

Apoptosis induced by adenovirus-mediated p14^{ARF} expression in U2OS osteosarcoma cells is associated with increased Fas expression[☆]

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Abstract

The INK4A/ARF locus on chromosome 9 is a tumor suppressor gene frequently mutated in human cancers. In order to study the effects of p14^{ARF} expression in tumor cells, we constructed a recombinant adenovirus containing p14^{ARF} cDNA (Adp14^{ARF}). Adp14^{ARF} infection of U2OS osteosarcoma cells which has wild type p53 and mutant p14^{ARF} revealed high levels of p14 (ARF) expression within 24 h. In addition, Adp14^{ARF}-mediated expressing of p14 (ARF) was associated with increased levels of p53, p21, and mdm2 protein. Growth inhibition assays following Adp14^{ARF} infection demonstrated that the growth of U2OS cells was inhibited relative to infection with control virus. Furthermore, TUNEL analysis as well as PARP cleavage assays demonstrated that Adp14^{ARF} infection was associated with increased apoptosis in U2OS cell line and that it was associated with Adp14^{ARF} induced overexpression of Fas and Fas-L. Addition of Fas-L neutralizing antibody NOK-1 decreased Adp14-mediated cell death, indicating that p14 (ARF) induction of the Fas pathway is associated with increased apoptosis. The finding that Adp14^{ARF} infection did not induce Fas expression in U2OS/E6 and MCF/E6 cells suggests that wild type p53 expression may be necessary for Adp14^{ARF}-mediated induction of Fas. The observation that overexpression of p53 by Adp53 infection in MCF-7 does not induce increased Fas protein levels nor apoptotic cell death suggests that p53 overexpression is required but not sufficient enough for apoptosis. These studies suggest there are other mechanisms other than induction of p53 in ARF-mediated apoptosis and gene therapy using Adp14^{ARF} may be a promising treatment option for human cancers containing wild type p53 and mutant or deleted p14 expression. © 2004 Elsevier Inc. All rights reserved.

Keywords: p14^{ARF}; Apoptosis; Fas; p53; Adenovirus

The *INK4a/ARF* gene locus produces two transcripts which use alternative first exons, E1 α and E1 β , each of which is joined through the identical splice acceptor site to sequences in exon 2 [1,2]. The exon 1 α -exon 2-exon 3 transcript encodes the CDK4/6-specific inhibitor p16^{INK4a}, while the exon 1 β -exon 2-exon 3 transcript

uses an alternative reading frame and encodes ARF. p16^{INK4a} and ARF have no homology at the amino acid level [3]. Translation from the mouse β cDNA yields a polypeptide of 169 amino acids with a molecular mass of 19 kDa (p19^{ARF}) while human ARF protein is predicted to be 132 amino acids long with a molecular mass of 14 kDa. Mouse and human ARF polypeptides are 45% identical in exon 1 β segments and 50% identical overall.

Homozygous deletion of the *INK4a/ARF* locus, which causes a loss of both the *INK4a* gene and the *ARF* gene, occurs frequently in a variety of human cancers. Thus, determining the individual contributions of each of the two genes to tumor suppression has been difficult [4,5]. However, mice lacking p19^{ARF} but retaining an

[☆] **Abbreviations:** ARF, alternative reading frame; MDM2, mouse double minute 2; CMV, cytomegalovirus; pfu, plaque-forming unit; FACS, fluorescence-activated cell sorting; TUNEL, terminal deoxynucleotidyl transferase-mediated dUTP nick end labeling; FITC, fluorescein isothiocyanate.

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intact p16^{INK4a} gene develop multiple tumors [5], indicating that ARF, like INK4a, is also a bona fide tumor suppressor. ARF overexpression results in cell cycle arrest, however, this is not associated with ARF binding to CDKs or direct inhibition of cyclin-CDK activity [3]. Furthermore, in contrast to INK4a, there is no evidence that ARF exerts its growth-inhibitory effect through the pRB pathway [6]. ARF does require the presence of wild type p53 for induction of cell cycle arrest and cell lines that lack p53 or have p53 mutations are resistant to growth arrest by ectopic expression of ARF [5].

Cell cycle arrest and apoptosis are two cellular responses induced by a wide variety of DNA damaging agents. Cell cycle arrest provides a critical opportunity for cells to repair damaged DNA, while cells with extensive, and presumably irreparable, DNA damage can be eliminated by apoptosis. It is not surprising that one or both of these processes are frequently altered in cancer cells [7–11].

Fas (CD95/APO1), a death-domain bearing member of the tumor necrosis factor receptor superfamily, is a transmembrane protein that, on specific engagement with Fas-ligand initiates an apoptotic response in cells. Oligomerization of the Fas receptors following cross-linking with Fas ligand or agonistic antibodies leads to recruitment of caspase-8 (FLICE/MACH/Mch5) to the receptor complex and triggers the apoptosis-inducing protease cascade by cleaving the caspase-1 zymogen [12,13]. Several studies have examined the relationship between Fas and p53 expression [14]. Overexpression of p53 resulted in Fas induction in lung cancer cells [15].

Others have reported that exogenous expression of p53 by microinjection can induce functional Fas protein that can be stimulated by the anti-Fas antibody to induce apoptosis [16]. We also previously reported that overexpression of wild type p53 can induce Fas protein and that this effect was observed predominantly in cells which express mutant p53 or null for p53 [17].

To study the function of p14^{ARF}, we have constructed a recombinant replication-deficient adenovirus expressing human p14^{ARF} (Adp14^{ARF}). Previously we reported that infection of cells with Adp14^{ARF} resulted in high levels of p14^{ARF} expression and induction of apoptosis in human breast cancer cell lines MCF-7 and MDA-MB-231 [18]. In this report, we provide evidences that Adp14^{ARF} induces apoptosis in human breast and osteosarcoma cancer cells are associated with upregulation of Fas.

Results

Adenovirus-mediated p14^{ARF} expression

Adp14^{ARF}-mediated p14^{ARF} expression was examined by Western blot analysis following infection of human osteosarcoma cancer cells U2OS which have wild type p53 and mutant type p14 with Adp14^{ARF}. As shown in Fig. 1A, 24 h after infection with Adp14^{ARF}, there was a marked increase in p14^{ARF} protein levels in U2OS cells. In U2OS cell lines, the increase in p14^{ARF} protein following infection with Adp14^{ARF} was dependent on the

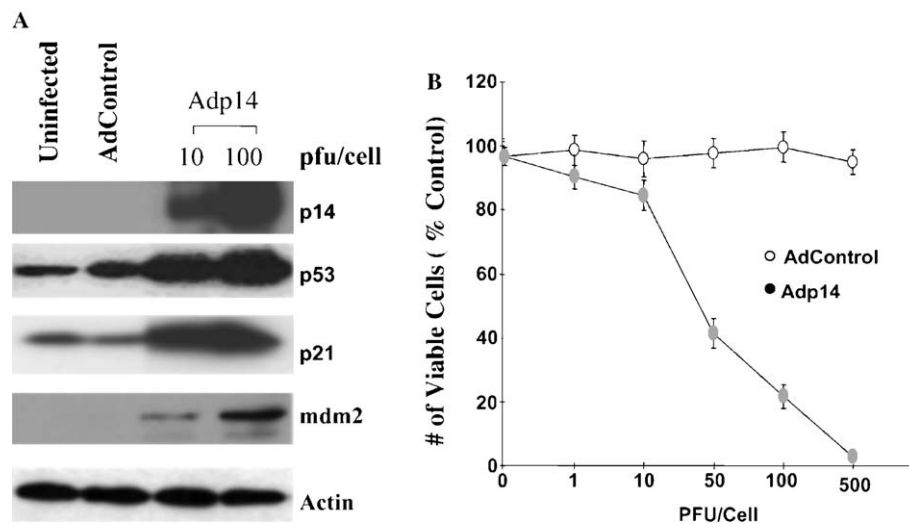


Fig. 1. Effects of Adp14^{ARF} on expression of p14 and associated proteins and on cell viability following Adp14 infection. (A) Western blot analysis of p14, p53, p21, and mdm2 in U2OS cells following Adp14 infection. Cells were infected with 100 pfu/cell of AdControl and varying doses of Adp14. After 24 h incubation at 37 °C, whole cell lysates were collected and subjected to Western blot analysis as described in Materials and methods. Cell lines and adenoviral vectors used are shown on top of the panel; mock infected cells are labeled uninfected (Ui). (B) Dose dependent effects of Adp14 on the survival of U2OS cells. Cells were infected with Adp14 and incubated for 3 days. Viable cells were counted by trypan-blue exclusion assay. Results are expressed as percentage of change in the number of viable cells from cultures without any treatments. Data points represent the mean values of four replicates, with bars indicating SEM.

dose of the virus. Previous studies had indicated that p14^{ARF} expression can alter the intracellular levels of p53 protein as well as the level of p21, a p53 inducible protein, in p53 WT cell lines. As shown in Fig. 1B, the levels of p53 in U2OS cells increased following infection with Adp14^{ARF}. Moreover, following 24h infection of U2OS cells with Adp14^{ARF}, there was marked increase in p21 expression as well (Fig. 1A). Similar results were obtained in MCF-7 cells (data not shown). Thus, p14^{ARF}-mediated increase in p53 protein was also associated with increased levels of p21 protein in U2OS and MCF-7 cells.

Since previous studies have indicated that the induction of p53 protein by p14^{ARF} resulted from the interaction of p14^{ARF} with mdm2, a p53 binding protein, we also examined mdm2 levels following Adp14^{ARF} infection. As shown in Fig. 1A, Adp14^{ARF} infection of U2OS cells also results in a marked increase in mdm2 protein levels.

Adp14^{ARF}-mediated cytotoxicity in U2OS osteosarcoma cells

The effect of Adp14^{ARF} infection on cytotoxicity of human cancer cells was assessed using a trypan blue exclusion assay. As shown in Fig. 1B, 3 days following infection with Adp14^{ARF}, there was marked cytotoxicity in both U2OS osteosarcoma cells. Following infection with Adp14^{ARF} at 500 pfu/cell, >80% of cells were not viable. In contrast, following infection with 500 pfu/cell of AdControl, >90% of cells were viable. The IC₅₀ for Adp14^{ARF} was between 10 and 50 pfu/cell for U2OS cells (Fig. 1B), while that for the control virus was over 5000 pfu/cell for U2OS cells (data not shown). Thus, Adp14^{ARF} is approximately 100 times more cytotoxic to U2OS osteosarcoma cells compared to control virus.

p14^{ARF} expression induces apoptosis in U2OS osteosarcoma cells

We next investigated the mechanism of Adp14^{ARF} induced cytotoxicity in U2OS cells. As shown in Fig. 2A, PI staining indicated that uninfected or AdControl-infected U2OS cells had a small sub-G1 population. In contrast, cells infected with Adp14^{ARF} exhibited a marked increase in the proportion of sub-G1 population of cells (23.6% versus 10.4% for control virus infected U2OS). Similar results were obtained using TUNEL assay for evaluation of Adp14^{ARF} induced apoptosis. As shown in Fig. 2B, Adp14^{ARF} infection of U2OS cells resulted in a marked shift of the fluorescent peak. These results are consistent with induction of apoptosis in U2OS cell lines following infection with Adp14^{ARF}.

The process of apoptosis involves the activation of a cascade of proteolytic enzymes belonging to the caspase family resulting in cleavage of several key cellular

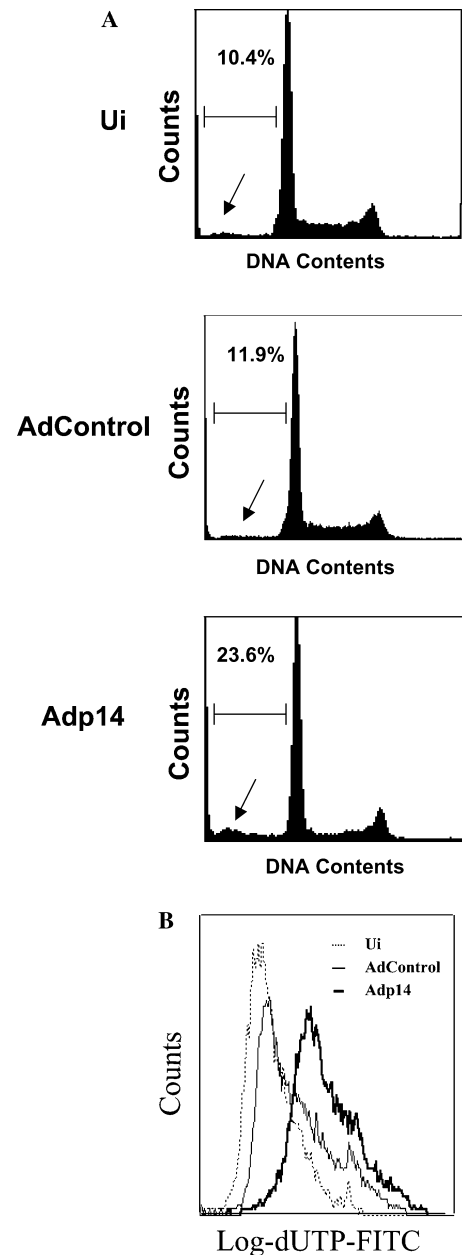


Fig. 2. Cell cycle and apoptosis analysis of Adp14 infected cells. (A) U2OS Cells (1×10^6) were infected with Adp14 (100 pfu), AdControl (100 pfu) or uninfected for 3 days and subjected to FACS analysis using propidium iodide. Cells were analyzed for apoptosis as represented by the appearance of a sub-G1 population. (B) Flow cytometric TUNEL analyses for apoptosis on U2OS cells. Cells (1×10^6) were infected with Adp14 (100 pfu), AdControl (100 pfu) or uninfected for 3 days and subjected to TUNEL analysis.

components including poly (ADP-ribose) polymerase (PARP) and CPP32. We therefore examined the cleavage of these proteins following Adp14^{ARF} infection. As shown in Fig. 3A, infection of U2OS cells with Adp14^{ARF} resulted in the appearance of a specific M_r 85000 cleavage product of PARP, which was not observed in uninfected cells or cells infected with a control virus (AdControl).

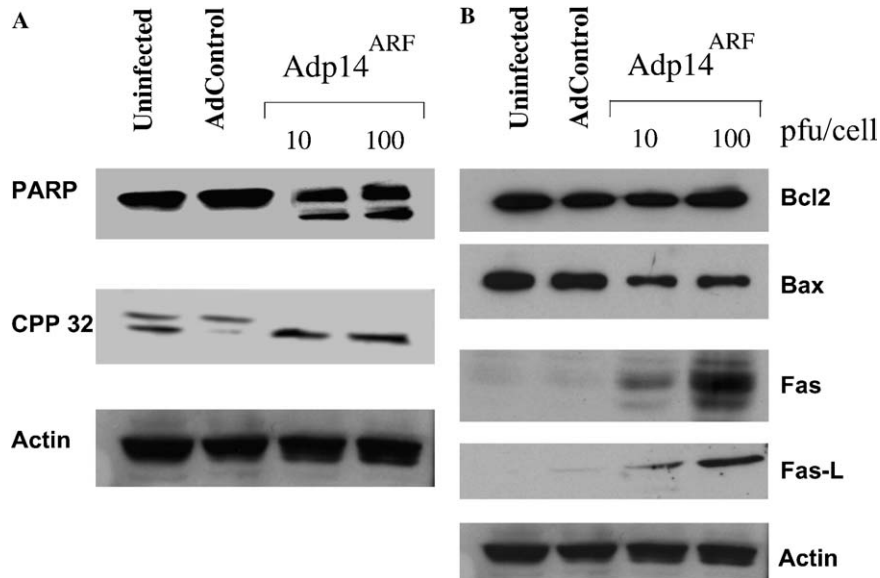


Fig. 3. Western blot analyses of apoptosis related proteins. Cells (5×10^6) were infected with 100 pfu/cell of AdControl and 10, 100 pfu/cell of Adp14. After 3 days of incubation, whole cells were collected and subjected to Western blot analysis. (A) PARP, CPP32, and actin in U2OS cells. (B) Bcl2, Bax, Fas, Fas-L, and actin in U2OS cells.

Since degradation of CPP32 has also been reported to be associated with apoptosis, we also examined the levels of CPP32 protein in uninfected U2OS cells or cells infected with AdControl virus or Adp14^{ARF} virus. As shown in Fig. 3A, the CPP32 cleavage protein level increased in U2OS cells following infection with increasing doses of Adp14^{ARF}, but remained unchanged in cells infected with AdControl. These results provide additional evidence that overexpression of p14 following Adp14 infection results in apoptosis in U2OS osteosarcoma cells.

Role of Fas induction in p14^{ARF}-mediated apoptosis

To elucidate the mechanism of p14 induced apoptosis in U2OS cells, we examined the expression of several proteins involved in apoptosis including Bcl2, Bax, Fas, and Fas-L by Western blot analysis. As shown in Fig. 3B, although Bcl2 expression was unchanged and Bax expression was little decreased, both Fas and Fas-L protein levels were increased in Adp14^{ARF} infected U2OS cells compared to uninfected or AdControl infected U2OS cells. These studies suggest a role for Fas/FasL in ARF-mediated cytotoxicity in U2OS cells.

Apoptosis and Fas expression by p14 is p53 dependent

To test the effect of p53 on Adp14 induced Fas expression, we made U2OS/E6 cells which were transfected with human papillomavirus E6 gene to inactivate p53 protein. Fig. 4A showed p53 protein was successfully deleted from U2OS cells. U2OS/E6 cells showed decreased FAS and Fas-L expression following infection

with Adp14^{ARF} compare with U2OS cells (Fig. 4B). U2OS/E6 cells were also more resistant to Adp14 compare with U2OS cells (data not shown). These results

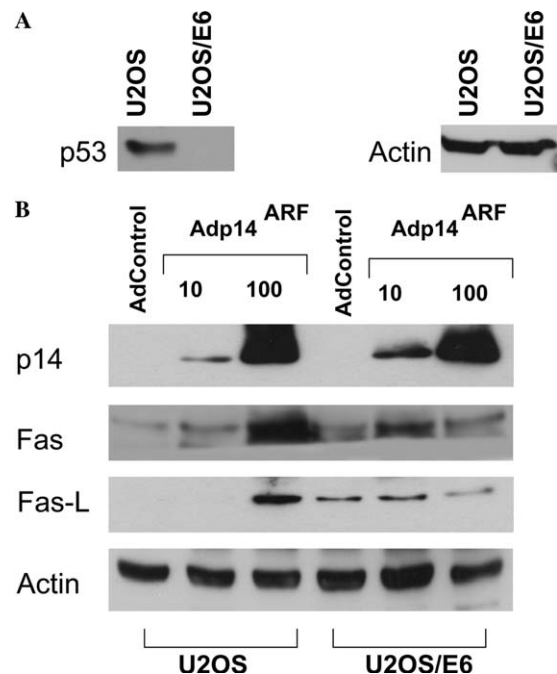


Fig. 4. Effects of Adp14 infection on Fas and Fas-L expression are associated with p53 status on cells. (A) U2OS Cells were infected with E6 virus to delete p53 protein. Western blot analysis showed p53 was successfully deleted from U2OS cells. (B) Western blot analyses of p14, Fas, Fas-L, and actin in U2OS and U2OS/E6 cells. Cells (5×10^6) were infected with 10 and 100 pfu/cell of Adp14 and 100 pfu/cell of AdControl. After 3 days of incubation, whole cells were collected and subjected to Western blot analysis.

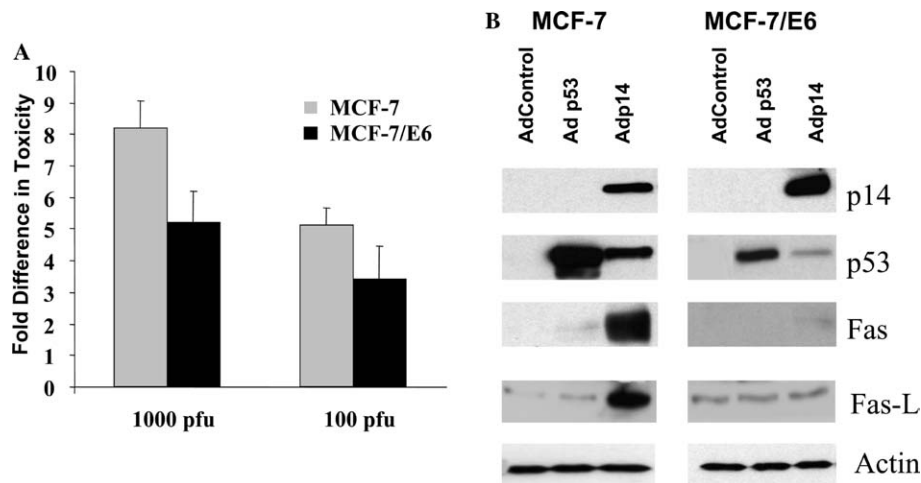


Fig. 5. Effect of Adp14 infection on cell cytotoxicity associated p53 status on cells. (A) Cells (1×10^3) were infected with 100 and 1000 pfu/cell of Adp14 to MCF-7 and MCF-7/E6 cells. After 5 days of incubation cell viability was accessed by MTT assay. The IC_{50} s for Adp14 and AdControl were calculated for each cell line assuming the survival rate of uninfected cells was 100%. Results shown are the ratios of the IC_{50} s following Adp14 infection relative to the IC_{50} s of AdControl in each cell line. Data points represent the mean values of four replicates, with bars indicating SEM. (B) Western blot analyses of p14, p53, Fas, Fas-L, and actin in MCF-7 and MCF-7/E6 cells. Cells (5×10^6) were infected with 100 pfu/cell AdControl, Adp53, and Adp14. After 3 days of incubation, whole cells were collected and subjected to Western blot analysis.

suggest p53 role on Fas and Fas-L expression after p14 overexpression. Shown in Fig. 5A are the cytotoxicity results of Adp14^{ARF} infection in MCF-7 cells which containing wild type p53 and MCF-7/E6 cells which were transfected with human papillomavirus E6 gene which binds to and inactivate p53. We have previously shown that Adp14^{ARF} infection results in increased cell death in MCF-7 cells relative to control virus [18]. As shown in Fig. 5A, the cytotoxicity of Adp14^{ARF} is diminished in cells containing reduced levels of wild type p53 (MCF-7/E6). Furthermore, the decreased toxicity of Adp14^{ARF} in MCF-7/E6 cells is also associated with decreased FAS expression in these cells following infection with Adp14^{ARF}.

We have previously shown that infection of MCF-7 cells with an adenoviral vector expressing wild type p53 (Adp53) resulted in cell cycle arrest but not apoptosis [17]. As shown in Fig. 5B, while Adp14^{ARF} infection in MCF-7 cells results in increased p53 expression and a concomitant increase in FAS expression, infection with Adp53 results in p53 overexpression without induction of FAS. Taken together these results indicate that wild type p53 overexpression may be necessary, but not necessarily sufficient for p14^{ARF}-mediated cytotoxicity (Fig. 5B). Furthermore, the pretreatment of MCF-7 cells with neutralizing antibody against Fas-L (NOK-1) reduced toxicity of Adp14^{ARF} (Fig. 6). These results suggest that the Fas pathway is involved in the induction of apoptosis in MCF-7 cells by Adp14^{ARF}. These studies also suggest that p14^{ARF}-mediated apoptosis involves other cellular change mechanisms in addition to p53 overexpression.

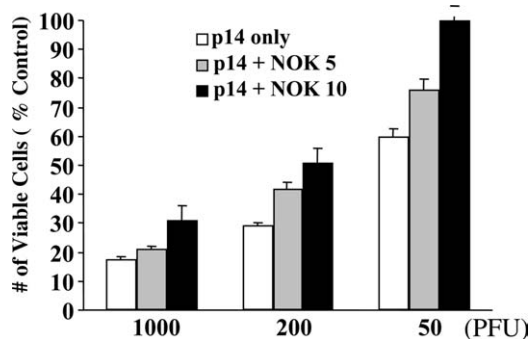


Fig. 6. Inhibition of Adp14-mediated cytotoxicity by anti Fas-L antibody in MCF-7 cells. Cells (1×10^3) were infected with 50, 200, and 1000 pfu/cell of Adp14. 5 and 10 μ g/ml of NOK-1 was added 24 h after Adp14 infection. After 5 days of incubation cell viability was accessed by MTT assay. Results are expressed as percentage of change in the number of viable cells from cultures without any treatments. Data points represent the mean values of four replicates, with bars indicating SEM.

Discussion

Recent studies have shown that the INK4A/ARF gene can function as a tumor suppressor gene and that p14^{ARF} can inhibit cell growth. In previous report we constructed an adenoviral vector expressing p14^{ARF} to study the function of p14^{ARF} overexpression in human breast cancer cells [18]. In this report we also found that overexpression of p14^{ARF} results in apoptosis in U2OS cells. This was demonstrated by TUNEL assays as well as cleavage of PARP and CPP32. The mechanism of p14^{ARF} inducing apoptosis in the cells involves overexpression of both FAS and FAS ligand and is inhibited at least in part, by incubation of cells with antibody directed against FAS ligand.

Previous studies by our laboratory have shown that overexpression of wild type p53 in MCF-7 cells using a recombinant adenoviral vector resulted in overexpression of p21 and increased G1 cell cycle arrest without induction of apoptosis [19]. Furthermore, we found that overexpression of wild type p53 in MCF-7 cells did not result in increased expression of FAS, while other studies showed that enhanced expression of FAS and wild type p53 expression in MCF-7 cells did result in increased apoptosis [17]. In this study, we report that overexpression of p14^{ARF} in U2OS and MCF-7 cells resulted in increased expression of both p53 and Fas and induction of apoptosis. These effects of p14^{ARF} expression involve other changes in cells in addition to induction of wild type p53 while previous studies had suggested the effect of p14^{ARF} is mediated through p53 [20]. The results proposed in this report as well as previous studies from our laboratory [18] suggest that the induction of wild type p53 by p53 adenoviral vector is by itself not sufficient to induce p53-mediated apoptosis in cells, indicating that other effects of p14^{ARF} are needed to induce apoptosis in addition to p53 regulation.

CDKIs, including p21 and p16, have been proposed as candidates for gene therapy of cancer [19,21]. p14^{ARF} may also represent a good choice for gene therapy of cancer as it can produce not only cell cycle arrest but also it can induce apoptosis in some cancer cells. Additional studies on the usefulness of Adp14^{ARF} as a vector for gene therapy in combination with chemotherapy should be explored.

Materials and methods

Cell culture. Human breast cancer cell line MCF-7 (ATCC HTB-22, wild type p53, mutant type p14), MCF-7/E6 (kindly provided by Dr. Albert J Fornace Jr, NCI, NIH, Bethesda, MD, USA), and osteosarcoma cell line U2OS (ATCC HTB96, wild type p53, mutant type p14) were cultured in Improved minimal essential medium (IMEM) (Life Technologies, Gaithersburg, MD, USA) containing 10% fetal bovine serum as described previously [22]. U2OS/E6 cells were obtained after transduction of human papillomavirus E6 and selection with neomycin for 10 days. p53 deletion was confirmed with Western blot analysis.

Construction of adenoviral vectors and infection procedure. Human p14^{ARF} cDNA that expresses full-length human p14^{ARF} protein was a kind gift from Dr. K.H. Vousden of the National Cancer Institute-Frederick Cancer Research and Development Center, Frederick, MD. A recombinant adenovirus containing human p14^{ARF} cDNA, designated Adp14^{ARF}, was constructed by homologous recombination using methods described previously [19,21,23]. Briefly, human p14^{ARF} cDNA was cloned into an adenovirus shuttle vector containing Ad5 backbone. Following transfection with Ad5 genomic DNA into 293 cells, recombinant plaques were screened for the presence of p14^{ARF} sequences by polymerase chain reaction (PCR) using the following primers: 5'-AGTGAGGGTTTTCGTGGTTTAC-3' (upstream) and 5'-ATGACCTGGTCTTCTAGGAAGCGG-3' (downstream). The control adenovirus (AdControl) used in this study is identical to Adp14^{ARF} except that it does not contain any transgene.

Infection with Adp14^{ARF} and the control virus (AdControl) was accomplished by incubating cells with various doses of each virus

(expressed as pfu/cell) in serum-free medium for 2 h, followed by the addition of medium containing 10% serum and further incubation at 37 °C for the required time.

Cytotoxicity assays. Cell viability was assessed following appropriate treatment by microscopic examination of trypan-blue stained cells or was measured using the standard MTT assay, as described [24]. Cells (1500/well) were plated in each 96-well plate and incubated at 37 °C overnight. Following exposure to different concentrations of Adp14 or AdControl for 7 days at 37 °C, MTT (50 µl of 2 mg/ml) in 1 × PBS was added to each well and the cells incubated 4 h at 37 °C. Following centrifugation of plates at 500g for 10 min, DMSO (120 µl) was added to each well and incubated for 1 h at room temperature on an orbital shaker. The OD₅₇₀ was determined using Ultra-microplate reader (ELx 808, Bio-tek instrument, Winooski, VT, USA). Cytotoxicity was expressed as the percentage of OD₅₇₀ relative to that of untreated cells.

Cell cycle analysis. Cells were plated in 10-cm dishes (1–2 × 10⁶ cells/dish) 1 day prior to infection. Cells were exposed to various concentrations of Adp14^{ARF} (0, 1, 10, and 100 pfu/cell) for increasing times at 37 °C. Cells were harvested, fixed with 2% paraformaldehyde, and stained for cell cycle analysis using propidium iodide (PI) [25]. DNA content was measured using a flow cytometer (FACSCalibur, Becton–Dickinson, Mountain View, CA, USA) and cell cycle analysis was performed with the ModFit software (Becton–Dickinson). Analysis of the sub-G1 (sub-diploid) population was done by measuring the fluorescence activity of propidium iodide (PI)-stained DNA of permeabilized and fixed cells on the FACSCalibur as detailed above [22,25].

$$\% \text{sub-G1 population} = \frac{\text{Number of cells below G1}}{\text{Total number analyzed}} \times 100\%.$$

TUNEL assay. For detection of apoptosis by the TUNEL method, cells were plated in 15-cm dishes. When cells were 70% confluent, Adp14^{ARF} was added for varying times. The cells were harvested by gentle pipetting and collected by centrifugation at 1000g for 5 min. Cells were washed with ice cold PBS and then fixed with 4% paraformaldehyde. TUNEL assay was performed using MEBSTAIN Apoptosis kit (Medical and Biological Laboratories, Nagoya, Japan) according to the manufacturer's instructions.

Western blot analysis. Cells were plated in 15-cm dishes. When cells were 70% confluent, adenoviral vectors were added. Following incubation at 37 °C, the cells were harvested by gentle scraping, and whole cell lysates were prepared and subjected to Western blot analysis as described previously [22,25]. Blots were probed with 2 µg/ml of antibody reactive with p14^{ARF} (C-18; Santa Cruz Biotechnology, Santa Cruz, CA, USA), p53 (DO-1; Santa Cruz Biotechnology, Santa Cruz, CA, USA), p21 (Ab-1; Oncogene, Cambridge, MA, USA), CPP32 (K-19; Santa Cruz Biotechnology, Santa Cruz, CA, USA), PARP (Ab-2; Calbiochem, San Diego, CA, USA), FAS (C-20; Santa Cruz Biotechnology, Santa Cruz, CA, USA), and FAS-L (N-20; Santa Cruz Biotechnology, Santa Cruz, CA, USA).

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