. $**p < 0.01$ using one-way ANOVA with Fisher's PLSD test.

μM , p53/AS completely blocked GAPDH protein upregulation, and p53/S was partially effective. Similar dose-dependent neuroprotective effects of p53/AS and p53/S on AraC-induced apoptosis were observed (data not shown). Therefore, oligonucleotide concentrations higher than $3 \mu\text{M}$ were not recommended. These results suggest that, during AraC-induced apoptosis of granule cells, GAPDH may function downstream of p53.

p53 gene knock-out in mice inhibits AraC-induced apoptosis and GAPDH expression in cerebellar granule cells

To further explore the role of p53 in mediating GAPDH induction, we used cerebellar granule cells from p53 knock-out mice. AraC treatment induced a concentration-dependent decrease in the viability of granule cells derived from wild type but had little or no effect in those cells derived from p53 $^{-/-}$ mice (Fig. 10*A*). Thus, after treatment with $1000 \mu\text{M}$ AraC for 24 hr, viabilities of granule cells derived from p53 $^{+/+}$ and p53 $^{-/-}$ mice were 32.1 ± 1.5 and $81.4 \pm 5.8\%$, respectively. These results confirm that p53 gene knock-out confers resistance to AraC-induced apoptotic cell death (Enokido et al., 1996b). Importantly, GAPDH protein levels were markedly elevated by AraC treatment in granule cells from p53 $^{+/+}$ mice but were only marginally increased in those cells from p53 $^{-/-}$ mice (Fig. 10*B*).

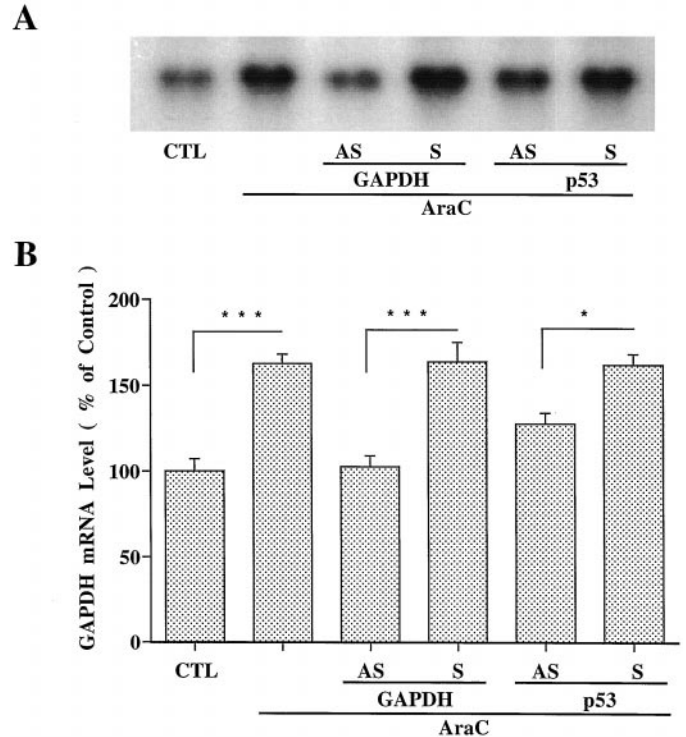


Figure 8. GAPDH and p53 antisense oligonucleotides inhibit AraC-induced GAPDH mRNA increase. Cerebellar granule cells were treated with AraC, and RNA was prepared as described in Figure 6. *A*, Northern blot results of GAPDH mRNA levels from a typical experiment. *B*, Levels of GAPDH mRNA expressed as percent of the control quantitated with a Betascope Model 603 blot analyzer. Each column represents mean \pm SEM from three independent experiments. *AS*, Antisense oligonucleotide; *S*, sense oligonucleotide; *CTL*, control. $*p < 0.05$; $***p < 0.001$ using one-way ANOVA with Fisher's PLSD test.

p53 gene transfection into PC12 cells upregulates GAPDH protein and triggers apoptosis

To obtain additional evidence that GAPDH is regulated by p53, we infected PC12 pheochromocytoma cells with the recombinant adenovirus containing AdWtp53, Ad β -gal, or AdControl. The infection efficiency determined by counting β -galactosidase-positive cells was dependent on the dose of the recombinant adenovirus and found to be $82.9 \pm 4.4\%$ with 20 pfu/cell of Ad β -gal (data not shown). The p53 gene transfection of PC12 cells resulted in a time-dependent increase in the levels of exogenous p53 protein in the cell homogenate (Fig. 11). Thus, the p53 protein levels were progressively increased from 12 to 48 hr after AdWtp53 infection. In contrast, infection with AdControl produced no detectable p53 protein. Interestingly, the levels of GAPDH protein in the particulate fractions were also markedly increased at 24 and 48 hr in cells infected with AdWtp53 but not AdControl. The p53 gene transfer into PC12 cells induced a time-dependent loss of cell viability with a detectable decrease at day 1, followed by a progressive decrease to $<20\%$ of the control at days 3 and 5 (Fig. 12*A*). A dramatic increase in internucleosomal DNA cleavage was also observed at 2 d after AdWtp53 infection (Fig. 12*B*). In contrast, infection with AdControl induced neither cell death nor detectable DNA fragmentation throughout the time course studied.

DISCUSSION

The tumor suppressor p53 serves as a critical regulator in the cell cycle and in the apoptosis of normal cells after exposure to

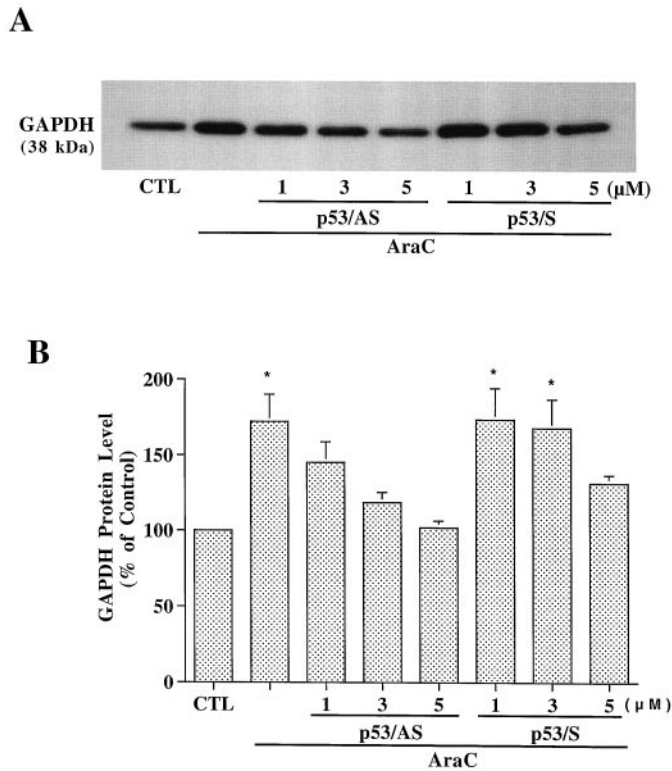


Figure 9. Antisense oligonucleotide to p53 suppresses AraC-induced GAPDH protein increase in the particulate fraction of cerebellar granule cells. Cells were treated with AraC as described in Figure 6. After exposure to AraC for 24 hr, cells were harvested and the particulate (pellet) fraction was prepared as described in Materials and Methods. An aliquot of 10 μ g of protein from each particulate fraction was used for Western blot analysis. *A*, Western blot results of GAPDH protein levels from a typical experiment. *B*, Quantified results of GAPDH protein levels expressed as percent of mean \pm SEM of the untreated control from three independent experiments. CTL, Control. * $p < 0.05$ compared with the untreated control using one-way ANOVA with Fisher's PLSD test.

ionizing radiation or DNA-damaging agents. It has been suggested that p53 may act as a "guardian of the genome." When DNA damage is mild, p53 halts cell cycle progression at the G1 phase through the induction of genes such as p21 and GADD45 (Pellegata et al., 1996) (for review, see Hughes et al., 1997), allowing DNA repair to occur before advancing through the cell cycle. However, when DNA damage is extensive and not repairable, p53 triggers apoptosis (Grasso and Mercer, 1997) (for review, see Yonish-Rouach, 1996; Hughes et al., 1997). These biochemical characteristics ensure that p53 plays a key role in controlling genomic stability (Lowe et al., 1993; Fukasawa et al., 1997). Elevated expression of p53 may initiate apoptosis by increasing the expression of the death gene Bax (Miyashita and Reed, 1995), decreasing the expression of the cytoprotective gene Bcl-2 (Selvakumaran et al., 1994) or inducing the transcription of redox-related genes that cause oxidative stress (Polyak et al., 1997).

When DNA is damaged in postmitotic neurons, p53 induction is associated with mechanisms underlying death rather than recovery (Sakhi et al., 1994; Wood and Youle, 1995; Enokido et al., 1996a; Jordan et al., 1997). Transfection studies have shown that overexpression of p53 per se is sufficient to induce neuronal apoptosis (Xiang et al., 1996; Jordan et al., 1997). In contrast, p53-deficient neurons are resistant to apoptosis induced by exci-

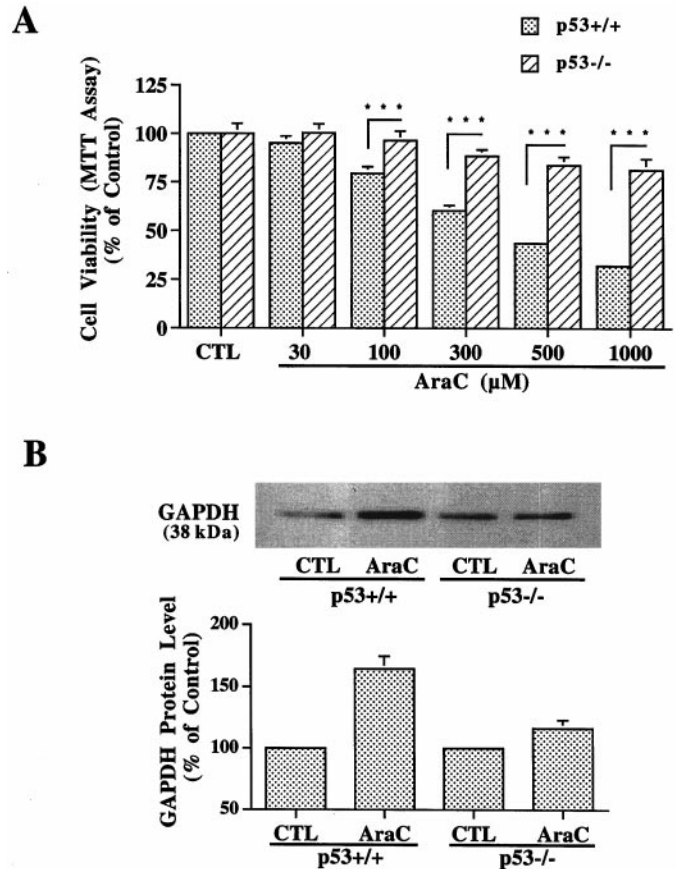


Figure 10. Knock-out of p53 gene suppresses AraC-induced neurotoxicity and GAPDH overexpression in mouse cerebellar granule cells. Granule cells were prepared from the cerebellum of 7-d-old wild-type (p53+/+) and p53 knock-out (p53-/-) mice (Donehower et al., 1992) according to the procedures described in Materials and Methods. *A*, At 12 hr after plating onto 96-well plates, various concentrations of AraC were added to the cultures, and the cell viability was determined 24 hr later by MTT assay. Results shown are mean \pm SEM from six independent experiments. *B*, At 24 hr after plating onto 60 mm dishes, GAPDH protein levels in wild-type and p53-deficient mouse granule cells were determined by Western blot analysis. The top represents Western blot from a typical experiment. The bottom represents quantified results expressed as mean \pm ranges from two independent experiments. Note that p53 gene knock-out resulted in not only inhibition of AraC toxicity but also suppression of AraC-induced GAPDH upregulation. CTL, Control. *** $p < 0.001$ compared with the corresponding results in p53-deficient mouse granule cells, using one-way ANOVA with Fisher's PLSD test.

totoxins such as glutamate (Morrison et al., 1996). The precise mechanism by which p53 modulates neuronal viability has not yet been determined. However, it has been suggested that p53 may function by upregulating Bax and/or by downregulating Bcl-2, whereas generation of the reactive oxygen species and activation of caspases may be further downstream events of p53-dependent apoptosis (Johnson et al., 1996; Lotem and Sachs, 1996).

In this study, we extensively characterized AraC-induced apoptosis with a major emphasis on the role of p53 in the apoptotic process. We found that levels of p53 mRNA and protein in cerebellar granule cells were rapidly and markedly increased after treatment with AraC. Pre-exposure of these cells to p53/AS robustly protected these neurons from AraC-induced chromatin condensation, nuclear fragmentation, and internucleosomal DNA cleavage. Concomitant with these effects, the viability of AraC-

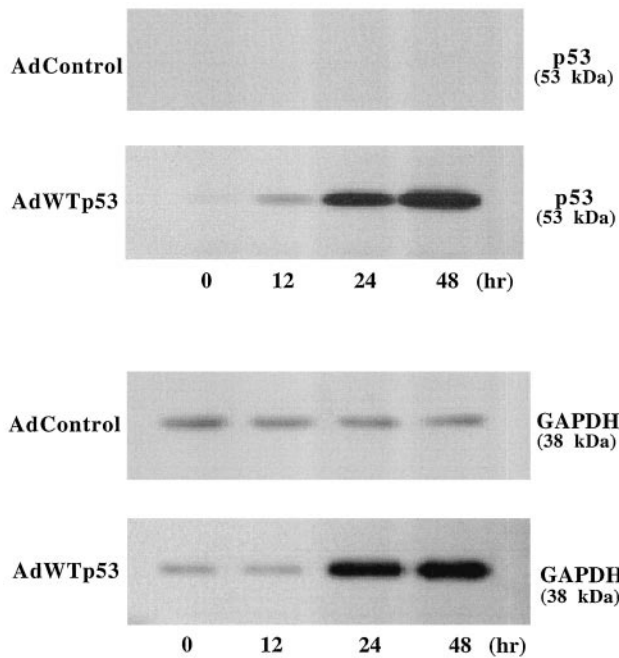


Figure 11. p53 gene transfection into PC12 cells induces GAPDH overexpression. PC12 cells in cultures were incubated at 37°C for different lengths of time after infection with 20 pfu/cell of either AdWTP53 or AdControl. Western blotting for p53 was determined at indicated times as described in Materials and Methods using 30 μ g of cell homogenate proteins. Western blotting for GAPDH was determined using 7 μ g of 2×10^5 g pellet proteins. Note that infection of cells with AdWTP53, but not AdControl, induced a dramatic time-dependent increase in p53 and GAPDH protein. The experiment has been repeated three times with similar results.

treated cells was markedly enhanced. Additionally, AraC-induced apoptosis was shown to be suppressed in granule cells from p53 null mice (Enokido et al., 1996b; the present study). These results suggest that p53 plays an essential role in mediating AraC-induced apoptotic death of cerebellar granule cells. The overexpression of p53 induced by AraC also suggests that this apoptotic insult involves extensive DNA damage before p53 induction, as suggested previously (Tomkins et al., 1994; Enokido et al., 1996a,b). It is interesting to note that AraC-induced apoptosis of granule cells is preceded by an increase in the levels of both mRNA and protein of p53, suggesting an involvement of transcriptional regulation of p53.

p53 upregulation was immediately followed by increased expression of Bax; the latter was blocked by p53/AS but not p53/S, reinforcing the well established notion that Bax is a p53-responsive target gene and is involved in p53-mediated neuronal death (Hughes et al., 1997; Xiang et al., 1998). Importantly, GAPDH mRNA and protein levels were increased after p53 overexpression, and GAPDH upregulation was effectively suppressed by p53/AS. Conversely, neither p53 nor Bax overexpression was affected by GAPDH/AS, which is neuroprotective by blocking GAPDH upregulation and the subsequent nuclear translocation (Ishitani and Chuang, 1996; Saunders et al., 1997; Ishitani et al., 1998). Moreover, our results show that p53 gene knock-out suppressed AraC-induced GAPDH upregulation in mouse cerebellar granule cells compared with that in wild-type neurons. These observations suggest that GAPDH overexpression is an event downstream of p53 expression and may function independently of Bax in mediating AraC-induced apoptosis of

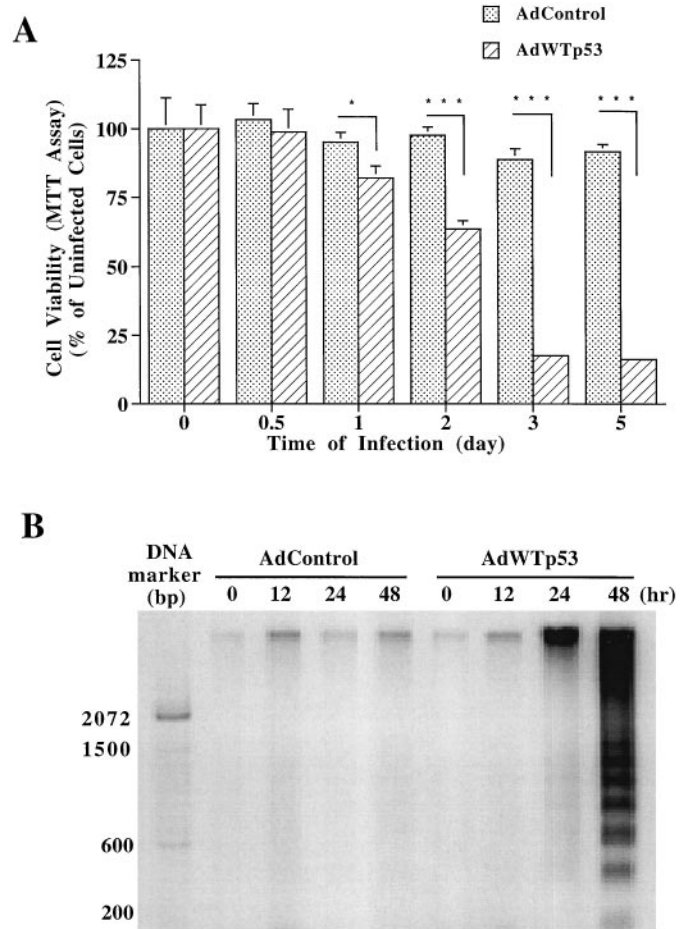


Figure 12. p53 gene transfection into PC12 cells induces delayed DNA fragmentation. Cells were infected with 20 pfu/cell of AdWTP53 or AdControl as described in the legend to Figure 11. At indicated times, cells were examined for viability using MTT assay (*A*) or for internucleosomal DNA cleavage using agarose gel electrophoresis (*B*). Results of MTT assay are expressed as percentage of mean \pm SEM of the respective zero time control from eight independent determinations. The results of DNA agarose gel electrophoresis are from a typical experiment of three such experiments. Note that infection of cells with AdWTP53, but not AdControl, induced a time-dependent loss of cell viability and an increase in the level of internucleosomally cleaved DNA in the soluble DNA preparation. * $p < 0.05$; *** $p < 0.001$ using one-way ANOVA with Fisher's PLSD test.

granule cells. Moreover, our results raise the possibility that GAPDH is a novel target of p53. In further support of this notion, we found that high-efficiency transfection of p53 gene into PC12 cells by infection with AdWTP53 resulted in dramatic overexpression of not only p53 but also GAPDH protein, and induction of these proteins preceded internucleosomal DNA cleavage and extensive cell death. In contrast, infection with AdControl produced neither p53 and GAPDH increase nor apoptosis.

Our results show that levels of Bax mRNA and protein were rapidly and markedly increased by AraC in cerebellar granule cells. Other studies have suggested that Bax may produce cell death by induction of the mitochondrial permeability transition, which can result in disruption of plasma membrane integrity or release of cytochrome C from mitochondria (Pastorino et al., 1998; Chen and Chuang, 1999). The released cytochrome C may bind to Apaf-1, causing caspase cleavage of substrate proteins and, ultimately, cell death (Liu et al., 1996). The involvement of

Bax in neuronal apoptosis seems to depend on the nature of stimuli. For example, Bax is required for apoptosis induced by deprivation of trophic factors and high potassium but is not involved in NMDA-induced excitotoxicity in some studies (Deckwerth et al., 1996; Miller et al., 1997).

The precise mechanism underlying the involvement of GAPDH in apoptosis is unclear. Because overexpressed GAPDH is translocated from the cytosol to the nucleus in neurons and non-neuronal cells (Saunders et al., 1997, 1999; Sawa et al., 1997; Ishitani et al., 1998; Shashidharan et al., 1999), one may surmise that GAPDH-dependent apoptosis is caused by a perturbation of nuclear functions. In this context, we found that AraC-induced GAPDH nuclear translocation occurs concurrently with a nuclear loss of dehydrogenase and uracil glycosylase activities (Saunders et al., 1999), suggesting that GAPDH undergoes structural and functional changes after transport to the nucleus. In the cytosol, GAPDH binds to glutamine repeats in huntingtin, a gene product of Huntington's chorea, and other polyglutamine-containing proteins associated with neurodegenerative diseases (e.g., ataxin for spinocerebellar ataxia) (Burke et al., 1996). Because N-terminal fragments of huntingtin and ataxin-3 are found in the intranuclear inclusions of cells in affected brain areas (Davies et al., 1997; DiFiglia et al., 1997; Paulson et al., 1997), it seems possible that GAPDH is a "carrier" mediating the transfer of these disease-related proteins to the nucleus, thus contributing to the progression of neuronal apoptosis.

Recently, it has been shown that the administration of *R*-deprenyl and its derivatives can protect cerebellar granule cells from AraC-induced apoptosis (Paterson et al., 1998). This neuroprotective effect could be attributable to the ability of these drugs to bind overexpressed GAPDH, thereby inhibiting the proapoptotic activity of GAPDH; such an effect was demonstrated in an independent study using neuroblastoma cells (Kragten et al., 1998). Clearly, molecular mechanisms underlying GAPDH-mediated apoptotic death in neurons and non-neuronal cells require future detailed investigation.

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