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p53-Dependent but ATM-Independent Inhibition of DNA Synthesis and G2 Arrest in Cadmium-Treated Human Fibroblasts

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Abstract

This study focused on the activation of cell cycle checkpoint responses in diploid human fibroblasts that were treated with cadmium chloride and the potential roles of ATM and p53 signaling pathways in cadmium-induced responses. The alkaline comet assay indicated that cadmium caused a dose-dependent increase in DNA damage. Cells that were rendered p53-defective by expression of a dominant-negative p53 allele or knockdown of p53 mRNA were more resistant to cadmium-induced inactivation of colony formation than normal and ataxia telangiectasia (AT) cells. Synchronized fibroblasts in S were more sensitive to cadmium toxicity than cells in G1 suggesting that cadmium may target some element of DNA replication. Cadmium produced a dose- and time-dependent inhibition of DNA synthesis. An immediate inhibition was associated with severe delay in progression through S phase and a delayed inhibition seen 24 h after treatment was associated with accumulation of cells in G2. AT and normal cells displayed similar patterns of inhibition of DNA synthesis and G2 delay after treatment with cadmium, while p53-defective cells displayed significantly less of the delayed inhibition of DNA synthesis and accumulation in G2 post-treatment. Total p53 protein and ser15-phosphorylated p53 were induced by cadmium in normal and AT cells. The p53 transactivation target Gadd45 α was induced in both p53-effective and p53-defective cells after 4 h cadmium treatment, and this was associated with an acute inhibition of mitosis. Cadmium produced a very unusual pattern of toxicity in human fibroblasts, inhibiting DNA replication and inducing p53-dependent growth arrest but without induction of p21^{Cip1/Waf1} or activation of Chk1.

Keywords

Cadmium; checkpoint; fibroblasts; DNA synthesis; p53; ataxia telangiectasia; Gadd45 α

Introduction

Cadmium is a systemic poison affecting many cellular functions. Cadmium contamination poses a serious health threat throughout the world, and cadmium has been classified as a human carcinogen by the International Agency for Research on Cancer (1994). Toxic responses to cadmium exposure include kidney damage, respiratory diseases, neurological disorders, and

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lung, kidney, prostate and testicular cancers (Abdulla and Chmielnicka, 1989; Waalkes, 2003).

Cadmium exposure can induce intracellular damage through several mechanisms. In cultured cells cadmium produces direct and indirect genotoxic effects such as DNA strand breaks, DNA-protein crosslinks, oxidative DNA damage, and chromosomal aberrations (Dally and Hartwig, 1997; Hwua and Yang, 1998; Misra *et al.*, 1998). Several cellular factors that respond to DNA damage to regulate proliferation also respond to cadmium exposure. The tumor suppressor gene p53 and proto-oncogenes c-Jun, c-Fos and c-Myc were transcriptionally activated by cadmium (Jin and Ringertz, 1990; Zheng *et al.*, 1996), and cadmium caused an irreversible G2/M arrest (Yang *et al.*, 2004). It was therefore of interest to determine whether cadmium-induced DNA damage causes cells to arrest cell division through p53-dependent cell cycle checkpoints.

Cell cycle checkpoints are biochemical surveillance pathways that actively delay or arrest cell cycle progression in response to DNA damage and activated oncogenes (Kaufmann and Paules, 1996; Kastan and Bartek, 2004). DNA damage checkpoints minimize the probability of replicating and segregating damaged DNA and, therefore, reduce the frequencies of mutations and chromosomal aberrations that are induced by genotoxic stress. p53 and ataxia telangiectasia-mutated (ATM) are important signaling molecules in checkpoint responses to DNA damage (Sancar *et al.*, 2004). ATM is a protein kinase that is activated by autophosphorylation in response to DNA damage and alterations in chromatin structure (Bakkenist and Kastan, 2003). As guardian of the genome and a substrate for ATM, p53 enforces G1 arrest through transactivation of p21^{Cip1/Waf1} and G2 delay through transrepression of growth-regulated genes such as cyclin B1 and Cdk1 (Agarwal *et al.*, 1998). Germline mutations in p53 and ATM are seen in the familial cancer syndromes, Li-Fraumeni (Malkin *et al.*, 1990) and ataxia telangiectasia (AT) (Savitsky *et al.*, 1995), respectively. Checkpoint responses to environmental carcinogens such as cadmium may suppress cancer development.

Cadmium has the propensity to replace zinc in biological material and several DNA repair factors including p53 (Meplan *et al.*, 1999), XPA (Asmuss *et al.*, 2000), and hMutS- α (Clark and Kunkel, 2004; Banerjee and Flores-Rozas, 2005) are inhibited by cadmium. Inhibition of DNA repair or other elements of DNA damage response such as apoptosis could sensitize cells to carcinogenesis by endogenous or exogenous stresses that damage DNA (Mukherjee *et al.*, 2004; Sancar *et al.*, 2004). Reduced DNA repair increases mutagenesis and clastogenesis by chemical carcinogens and radiations, and reduced apoptosis increases the yields of cells that survive with mutations and chromosomal aberrations.

In the present study, the effects of cadmium chloride (CdCl₂) on DNA and cell division were examined in diploid human fibroblasts. Although fibroblasts are not targets of cadmium toxicity *in vivo*, they represent an excellent *in vitro* model for elucidating mechanisms of DNA damage response. Unlike transformed and cancer cell lines, diploid human skin fibroblasts express the full repertoire of repair and cell cycle checkpoint gene products that respond to carcinogen-induced DNA damage. Exposure of human fibroblast lines to cadmium caused DNA damage and induced a concentration- and time-dependent inhibition of DNA synthesis and mitosis. Cadmium induced a very unusual pattern of toxicity in fibroblasts, with p53-dependent inactivation of colony formation in the absence of induction of p21^{Cip1/Waf1}, and inhibition of DNA replication without activation of Chk1.

Materials and Methods

Cell lines and culture conditions

The normal human fibroblast strains, F1, F3 and F10, were derived from neonatal foreskin and established in culture according to established methods (Maher and McCormick, 1976; Boyer *et al.*, 1991). AT fibroblasts were isolated from the skin of an affected individual. The original fibroblast strain (GM02052A) was obtained from the NIGMS Human Genetic Cell Repository. Immortalized cell lines from these strains of human fibroblasts were obtained by ectopic expression of the human telomerase reverse transcriptase (hTERT), as previously described (Deming *et al.*, 2002; Heffernan *et al.*, 2002). The immortalized normal human fibroblasts were denoted as F1-hTERT, F3-hTERT, F10-hTERT and the telomerase-expressing AT cell line as GM02052A.

Fibroblasts were cultured in DMEM (Invitrogen) supplemented with 2 mM L-glutamine (Invitrogen) and 10% or 20% (AT cells) fetal bovine serum (Hyclone). All cell lines were maintained at 37°C in a humidified atmosphere of 5% CO₂, and were tested and shown to be free of mycoplasma contamination using a commercial kit (Gen-Probe).

Plasmids and viruses

F10-hTERT fibroblasts were engineered to express a dominant-negative form of p53 (a histidine to glutamine substitution at amino acid 179 in p53) by infection with a replication-defective retrovirus carrying the mutant p53 cDNA (a generous gift from Dr. Howard Liber, Colorado State University) and the neomycin-resistance gene under the control of the same viral promoter (pLXIN) (Simpson *et al.*, 2005). The resulting derivative line was denoted as F10-hTERT-p53H179Q. An isogenic cell line carrying the empty vector (F10-hTERT-LXIN) was derived in parallel.

The retroviral short interfering RNA (siRNA) vector to inactivate p53 was purchased from Oligoengine. Vesicular stomatitis virus glycoprotein G-pseudotyped, replication-defective retroviruses were produced as previously described following transient transfection of viral vector and helper plasmids into HEK 293T cells (Olsen *et al.*, 1993; Comstock *et al.*, 1997; Johnson *et al.*, 1998). As F10-hTERT was initially selected in 200µg/µl puromycin, stable expression of the p53 RNAi was done by growth in 400µg/µl puromycin. Batch cultures of puromycin-resistant F10-hTERT-p53RNAi were expanded and shown to display 90% inactivation of p53-dependent DNA damage G1 checkpoint function (results not shown).

Cell treatment with cadmium

A stock solution of cadmium chloride (Alfa Aesar) was prepared at 10 mM in sterile H₂O. Cadmium was added directly to culture medium. Cells were exposed to cadmium for 4 hours at concentrations ranging from 40–80 µM. After treatment, medium was removed, cells were rinsed with phosphate-buffered saline (PBS) and fresh medium replaced. A sham-treatment control was incorporated in each assay using the same manipulations but without cadmium. All experiments were performed in triplicate in independent trials to assess reproducibility.

Colony formation ability

Colony formation was measured in logarithmically growing cells, plated at 500 cells per 100 mm diameter dish and incubated for 8 hours before the 4 h cadmium treatment. Cells were cultured for 2 weeks, changing medium twice each week. Colonies were fixed and stained with a solution of 40% methanol and 0.05% crystal violet. Colonies of 50 or more cells were counted. Three individual dishes were assayed per treatment and the mean values were used to estimate cytotoxicity. Cytotoxicity was determined as the relative colony-forming ability (the cloning

efficiency of the treated cells divided by the cloning efficiency of the untreated cells, multiplied by 100).

Comet assay

The comet assay to detect DNA damage was performed using the method of Sasaki *et al.* with some modifications (Sasaki *et al.*, 1997). Briefly, fibroblasts were exposed to 40, 60 and 80 μM cadmium for 4 h. At the end of the incubation with cadmium, cells were removed from the plates with trypsin. Trypsin was inactivated with serum-containing medium and cells were collected by sedimentation and resuspension in PBS. Ten microliters of the cell suspension ($\sim 100,000$ cells/ml) were diluted in 70 μl low-melting-point agarose (0.75% w/v in PBS). The resulting suspension were embedded in previously prepared normal-melting-point agarose (1% w/v in PBS) on frosted slides followed by the addition of 75 μl of normal-melting-point agarose (1% w/v in PBS). The slides were then immersed in lysis buffer (2.5 M NaCl, 100 mM Na₂EDTA, 10 mM Tris-HCl, pH 10) for 1 h at 4°C in the dark. After lysis slides were placed in alkaline electrophoresis buffer (0.3 M NaOH, 1 mM Na₂EDTA) for 20 min at 4°C to denature DNA and express alkali-labile sites. Electrophoresis was carried out at 4°C for 20 min at 24 V and approximately 300 mA. The slides were then washed twice in neutralizing buffer (0.4 M Tris-HCl, pH 7.4) for 5 min. The DNA was stained by adding 45 μl of ethidium bromide (20 $\mu\text{g}/\text{ml}$) to each slide. More than fifty nuclei on each slide were examined for the presence of comet tails at 200 X magnification using a fluorescence microscope (Zeiss Axioskop) equipped with a green filter. The percent of nuclei with a DNA tail of $\geq 25 \mu\text{m}$ was used as the measure of DNA damage.

G1 and G2 checkpoint analyses

G1 and G2 checkpoint functions were analyzed at 0, 6 and 24 h after the 4 h cadmium treatment or sham treatment. G1 checkpoint function was assessed by measuring incorporation of 5'-bromo-2'-deoxyuridine (BrdU) by S phase cells (Kaufmann *et al.*, 1995) and G2 checkpoint function was assessed by measuring mitotic cells with 4N DNA that stained positive for mitosis-specific phospho-histone H3 (Juan *et al.*, 1998; Deming *et al.*, 2002). Summit (Cytomation Inc.) flow cytometry analysis software was used to quantify the numbers of unlabeled cells with 2N (G0/G1) and 4N DNA content (G2), BrdU-labeled cells with 2–4N DNA content (S), and 4N cells with phospho-histone H3 (M).

Cell synchronization

F1-hTERT, F3-hTERT and F10-hTERT fibroblasts were synchronized as previously described (Cordeiro-Stone *et al.*, 1986; Kaufmann and Wilson, 1990). Briefly, cells were plated at a density of $1.3 \times 10^4/\text{cm}^2$ and allowed to grow for 8 days to confluence arrest (G0 phase). Confluence-arrested cells were trypsinized, reseeded at 500 cells per 100 mm dish and incubated for 8 h to allow them to reenter the cell cycle and reach G1 or for 20 h to allow them to reach S. The cells were then treated with different concentrations of cadmium for 4 h followed by washing with PBS and incubation in fresh medium for two weeks to observe colony formation ability.

Cells were similarly released from confluence-arrest, reseeded at 1 million cells per 100 mm dish and incubated with the DNA polymerase inhibitor, aphidicolin (APC), at a concentration of 2 $\mu\text{g}/\text{ml}$ for 24 h to collect cells at the beginning of S phase (Kaufmann and Wilson, 1990). APC was washed out and cells incubated in fresh medium containing cadmium for 6 h to observe the increase in DNA content by flow cytometry.

Western immunoblot analyses—Logarithmically growing cells were seeded at 10^6 per 150-mm dish and incubated for 40 h. Cultures were treated as described above and incubated in fresh medium for 0, 2, 6 and 24 h at 37°C. Cells were harvested by trypsinization, washed

once in phosphate-buffered saline, and resuspended in lysis buffer (10 mM sodium phosphate buffer [pH 7.2], 1 mM EDTA, 1 mM EGTA, 150 mM NaCl, and 1% NP-40, supplemented with 10 mM 4-(2-aminoethyl) benzenesulfonyl fluoride, 10 mM β -glycerophosphate, 10 mM sodium orthovanadate, and 10 μ g/ml leupeptin and aprotinin). Protein concentrations were determined using the Bio-Rad DC protein assay (Bio-Rad Laboratories). Samples containing equal amounts of protein were mixed with an equal volume of 2 \times Laemmli sample buffer (125 mM Tris-HCl [pH 6.8], 4% sodium dodecyl sulfate (SDS), 20% glycerol) containing 5% β -mercaptoethanol, boiled, and proteins separated by SDS-polyacrylamide gel electrophoresis (SDS-PAGE). Proteins were transferred to nitrocellulose and probed with antibodies against p21^{Cip1/Waf1} (Ab-11, Lab vision), p53 (Ab-6, Lab Vision), phospho-ser15 p53 (Cell Signaling), Chk1 (G4, Santa Cruz Biotechnology), phospho-Ser317 Chk1 (Cell Signaling), Chk2 (Upstate Biotechnology Inc.), phospho-Thr68 Chk2 (Cell Signaling), actin (Santa Cruz Biotechnology), caspase3 (Upstate Biotechnology Inc.), PARP (Upstate Biotechnology Inc.), and GADD45 α (Santa Cruz Biotechnology).

DNA synthesis assay—Logarithmically growing cells were plated at a density of 2×10^5 cells per 60 mm dish and grown at 37°C for 24 h. The time course of inhibition of DNA synthesis in all cell lines was measured by adding 40 and 80 μ M cadmium to culture medium and at various times later pulse-labeling DNA with 10 μ Ci/ml [³H]-thymidine for 30 min. During the 4 h treatment time, DNA synthesis was measured every hour. After the 4 h treatment, the cadmium was removed and DNA synthesis was measured at 1, 2, 6, 12 and 24 h later. The DNA synthesis assay was described in detail elsewhere (Kaufmann *et al.*, 2003) with some modifications. Briefly, radioactive medium was removed and plates washed twice with cold PBS before adding 3 ml of cold 4% trichloroacetic acid (TCA) and incubating at 4°C for 30 min. After washing the plates with cold 4% TCA at 4°C, the fixed cells were dissolved in 0.4 M NaOH and transferred to test tubes. The Abs₂₆₀ was measured to estimate nucleic acid content. Acid-insoluble material was then collected on GF/C microfiber glass filters for measurement of radioactivity by liquid scintillation spectrometry. Net ³H radioactivity was normalized for cell number (the concentration of nucleic acids). Mean ³H/nucleic acid ratios from triplicate cultures were determined as DNA synthesis rates.

Statistical analysis

Statistical evaluation was done using the SPSS 11.5 software (SPSS Inc., Lead Technologies, Chicago, IL, USA). In all cases, a P value < 0.05 was considered to represent a significant difference. All data represent the means \pm SEM of three or more replicates. Student's t-test or ANOVA followed by Dunnett's multiple comparison test was used, as appropriate, to define differences between experimental groups and controls.

Results

Induction of DNA damage by cadmium

The immediate effects of 4 h cadmium treatment on DNA integrity in fibroblasts are shown in Figure 1. DNA damage was assessed by measuring the percent of nuclei with comet tails ≥ 25 μ m. Treatment with 4.5 Gy of IR as a positive control to produce DNA single- and double-strand breaks substantially increased the fraction of nuclei with tails. Incubation with cadmium resulted in a dose-dependent increase of comets of tail length ≥ 25 μ m in normal, p53-defective and AT cells. Significant differences were observed between the untreated cells and those treated with 60 and 80 μ M of cadmium. These results support previous studies showing induction of DNA damage in cadmium-treated human lung fibroblasts and HEPG2 cells (Fatur *et al.*, 2002; Mouron *et al.*, 2004).

Cytotoxicity of cadmium in normal, AT and p53-defective cells

The cytotoxicity of cadmium was determined by colony formation assay. Treatment with cadmium inhibited single cell colony formation by the F1-hTERT, F3-hTERT, F10-hTERT and GM02052A cell lines with similar dose-kinetics (Fig. 2A). For these lines a concentration of 60 μM cadmium inactivated colony formation by about 40%. p53-defective cells were resistant to cadmium in comparison with normal cells, as shown in Figure 2B. The colony formation efficiencies of cells that were treated with 80 μM cadmium were 75% for F10-hTERT-p53-RNAi and 50% for F10-hTERT-p53-H179Q as compared to 21% for F10-hTERT and 20% for F10-hTERT-LX1N.

Sensitivity of G1 and S phase cells to cadmium

In order to compare sensitivity in G1 and S phase cells, F1-hTERT, F3-hTERT and F10-hTERT fibroblasts were released from G0 and the proliferation kinetics of the three cell lines were measured (Fig. 3A). From 8 to 12 h after release from confluence arrest, at least 80% cells were in the pre-replicative G1 phase. Very low fractions of cells were in S or M during this time. After 12 h, cells entered the S phase so that by 20 h about 60% of cells were in S. The fractions of cells in G2 and M remained low at 20 h, but by 24 h increased fractions of G and M cells were seen. According to these results, cells were treated at 8 or 20 h after release when proliferating cells were predominantly in G1 or S. All three cell lines were found to be more sensitive to cadmium when in S than when in G1 (Fig. 3B). In S phase cultures 80 μM cadmium inhibited colony formation by 81–86% while in G1 phase cultures this concentration inactivated colony formation by only 35–41%.

Cadmium inhibits DNA synthesis

A previous study using Chinese hamster ovary cells showed that cadmium inhibited replicative DNA synthesis (Banfalvi *et al.*, 2000). To examine the inhibition of DNA synthesis in diploid human fibroblasts, incorporation of [³H]-thymidine was monitored during and after incubation with 40 or 80 μM cadmium. Initially the effect of cadmium on DNA synthesis was measured in exponentially growing cells. Normal cells, p53-defective cells and AT cells were all sensitive to inhibition of incorporation of [³H]-thymidine by cadmium (Fig. 4). Cadmium produced a dose- and time-dependent inhibition of DNA synthesis during 4 h incubation. At the end of the 4 h treatment with 80 μM , DNA synthesis in F10-hTERT, F10-hTERT-LX1N, F10-hTERT-p53-RNAi and F10-hTERT-p53-H179Q was 7%, 17%, 12% and 15% of control, respectively. After the 4 h incubation, cadmium was removed and DNA synthesis was measured at various times later. Fig. 4 shows that p53 affected the kinetics of DNA synthesis post-treatment. In normal cells DNA synthesis recovered to 50–80% of control at 6–12 h and then decreased again by 18 and 24 h after treatment with cadmium. In contrast, while p53-defective cells also recovered DNA synthesis they did not display the secondary inhibition. At 24 h after treatment with 80 μM cadmium, DNA synthesis rates in F10-hTERT and F10-hTERT-LX1N were 17% and 21% of control in comparison with 63% and 52% in F10-hTERT-p53-RNAi and F10-hTERT-p53-H179Q, respectively. Thus, relative DNA synthesis rates 24 h after treatment with cadmium were similar to relative colony formation efficiencies.

It was conceivable that cadmium simply interfered with the uptake of [³H]-thymidine and its conversion to [³H]-TTP. DNA synthesis was measured using a method that did not require uptake and metabolic conversion of a labeled precursor. Aphidicolin was used to collect cells at the beginning of S phase (Kaufmann and Wilson, 1990). After release from aphidicolin the increase in nuclear DNA content served as a measure of DNA synthesis. The flow cytometric profiles of DNA content in untreated and cadmium-treated cells are shown in Fig. 5A. The DNA content of control cells increased from 2N to 3–4N by 6 h after release as an index of DNA synthesis. Incubation of released cells with 40 or 80 μM cadmium for 6 h inhibited DNA synthesis as evidenced by the reduced fraction of nuclei with 3–4N DNA, and this effect was

greater with the higher dose. F1-hTERT, F3-hTERT and F10-hTERT behaved similarly in this assay with 80 μ M cadmium producing a severe (>90%) inhibition of DNA replication (Fig. 5B).

Rapid and delayed G2 arrest in cadmium-treated fibroblasts

To obtain a more comprehensive view of the effects of cadmium on cell proliferation, flow cytometry was used to quantify cells in G1, S, G2 and M. Analysis of BrdU incorporation demonstrated the expected moderate-to-severe inhibition of DNA replication after 4 h treatment with cadmium (Fig. 6A). Instead of the well-defined arc of positively stained nuclei arrayed between 2N and 4N DNA content as seen in controls, the cadmium-treated cells displayed reduced intensity of labeling with BrdU throughout S phase. This pattern has been observed before in human cells after treatment with cytotoxic fluences of ultraviolet light that severely inhibit DNA replication (Cistulli and Kaufmann, 1998;Cordeiro-Stone *et al.*, 2002). The analysis of cell cycle progression after cadmium exposure also demonstrated both rapid and delayed G2 arrest in cadmium-treated fibroblasts (Fig. 6A). At the end of the 4 h treatment with cadmium, the fraction of mitotic cells was reduced in a dose-dependent fashion with 80 μ M producing nearly 80% reduction in the fraction of mitotic cells. This reduction would appear to be due to a G2 arrest blocking the movement of cadmium-damaged cells into mitosis. AT and p53-defective cells showed a similar reduction of mitotic cells after 4 h cadmium treatment (Fig. 6B). Thus, the rapid G2 arrest induced by cadmium was independent of ATM and p53.

By 6 h after the 4 h cadmium treatment, the mitotic index of all cell lines recovered with mitotic indices equal to or greater than untreated cells (Fig. 6A, B). Incorporation of BrdU by S phase cells also recovered as quantitatively demonstrated in Figure 4. However, by 24 h after the 4 h cadmium treatment, a significantly increased fraction of normal cells accumulated in G2 phase, with a corresponding significant secondary decrease in mitotic cells. Analysis of BrdU incorporation at 24 h revealed that after the severely toxic dose of 80 μ M, 4% of cells with 2–4 N DNA content were unable to incorporate BrdU, suggestive of full replicative arrest. The p53-defective cells displayed significantly less accumulation in G2 phase, and no inhibition of mitosis at 24 h after treatment (Fig. 6A, B).

Western immunoblot analysis of cell cycle checkpoint protein expression

To probe the mechanisms of cadmium-induced cell cycle arrest, western immunoblot analyses were performed (Fig. 7). Cadmium increased p53 protein and phospho-ser15-p53 in both normal and AT cells (Fig. 7A) indicating that activation of p53 was independent of ATM. Gadd45 α protein levels were increased immediately after cadmium treatment in all cell lines but declined by 6 h after treatment and were undetectable at 24 h (Fig. 7B). In spite of the severe inhibition of DNA replication and G2 arrest induced by cadmium, the levels of Chk1, Chk2, phospho-Ser317 Chk1, phospho-Thr68 Chk2 and p21^{CIP1/WAF1} did not change in cadmium-treated normal human fibroblasts (Fig. 7C). There was no evidence of apoptosis at 24, 48 and 66h after treatment with 80 μ M cadmium as indicated by expression only of intact PARP and caspase 3 (not shown).

Discussion

Cadmium has been found to be carcinogenic to humans and other mammals. However, the mechanisms involved in the carcinogenicity are still unclear (Waalkes, 2003; Waisberg *et al.*, 2003). Indirect genotoxic effects of cadmium have been discussed, e.g. interference with DNA repair and DNA replication processes (Hartwig, 1994; Dally and Hartwig, 1997). We tested the hypothesis that cadmium could cause DNA damage and induce DNA damage checkpoint responses in human fibroblast lines with selected genetic alterations. The survey

of the effects of cadmium on DNA metabolism in human fibroblasts confirmed the induction of DNA damage and demonstrated a role for p53 in the cellular responses to cadmium. Cadmium-induced cytotoxicity and the delayed inhibition of DNA replication and mitosis were p53-dependent, and the treatment with cadmium induced p53 protein accumulation and phosphorylation of p53 at ser 15. A high concentration of cadmium also produced a severe inhibition of DNA synthesis in S phase cells and reduced the fraction of mitotic cells seemingly by GADD45 α -associated G2 arrest. The results demonstrated profound disturbance in human DNA metabolism and cell division as provoked by acute exposure to toxic concentrations of cadmium.

DNA damage checkpoints are biochemical pathways that delay or arrest cell cycle progression in response to DNA damage (Kaufmann and Paules, 1996; Kastan and Bartek, 2004). It is assumed that checkpoints maintain cell cycle arrest until damaged DNA is repaired when the arresting signal is attenuated or reversed to permit resumption of cell cycle progression (Abraham, 2001). For the rapid G2 arrest, DNA damage likely occurred in G2 cells, as a significant decrease in the mitotic index was observed immediately after 4 h cadmium treatment (Fig. 6A). The rapid induction of Gadd45 α could explain the cadmium-induced G2 arrest. Gadd45 α is known to inhibit mitosis-promoting-factor (cyclin B1/Cdk1) kinase activity and enforce a G2 checkpoint response to DNA damage (Wang *et al.*, 1999; Zhan *et al.*, 1999; Yang *et al.*, 2000). Upon removal of cadmium, Gadd45 α levels diminished and mitosis resumed. Inactivation of p53 did not abrogate the rapid G2 arrest induced by cadmium nor the induction of Gadd45 α (Fig. 6B). Cadmium is known to induce phosphorylation of C-Jun apparently through activation of c-Jun NH₂-terminal kinase (Lee *et al.*, 2001) and p53-independent induction of Gadd45 α after UV-treatment was shown to include mitogen-activated protein kinase interaction with Oct1 and NF-YA transcription factors (Tong *et al.*, 2001). Thus, the ATM- and p53-independent induction of Gadd45 α and G2 arrest by treatment with cadmium appears to be mediated by mitogen-activated protein kinase signaling.

Cadmium is also a spindle poison; it depolymerizes microtubules and actins (Li *et al.*, 1993; Wang and Templeton, 1996) and previous studies have shown that G2/M phase cells are more sensitive to challenge with cadmium (Chao and Yang, 2001). Mitotic arrest may thus occur in the presence of cadmium. If the mitotic spindle is a major target for cadmium, a high proportion of cells should be stopped in mitosis after treatment with cadmium. We found that the proportion of mitotic cells decreased in a concentration-dependent manner during cadmium exposure (Fig. 6) implying that cells completed mitosis in the presence of cadmium and the mitotic compartment emptied behind the G2 arrest. This result suggests that microtubules in the mitotic spindle apparatus may not be a major target for cadmium action in diploid human fibroblasts.

p53 protein and phospho-ser15-p53 were induced by cadmium in normal and AT cells. ATM and ATR phosphorylate ser15 of p53 directly and ser20 through activation of Chk2 or Chk1 (Banin *et al.*, 1998; Ryan *et al.*, 2001). Phosphorylation of p53 inhibits its export and degradation, thus increasing its level of expression (Zhang and Xiong, 2001). Cadmium has been shown to induce p53 and phospho-ser15 phosphorylation in MCF-7 cells (Matsuoka and Iqisu, 2001). Cadmium is also known to replace zinc in the p53 zinc-finger domain, altering the structure of p53 and inhibiting DNA binding (Meplan *et al.*, 1999). Accordingly, cadmium inhibited the induction of p53 and p21^{Cip1/Waf1} in human cells treated with the carcinogen, benzo[a]pyrene diolepoxide I (Mukherjee *et al.*, 2004). The cadmium-induced inhibition of p53 was dose-dependent and at 10–30 μ M concentrations, p53 transactivation activity in mouse cells was inhibited by 65–85% (Meplan *et al.*, 1999). We anticipate that metallothionein expression should reduce the concentration of free cadmium inside cells so that p53 function was only attenuated, not fully ablated in normal human fibroblasts. Moreover, the effect of cadmium on p53 was reversible such that by 12 h after removal from culture medium, p53

binding to DNA was restored by 50% (Meplan *et al.*, 1999). Thus, although induction of p21^{Cip1/Waf1} was not evident in cadmium-treated cells, some function of p53 remained to inactivate clonal expansion and produce the delayed inhibition of DNA synthesis and G2 arrest.

By 24 h after treatment, cadmium induced a p53-dependent but ATM-independent G2 arrest. Gadd45 α had returned to control levels and p21^{Cip1/Waf1} was unchanged in cadmium-treated cells (Fig. 7), so Gadd45 α and p21^{Cip1/Waf1} do not appear to contribute to the delayed G2 arrest. p53 activates the transcription of many genes that mediate its downstream functions (Zhou *et al.*, 2006) and can also repress the transcription of different genes through several mechanisms (Taylor and Stark, 2001). Leach *et al.* reported that the activation of wild-type p53 resulted in a down-regulation of Wee1 expression and dephosphorylation of Cdk1 under conditions of p53-induced G2/M growth arrest and p53-mediated apoptosis (Leach *et al.*, 1998). However, Wee1 expression did not change in cadmium-treated cells (data not shown). p53 may suppress the G2/M transition by negatively regulating the expression of cyclin B1, Cdk1, and topoII-alpha (Passalaris *et al.*, 1999; Taylor and Stark, 2001). Cyclin B1 and Cdk1 are subunits of mitosis-promoting-factor that is required for entry to mitosis, and topoII-alpha is required for timely chromatid decatenation to allow bypass of a decatenation checkpoint that acts in G2 cells (Deming *et al.*, 2001; Deming *et al.*, 2002). Further experiments are needed to determine whether p53 trans-repression of these G2-regulated target genes accounts for the cadmium-induced delayed G2 arrest.

The mechanisms of inhibition of DNA replication by cadmium remain to be determined. Clark and Kunkel reported that cadmium did not inhibit *in vitro* plasmid DNA replication by human cell extracts under conditions that mismatch repair was inhibited (Clark and Kunkel, 2004). This result implies that cadmium is not directly toxic to the basal DNA replication machinery including DNA primase, DNA polymerases, DNA ligase and required accessory proteins such as RPA. The result also indicates that mismatch repair and DNA replication can be uncoupled *in vitro* and that DNA replication does not require mismatch repair. One mechanism for inhibition of DNA replication without direct inhibition of the replication machinery is inhibition of dNTP precursor production. Hydroxyurea inhibits DNA replication by this mechanism. The observation that treatment with HU induced phosphorylation of Chk1 while cadmium did not (Fig. 7C) argues against an inhibitory effect of cadmium on DNA precursor production. Further study will be required to determine whether cadmium inhibits any of the enzymes that are required for initiation of DNA replication at replicon origins. As the known mechanisms of inhibition of replicon initiation by DNA damage require activation of Chk1 and Chk2 (Falck *et al.*, 2001; Heffernan *et al.*, 2002), the failure of cadmium to activate Chk1 and Chk2 suggests that the effect of cadmium is not through a recognized S checkpoint.

It is of interest to compare and contrast the mechanisms of growth arrest by cadmium and another carcinogenic metal, chromium. Both metals appear to induce oxidative stress (Valko *et al.*, 2006), activate the mitogen-activated protein kinases p38, JNK and ERK (Chuang and Yang, 2001), and induce inhibition of DNA replication and mitosis (Wakeman *et al.*, 2005). However, while the effects of cadmium on DNA synthesis and mitosis were independent of ATM signaling, chromium activated ATM apparently by inducing DNA dsb in S phase cells (Ha *et al.*, 2004). In keeping with the induction of DNA dsb and activation of ATM, chromium also activated Chk2 (Ha *et al.*, 2003); cadmium did not activate Chk2 even when DNA replication was severely inhibited. Chromium caused ATM-dependent apoptosis in human fibroblasts and AT fibroblasts were more sensitive to inactivation of colony formation by chromium (Ha *et al.*, 2003); cadmium did not induce apoptosis in foreskin fibroblasts and AT fibroblasts were not hypersensitive to cadmium. The enhanced growth arrest and inactivation of colony formation in chromium-treated AT cells may reflect signaling from ATR to p53 (Wakeman and Xu, 2006). Chromium induced GADD45 mRNA (Ceryak *et al.*, 2004) and cadmium induced GADD45 α protein consistent with both compounds causing oxidative stress

and activating the stress-responsive mitogen-activated protein kinases. This comparison reveals similarities and dissimilarities in the mechanisms of action of cadmium and chromium suggesting that unique properties of the metal salts may contribute to their toxicities. The facility with which cadmium replaces zinc in important proteins such as p53, XPA and hMSH2 may contribute to its biological effects.

In summary the results presented here suggest a model in which cadmium-induced DNA damage or oxidative stress causes p53-independent induction of GADD45 α to produce a rapid G2 arrest, and p53-dependent trans-repression of downstream target genes to produce a delayed G2 arrest.

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Abbreviations

APC	aphidicolin
AT	ataxia telangiectasia
ATM	ataxia telangiectasia-mutated
BrdU	5'-bromo-2'-deoxyuridine
dsb	double-strand breaks
hTERT	human telomerase reverse transcriptase
SDS	sodium dodecyl sulfate
TCA	trichloroacetic acid

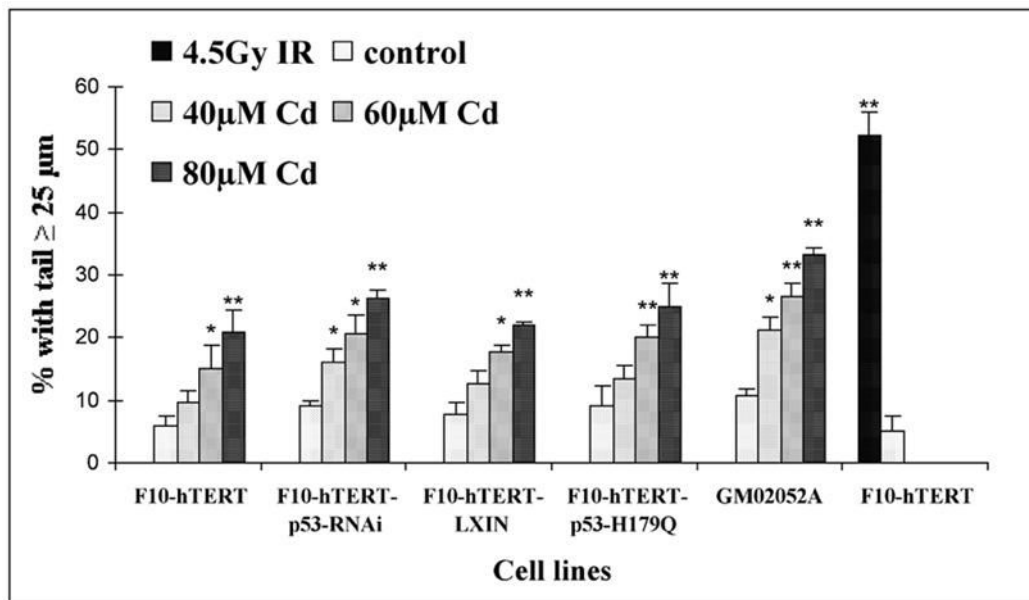


Figure 1.

Cadmium-induced DNA damage in human fibroblasts. At the end of the 4 h incubation with 40, 60 and 80 μM cadmium, fibroblasts were removed from the plates with trypsin and collected for comet assay. The percentages of cells with a comet tail $\geq 25 \mu\text{m}$ were determined as a measure of DNA damage. Fifty cells were analyzed per slide. Mean values of three independent experiments \pm S.E. are presented. (*) Denotes values that were significantly different from the untreated control culture (* $p < 0.05$; ** $p < 0.01$). The culture that was treated with 4.5 Gy IR was harvested immediately after irradiation.

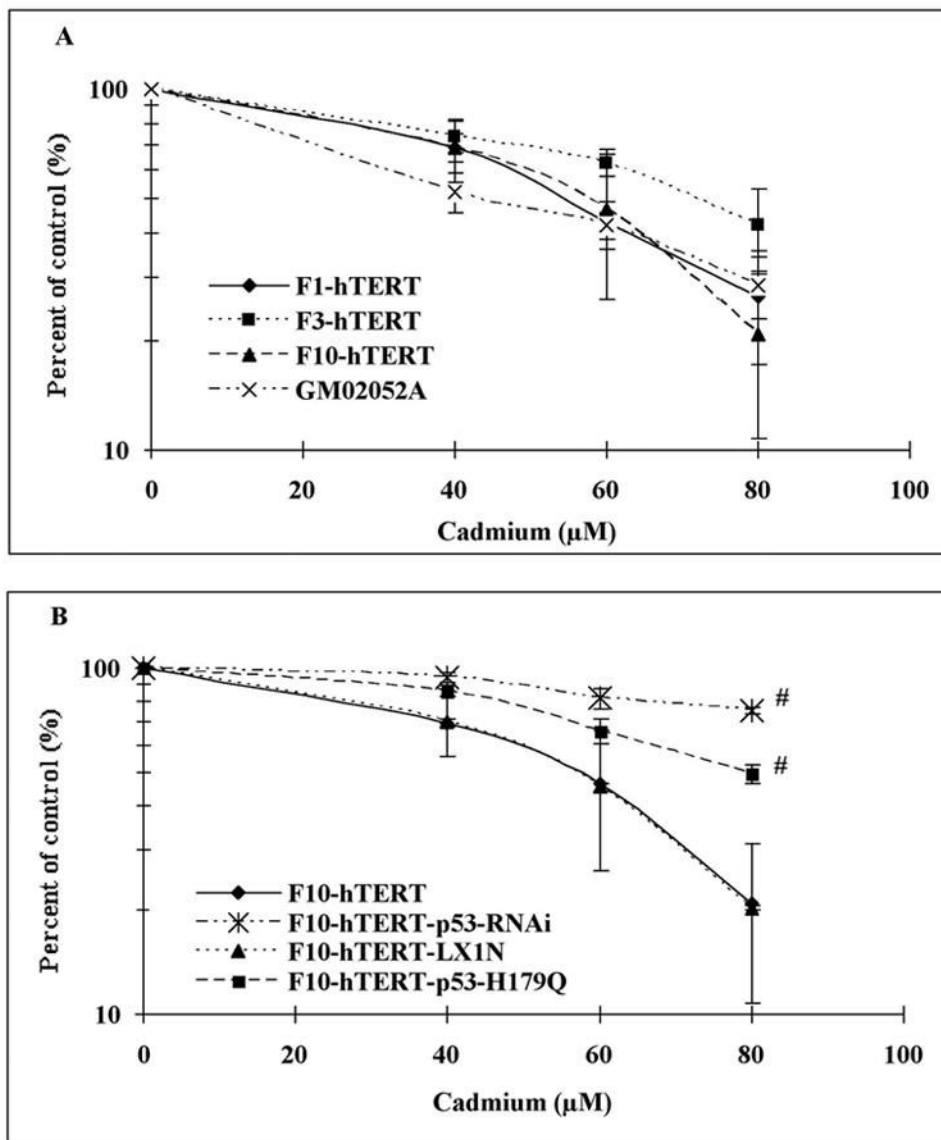


Figure 2. Inactivation of clonogenic survival by cadmium. Colony formation was measured in logarithmically growing cells, plated at 500 cells per 100 mm dish and incubated 8 h before the 4 h cadmium treatment. Fibroblast lines were treated with 0, 40, 60 and 80 μM cadmium, and after a 14- day incubation colonies were stained with crystal violet and counted. Results show the mean colony formation in cadmium-treated cultures relative to the untreated control (Mean \pm SE, n=3). (A) Colony formation by F1-hTERT, F3-hTERT, F10-hTERT and GM02052A after cadmium treatment. (B) Colony formation in normal and p53-defective cells after cadmium treatment. (#) denotes both F10-hTERT-p53-RNAi and F10-hTERT-p53-H179Q were significantly different than F10-hTERT and F10-hTERT- LX1N ($p < 0.05$).

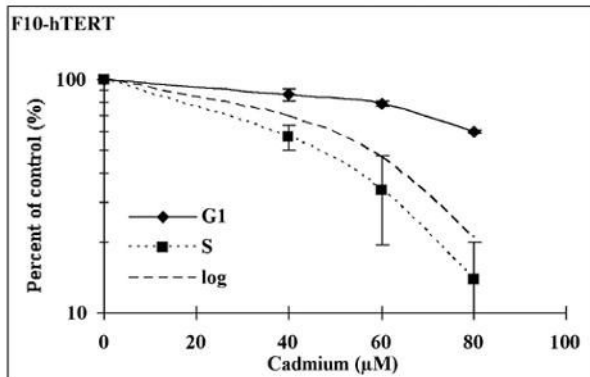
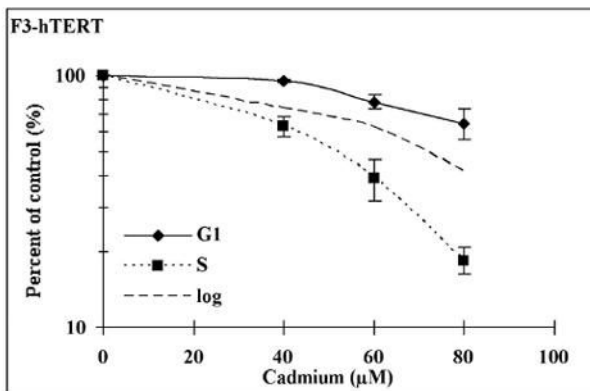
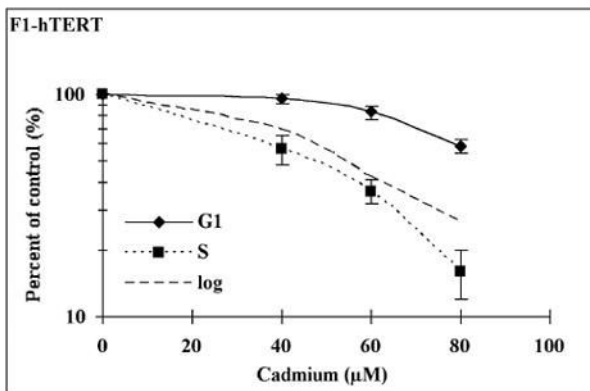
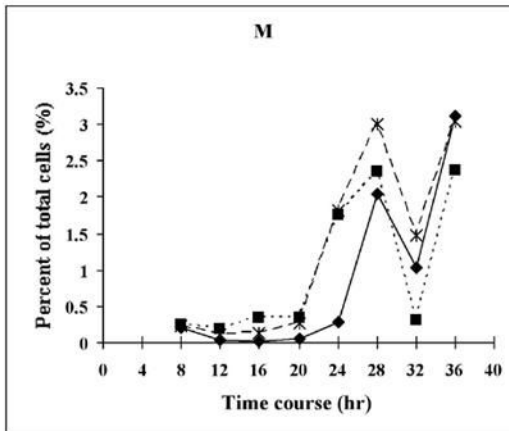
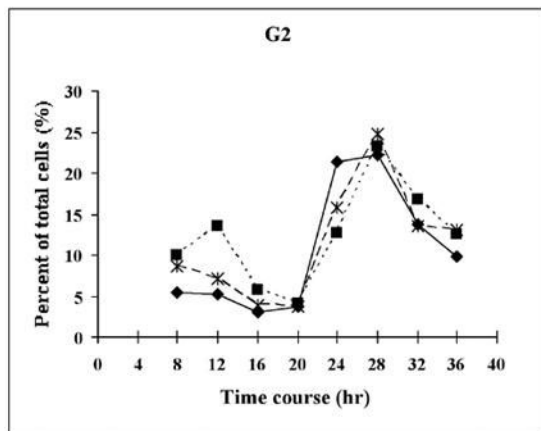
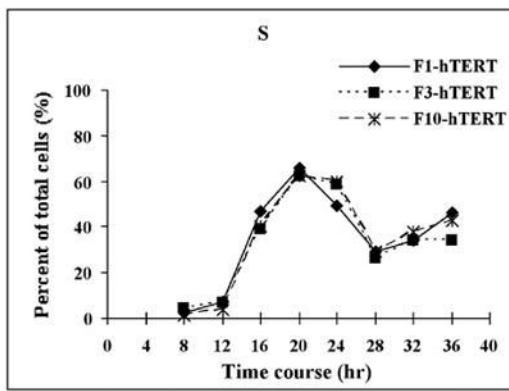
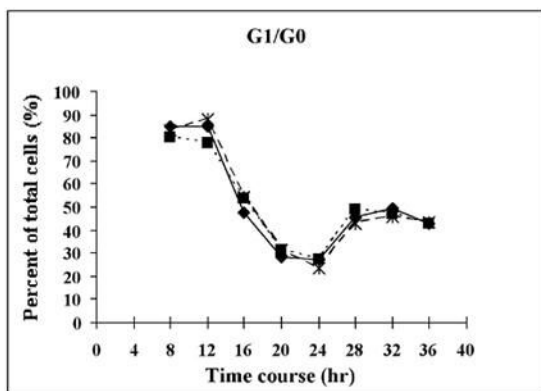


Figure 3.

Sensitivity to cadmium of G1 and S phase cells. Cells were allowed to grow for 8 days to confluence arrest (G0 phase). (A) F1-hTERT, F3-hTERT and F10-hTERT fibroblasts were released from G0 and the proliferative activity of the three cell lines was measured by flow cytometry. (B) Confluence-arrested cells were released for 8 h to allow them to reach G1 and 20 h to allow them to reach S phase. The cells were then treated with different concentrations of cadmium for 4 h. The colonies were counted after a 14-day incubation. Results show the mean relative colony formation in cadmium-treated cultures (Mean \pm SE, n=3).

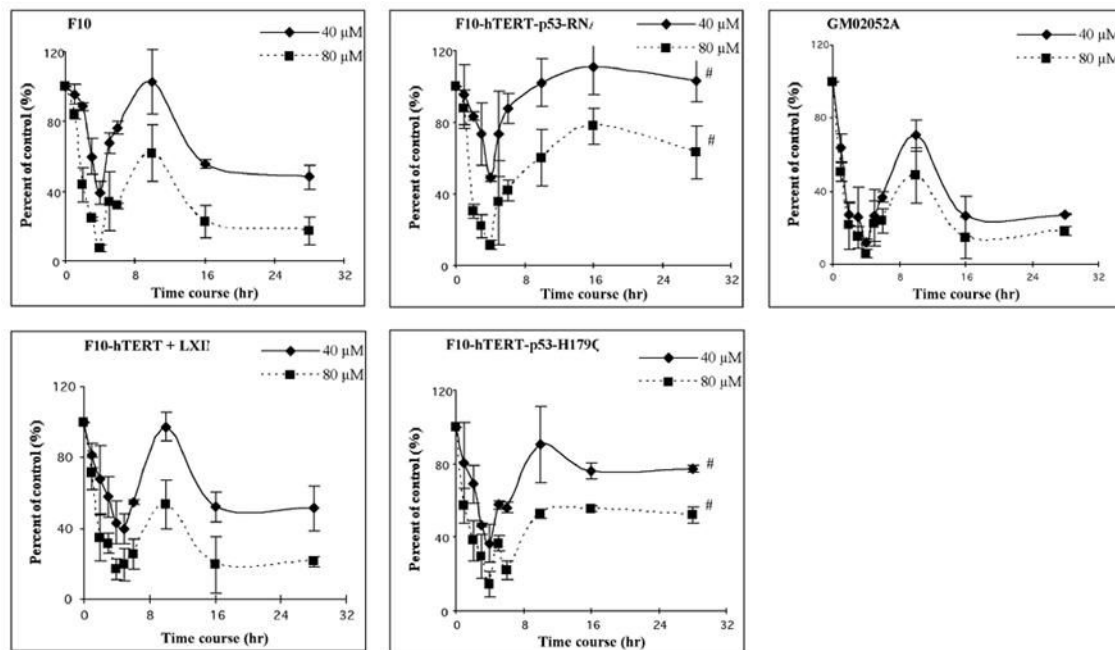
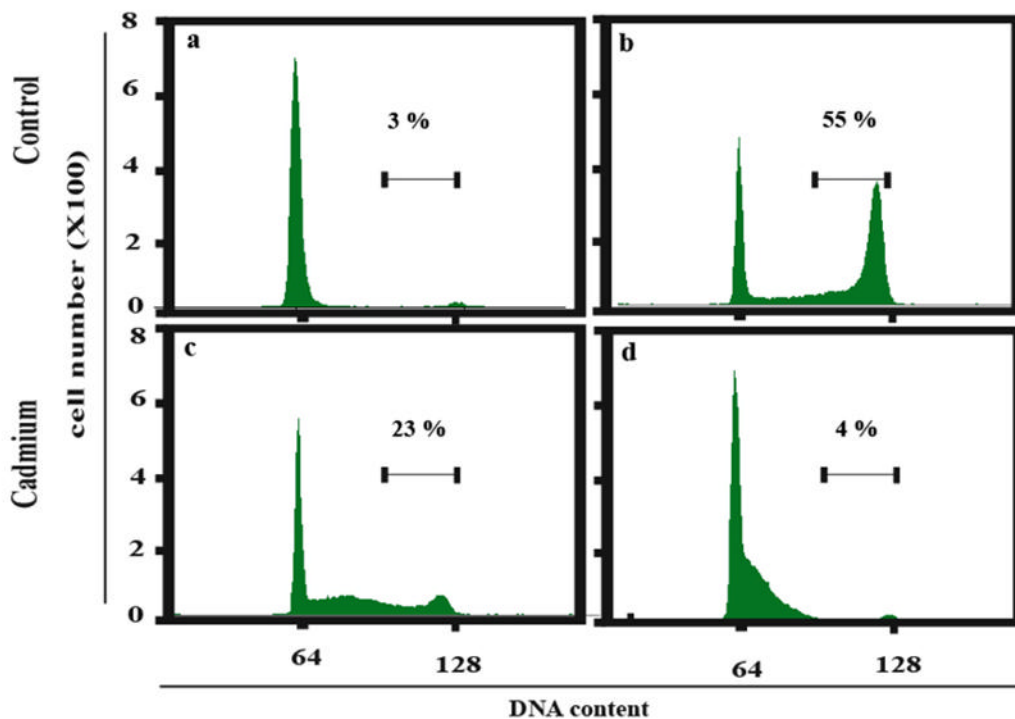


Figure 4.

Cadmium inhibits DNA synthesis. The human fibroblast lines, F10-hTERT, F10-hTERT-LX1N, F10-hTERT-53-RNAi, F10-hTERT-p53-H179Q and GM02052A, were treated with cadmium during log-phase growth. During the 4 h treatment, DNA synthesis was measured every hour by incubation with [3 H] thymidine for the final 30 min. After the 4 h treatment, the cadmium was removed, fresh medium added and DNA synthesis was measured at 1, 2, 6, 12 and 24 h later. Net [3 H] radioactivity was normalized to cell number. Normalized [3 H] CPM were graphed as a percent of time-matched, sham-treated controls. (#) denotes both F10-hTERT-p53-RNAi and F10-hTERT-p53-H179Q were significantly different than F10-hTERT and F10-hTERT- LX1N ($p < 0.05$).

A



B

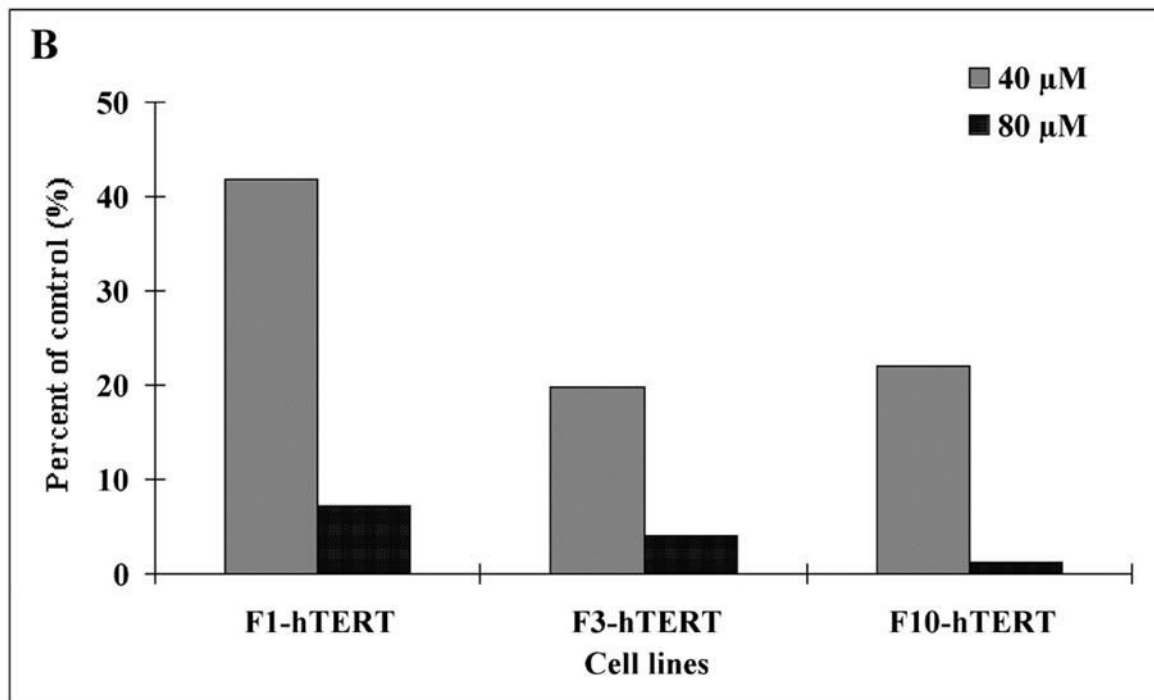
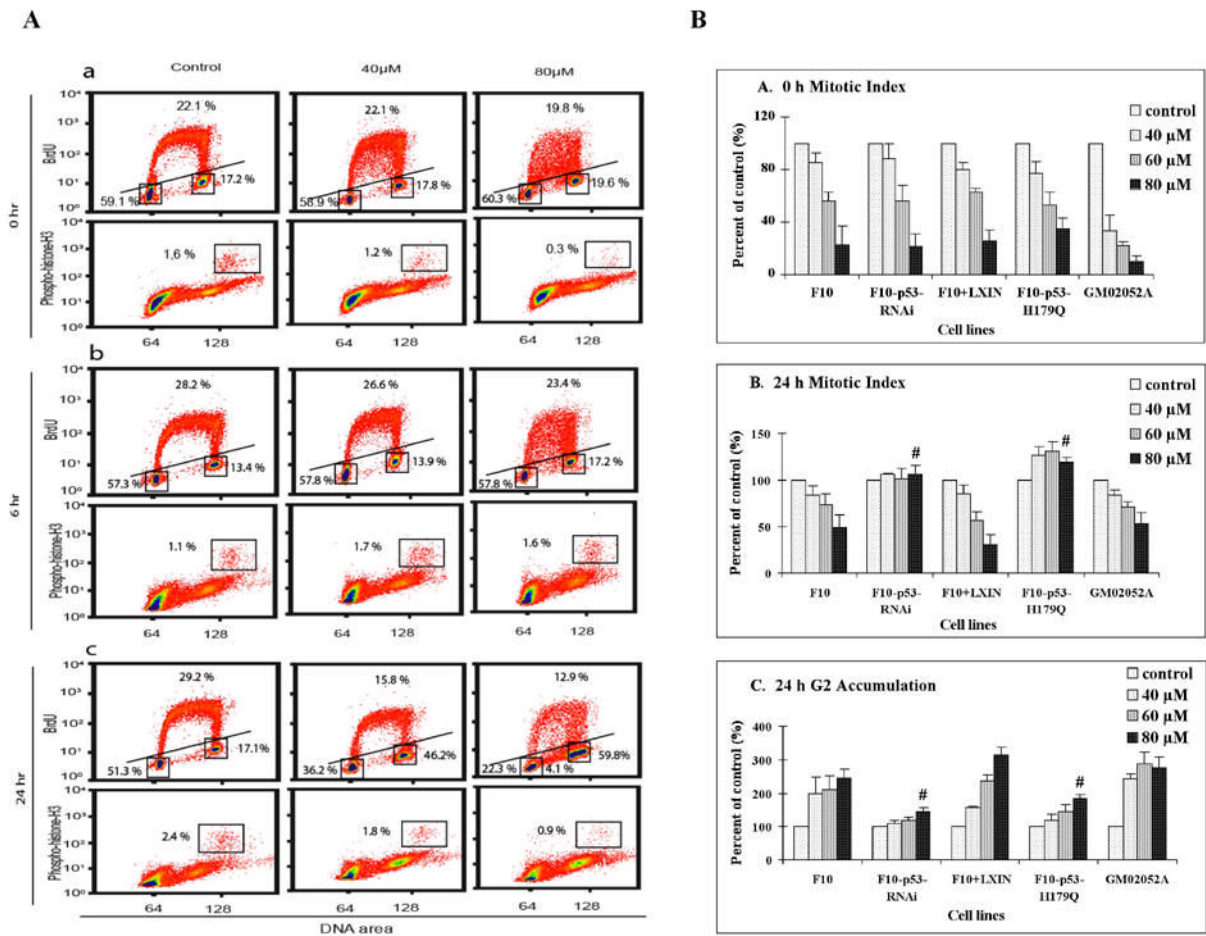


Figure 5. Inhibition of DNA synthesis in S phase cells. Cells were released from confluence-arrest, reseeded at 1 million cells per 100 mm dish and incubated with aphidicolin for 24 h to collect cells at the beginning of S phase. Aphidicolin was then removed and cells incubated in fresh

medium with or without cadmium for 6 h to observe the increment of DNA content during S phase progression. Propidium iodide was used to stain DNA. The stained nuclei were analyzed by flow cytometry to determine their DNA content. (A) F1-hTERT cells were released from synchronization and treated with various concentrations of cadmium. Numbers above the bar (3–4 N DNA content) represent the fraction of total cells in this region. (a) Cells synchronized for 24 h with aphidicolin. (b) Cells incubated for 6 h without cadmium after removal of aphidicolin. (c) Cells incubated 6 h with 40 μM cadmium after removal of aphidicolin. (d) Cells incubated 6 h with 80 μM cadmium after removal of aphidicolin. (B) The percentage of nuclei with 3–4 N DNA content after 6 h incubation with 40 or 80 μM cadmium was expressed as a percentage of the sham-treated control.

**Figure 6.**

Inhibition of DNA synthesis and mitosis by cadmium. Logarithmically growing cells were treated with cadmium for 4 h. At 0, 6 and 24 h after 4 h cadmium or sham treatment, cells were incubated with 10 μ M BrdU for 2 h, then cells were harvested for analysis of BrdU-incorporation by flow cytometry. At 0, 6 and 24 h after cadmium- or sham-treatment, cells were also harvested and processed for determination of mitotic index by flow cytometry. Propidium iodide was used to stain DNA (X-axis). Anti-BrdU-FITC was used to stain BrdU-labeled nuclei and antibody specific for phospho-histone H3 was used to stain mitotic cells. (A) F10-hTERT cells were analyzed 0, 6 and 24 h after cadmium treatment. Boxes and lines separate G1, S, G2 and M cells with the percentages of cells in these phases noted. (B) The cell cycle phase distributions of F10-hTERT, F10-hTERT-p53-RNAi, F10-hTERT-LX1N, F10-hTERT-p53-H179Q and GM02052A cells were analyzed after cadmium treatment. Results show the mean \pm SE of three independent experiments. (a) Relative mitotic index immediately after cadmium treatment. (b) Relative mitotic index 24 h after cadmium treatment. (c) Relative percentage of cells in G2 24 h after cadmium treatment. (#) denotes both F10-hTERT-p53-RNAi and F10-hTERT-p53-H179Q were significantly different than F10-hTERT and F10-hTERT-LX1N ($p < 0.05$).

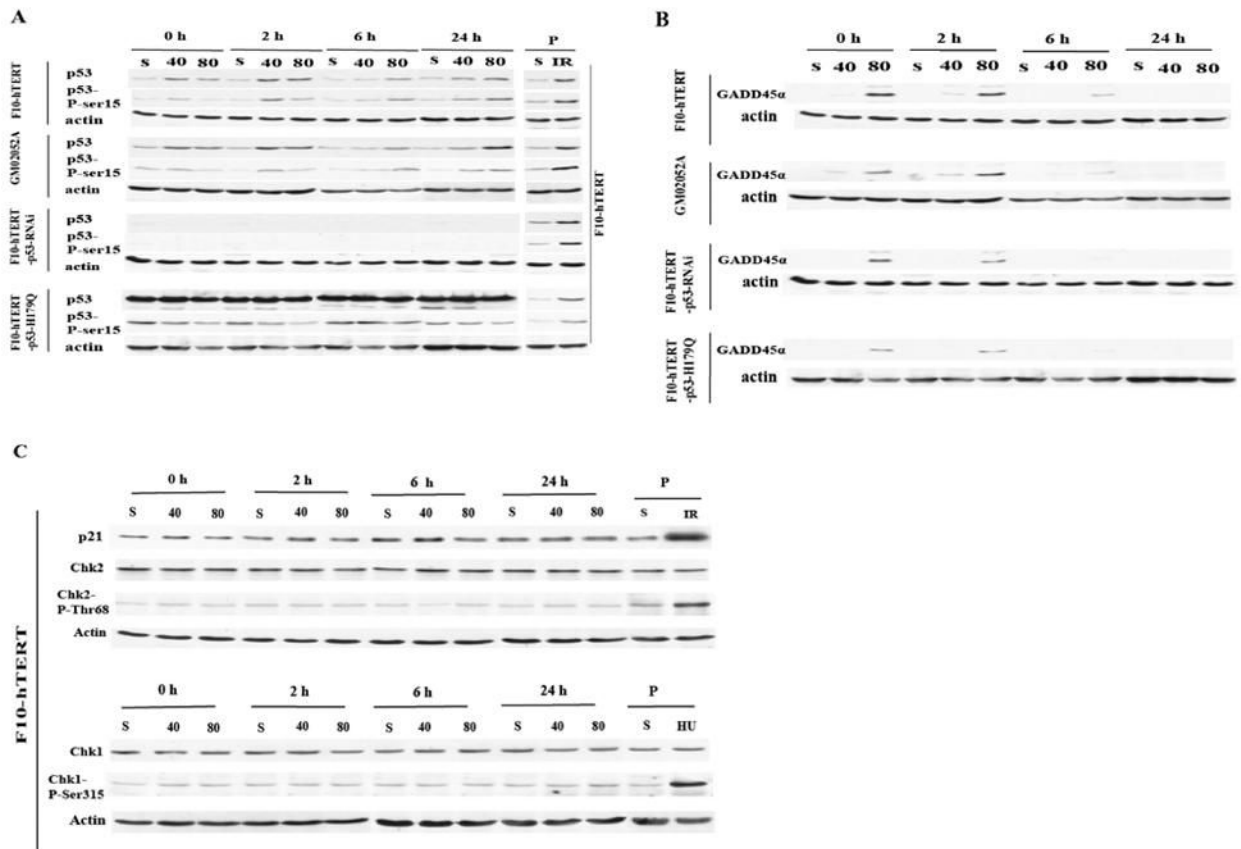


Figure 7.

Western immunoblot analysis of DNA damage response in cadmium-treated human fibroblasts. Cells were incubated in fresh medium for 0, 2, 6 and 24 h after 4 h treatment with 40 or 80 μ M cadmium or solvent control (S), then harvested and lysed. Protein extracts were prepared and 100 μ g of total protein analyzed for expression of selected proteins or phosphorylated species. (A). Expression of p53 and phospho-ser-15-p53. The positive control (P) was F10-hTERT cells harvested 6 h after 1.5 Gy IR. (B). Expression of Gadd45 α . (C). Expression of p21^{Cip1/Waf1}, Chk2, phospho-Thr68-Chk2, Chk1 and phospho-ser317-Chk1. The positive control (P) for p21^{Cip1/Waf1}, Chk2 and phospho-Thr68-Chk2 was F10-hTERT cells harvested 6 h after 1.5 Gy IR. The positive control (P) for Chk1 and phospho-ser317-Chk1 was F10-hTERT cells harvested after 24 h incubation with 2 mM hydroxyurea.