

# Cyclopentenyl Cytosine Induces Senescence in Breast Cancer Cells through the Nucleolar Stress Response and Activation of p53<sup>S</sup>

Min Huang, Patrick Whang, Patrick Lewicki, and Beverly S. Mitchell

*Department of Medicine, Divisions of Oncology and Hematology, and the Stanford Cancer Center, Stanford University School of Medicine, Stanford, California*

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## ABSTRACT

The induction of senescence has emerged as a potentially important contributor to the effects of chemotherapeutic agents against tumors. We have demonstrated that depletion of CTP induced by cyclopentenyl cytosine (CPEC; NSC 375575), a specific inhibitor of the enzyme CTP synthetase, induces irreversible growth arrest and senescence characterized by altered morphology and expression of senescence-associated  $\beta$ -galactosidase activity in MCF-7 breast cancer cells expressing wild-type p53. In contrast, differentiation in the absence of senescence resulted from CPEC treatment in MDA-MB-231 breast cancer cells that express a mutated p53. Both senescence of MCF-7 cells and differentiation of MDA-MB-231 cells were prevented by repletion of CTP through the cytidine salvage pathway. Senescence in MCF-7 cells was associated with

a G<sub>2</sub>- and S-phase arrest, whereas differentiation in MDA-MB-231 cells was associated with arrest in G<sub>1</sub> phase at 5 days. Mechanistic studies revealed that CTP depletion induced a rapid translocation of nucleolar proteins, including nucleostemin and nucleolin into the nucleoplasm. This nucleolar stress response resulted in a sustained elevation of p53 and the p53 target genes, p21 and Mdm2, in cells with wild-type p53. Furthermore, short interfering RNA-induced knockdown of p53 in MCF-7 cells treated with CPEC prevented cellular senescence and increased apoptotic cell death. We conclude that CTP depletion and the resulting nucleolar stress response results in a senescence-like growth arrest through activation of p53, whereas cells with mutated p53 undergo differentiation or apoptotic cell death.

## Introduction

Transformed cells retain the ability to undergo premature or replicative senescence in response to various stimuli. Senescent cells undergo irreversible cell cycle arrest and display characteristic phenotypic alterations, including an enlarged and flattened morphology and expression of senescence-associated  $\beta$ -galactosidase activity (SA- $\beta$ -gal) (Dimri et al., 1995). Some anticancer agents induce a rapid, non-telomere-dependent form of senescence, often termed premature senescence, in cancer cells at concentrations that do not cause acute cell death in vitro or in vivo (Hornsby, 2007). These agents include phosphoinositide-3 kinase inhib-

itors (Collado et al., 2000), retinoic acid (Roninson and Dormanovic, 2003), microtubule-stabilizing agents (Arthur et al., 2007), Sirt1 (Ota et al., 2006), and DNA topoisomerase inhibitors (Michishita et al., 1998), doxorubicin (Vigneron et al., 2005) and cisplatin (Varna et al., 2009). Induction of SA- $\beta$ -gal staining after chemotherapy has been observed in cells obtained from patients with breast cancer (te Poele et al., 2002), and senescence-like growth arrest has been proposed as a determinant of in vivo tumor response to both selective chemotherapeutic agents and ionizing radiation (Kahlem et al., 2004).

Multiple mechanisms have been implicated in the induction of senescence, including telomere shortening induced by inhibition of TERT activity (Kim et al., 2003), induction of p53 (Chang et al., 1999), increased expression of cyclin-dependent kinase inhibitors such as p16 INK4a (Uhrbom et al., 1997), p21 (Fang et al., 1999) and p27 (Collado et al., 2000), and dephosphorylation of retinoblastoma protein (Xu et al., 1997).

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**ABBREVIATIONS:** SA- $\beta$ -gal, senescence-associated  $\beta$ -galactosidase activity; CPEC, cyclopentenyl cytosine; EGFP, enhanced green fluorescent protein; NS, nucleostemin; PBS, phosphate-buffered saline; PI, propidium iodide; Pol I, RNA polymerase I; FITC, fluorescein isothiocyanate; BSA, bovine serum albumin; siRNA, small interfering RNA.

Cyclopentenyl cytosine (CPEC) is an analog of cytidine (Kang et al., 1989) and enters cells preferentially through the equilibrative nucleoside transporters, including equilibrative nucleoside transporters 1 and 2 (Huang et al., 2004). CPEC is intracellularly metabolized into its mono-, di-, and triphosphate forms (Kang et al., 1989). In its triphosphate form, it serves as a specific noncompetitive inhibitor of *CTP synthetase*, which catalyzes the conversion of UTP to CTP as the predominant pathway for CTP synthesis in proliferating cells (Schimmel et al., 2007). As a result, the intracellular CTP pool is depleted rapidly (Kang et al., 1989; Huang et al., 2004). Previous studies have established the antitumor effects of CPEC using a number of tumor cell lines (Schimmel et al., 2007) and several tumor models (Schimmel et al., 2007; Van Bree et al., 2009). In addition, CPEC was shown to act at nanomolar concentrations in vitro as a chemosensitizer to enhance the anticancer effects of several nucleoside analogs such as 1- $\beta$ -D-arabinofuranosylcytosine and 2',2'-difluorodeoxycytidine (gemcitabine) in a variety of tumor cell lines (Schimmel et al., 2007), although no enhanced efficacy of CPEC was found when combined with gemcitabine and radiation in two animal tumor models (Van Bree et al., 2009).

In the present study, we have found that CTP depletion results in a pronounced nucleolar stress response. The effects of CTP depletion on cell proliferation, differentiation, apoptosis, and cellular senescence and related signaling pathways were assessed. These data provide evidence that nucleolar stress triggered by CTP depletion serves to regulate critical events leading to the induction of senescence or differentiation/apoptosis depending on the p53 status of the cell.

## Materials and Methods

**Cell Culture Conditions and Reagents Used.** The p53 wild-type MCF-7 human breast tumor cell line and p53 mutant MDA-MB-231 cells were obtained from the American Type Culture Collection (Manassas, VA). U2OS cells stably expressing EGFP-tagged nucleostemin (NS) have been described previously (Huang et al., 2009). U2OS, MCF-7, and MDA-MB-231 cells were grown in Dulbecco's modified Eagle's medium supplemented with 10% fetal calf serum (HyClone Laboratories, Logan, UT), 1 mM sodium pyruvate, 0.1 mM nonessential amino acids, and 100 U/ml penicillin and streptomycin at 37°C in a humidified, 5% CO<sub>2</sub> atmosphere. Nutlin-3 was obtained from EMD Biosciences (San Diego, CA). 4',6-Diamino-2-phenylindol was purchased from Invitrogen (Carlsbad, CA). Mouse monoclonal anti-nucleolin (MS-3) and anti-p53 (DO-1) were from Santa Cruz Biotechnology (Santa Cruz, CA). Rabbit polyclonal anti-NPM was from Cell Signaling Technology Inc. (Danvers, MA). Goat anti-nucleostemin polyclonal antibody was from R&D Systems (R&D Systems, Minneapolis, MN). Anti-actin monoclonal antibody was from Sigma-Aldrich (St. Louis, MO). Full-length Av polyclonal EGFP antibody was from Clontech (Mountain View, CA). Fluorescein (FITC)-conjugated donkey anti-rabbit F(ab')<sub>2</sub> and rhodamine-conjugated goat anti-mouse secondary antibodies were purchased from Jackson ImmunoResearch Laboratories Inc. (West Grove, PA). Small interfering RNAs (siRNAs) SMARTpool against p53 was obtained from Dharmacon RNA Technologies (Lafayette, CO). CPEC was kindly provided by Dr. Lee Graves (University of North Carolina at Chapel Hill, Chapel Hill, NC).

**Assay of SA- $\beta$ -gal.** SA- $\beta$ -gal staining was performed as described previously (Dimri et al., 1995). At the appropriate times after treatment, cells were washed twice in phosphate-buffered saline (PBS), fixed in 2% formaldehyde/0.2% glutaraldehyde for 5 min at room temperature, washed again in PBS, and incubated with  $\beta$ -galactosidase staining solution containing 1 mg/ml 5-bromo-4-chloroindol-3-yl

$\beta$ -D-galactopyranoside, 40 mM citric acid/sodium phosphate, pH 6, 5 mM potassium ferrocyanide, 5 mM ferricyanide, 150 mM NaCl, and 2 mM MgCl<sub>2</sub>. After overnight incubation at 37°C, cells were photographed. The extent of senescence was quantified based on the mean number of cells displaying both changes in morphology as well as a percentage of blue-green staining in 8 fields that contained at least 50 cells per field for each experimental condition. All treatments were carried out in triplicate and results represent the mean  $\pm$  S.D.

**BrdU Incorporation and Cell Cycle Analysis.** Cells were incubated in medium containing DMSO or CPEC at the concentrations and times indicated. Cells were then incubated with 100  $\mu$ M BrdU (Sigma-Aldrich) at 37°C for 2 h before harvest. Approximately 10<sup>6</sup> cells were harvested with trypsin, washed in ice-cold PBS, and the pellets were resuspended and fixed with 70% ethanol at 4°C overnight. Cells were washed with PBS, resuspended in 1 ml of 2 N HCl supplemented with 0.5% Triton X-100 and incubated for 30 min at room temperature. After neutralizing the samples with 1 ml of 100  $\mu$ M borate buffer (Na<sub>2</sub>B<sub>4</sub>O<sub>7</sub>, pH 8.5), the cells were washed with PBS followed by PBS containing 0.5% Tween 20 and 1% BSA. The cells were resuspended in 100  $\mu$ l of PBS containing 5  $\mu$ l of FITC-conjugated anti-BrdU antibody (Invitrogen), 0.5% Tween 20, and 1% BSA, and incubated for overnight at 4°C. Cells were then washed with PBS containing 0.5% Tween 20, and 1% BSA and resuspended in 0.5 ml of PBS containing 5  $\mu$ g/ml propidium iodide, 0.5% Tween 20, and 1% BSA, and incubated for 30 min at room temperature in the dark. Before analysis, each sample was filtered through a 35- $\mu$ m nylon mesh. The percentage of cells in each phase of the cell cycle and the percentage of cells incorporating BrdU was assessed by dual-color flow cytometric analyses using a FACScan (BD Biosciences, San Jose, CA) and analyzed using FlowJo software (Tree Star, Inc., Ashland, OR). A minimum of 5  $\times$  10<sup>4</sup> events were collected for each sample.

**Immunocytochemistry.** U2OS, U2OS cells stably expressing EGFP-NS, MCF-7, and MDA-MB-231 cells were grown on coverslips in 24-well plates in Dulbecco's modified Eagle's medium complete medium for 12 h, then treated with CPEC or vehicle control at the concentrations and times indicated. The cells were fixed in 4% phosphate-buffered saline (PBS)-paraformaldehyde solution for 20 min followed by permeabilization with 0.1% Triton X-100 in PBS for 15 min. After blocking with 5% bovine serum albumin in PBS for 30 min, coverslips were inverted onto a 30  $\mu$ l of drop of the primary antibodies diluted in PBS containing 0.1% Triton X-100 and 5% BSA (1:50 for nucleolin, Mdm2, and nucleostemin). After 1-h incubation with antibody and subsequent washing with PBS three times (10 min each), coverslips were inverted again onto 30  $\mu$ l of drops of the secondary antibodies diluted in PBS containing 5% BSA and 0.1% Triton X-100 [rhodamine-conjugated secondary antibody (1:300); fluorescein isothiocyanate-conjugated secondary antibody (1:50)]. After washing, coverslips were stained with 4',6-diamino-2-phenylindol (300 nM) for 5 min, and washed with PBS three times (5 min each), mounted in Gel Mount Aqueous Mounting Medium (Sigma-Aldrich) and examined using fluorescent microscopy.

**Western Blotting.** Immunoblot analysis was performed as described previously (Huang et al., 2009).

**Apoptosis Assay.** Quantitation of apoptosis by annexin V/propidium iodide (PI) staining was performed as described previously (Huang et al., 2002). In brief, MCF-7 cells were transfected with p53 SMARTpool siRNA (100 nM) and control siRNA (100 nM) by Amaxa Electroporation (Amaxa Biosystems, Gaithersburg, MD). Twenty-four hours after electroporation, cells were exposed to CPEC or DMSO vehicle control for 5 days. Apoptotic cell death was determined using the BD ApoAlert Annexin V-FITC Apoptosis Kit (BD Biosciences) according to the manufacturer's instructions, and cells were analyzed using flow cytometry.

**Oil Red Staining.** Cells grown in 12-well plates were washed with PBS, fixed with paraformaldehyde, washed three times with PBS, and stained with Oil Red O (Sigma-Aldrich) for 10 min according to published procedures (Rydén et al., 2003). Stained cells were

then examined and photographed. The percentage of cells staining positive for lipid droplet accumulation was determined by counting the number of cells containing 10 or more Oil Red O-stained lipid droplets.

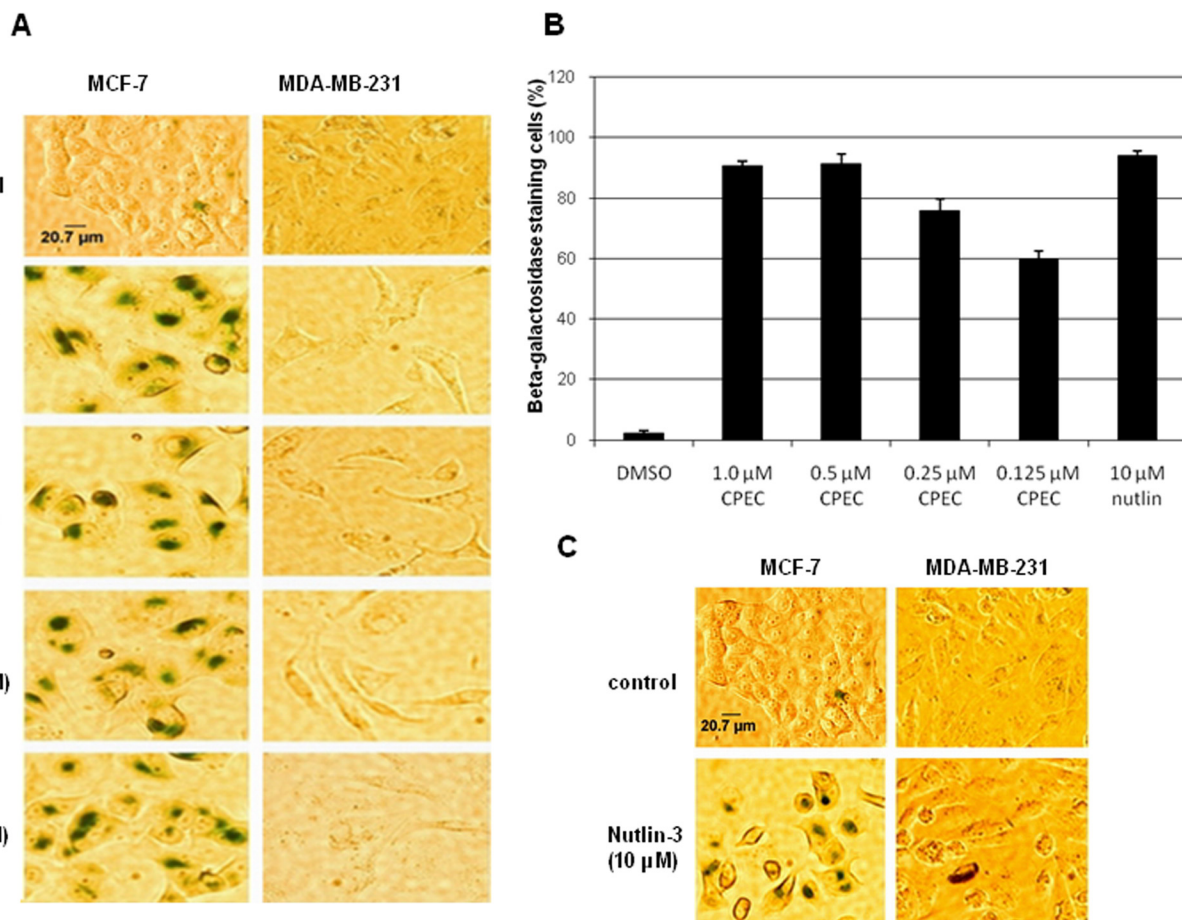
**Silencing of Endogenous p53 by siRNAs.** siRNA SMARTpool targeting p53 and a scrambled control oligonucleotide (5'-AAAGT-CATCGTGACTCGACG-3') were described previously (Huang et al., 2009). The oligonucleotides were electroporated into MCF-7 cells using the Amaxa Electroporation kit V (Amaxa) according to the manufacturer's instruction. Cells/sample ( $2 \times 10^6$ ) were resuspended in 100  $\mu$ l of Nucleofector solution at room temperature and electroporated into MCF-7 cells using 100 or 200 pmol SMARTpool p53-siRNA, scrambled control-siRNA (Program P-20 for MCF-7 on the Amaxa Nucleofector Device). Twenty-four hours after electroporation of p53-siRNA, MCF-7 cells were exposed to 1  $\mu$ M CPEC or vehicle control. Cells were either processed after 8 h of CPEC exposure for Western blot analysis of p53, Mdm2, or p21 or were left for analysis of senescence after 5 days of CPEC exposure.

**Statistical Analysis.** Results of SA- $\beta$ -gal were expressed compared statistically between the control and CPEC-treated results with the paired Student's *t* test with mean  $\pm$  S.D. A *p* value of  $<0.05$  was considered statistically significant.

## Results

### Differential Effects of CPEC Treatment on MCF-7 and MDA-MB-231 Cells. The effect of CPEC on growth of

MCF-7/MDA-MB-231 cells was monitored over a 5-day interval. CPEC treatment inhibited growth in both cell types. After 5 days, MCF-7 cells displayed striking morphological changes consistent with cellular senescence, including cell enlargement, flattening, and a decreased nucleo-cytoplasmic ratio, whereas vehicle-treated control cells remained unchanged. MDA-MB-231 cells harboring mutant p53 exhibited similar morphological changes after CPEC treatment (Supplemental Fig. 1). We then asked whether the growth arrest phenotype was caused by an induction of senescence, quiescence, or terminal differentiation. As shown in Fig. 1A, CPEC treatment of MCF-7 cells significantly increased the percentage of SA- $\beta$ -gal-positive cells ( $P < 0.001$ ) in a concentration-dependent manner after 5 days (Fig. 1, A and B). Nutlin-3a, an inhibitor of Mdm2, was used as a positive control (Fig. 1C). In contrast, no increase in SA- $\beta$ -gal positivity over background was observed in MDA-MB-231 cells treated with either CPEC or Nutlin-3a (Fig. 1, A and C). Next, we investigated whether the growth arrest found in MDA-MB-231 cells was attributable to either quiescence or differentiation. CPEC-treated MDA-MB-231 cells were stained with oil-red to assay for lipid droplets as a reflection of differentiation of breast cancer cells (Adan et al., 2003). As shown in Fig. 2A, more than 80% of MDA-MB-231 cells exhibited a marked



**Fig. 1.** Effect of CPEC on SA- $\beta$ -gal induction in MCF-7 and MDA-MB-231 cells. A and C, MCF-7 or MDA-MB-231 cells were seeded in 12-well plates at a density of  $10^5$  cells/well and continuously exposed to CPEC, 10  $\mu$ M nutlin-3a, or vehicle control for 5 days. Control cells were grown for 3 days and then passaged. Cells were washed, fixed, and stained at pH 6.0 as described under *Materials and Methods*. A, comparison of CPEC effects on induction of SA- $\beta$ -gal in MCF-7 and MDA-MB-231 cells. Scale bar represents 20.7  $\mu$ m for control MCF-7 cells. B, bar graph showing the percentages of  $\beta$ -galactosidase-positive cells (mean  $\pm$  S.D.) in five randomly selected fields of MCF-7 cells containing at least 50 cells per field. C, comparison of effects of Nutlin-3a on the two cell lines. Scale bar represents 20.7  $\mu$ m for control MCF-7 cells.

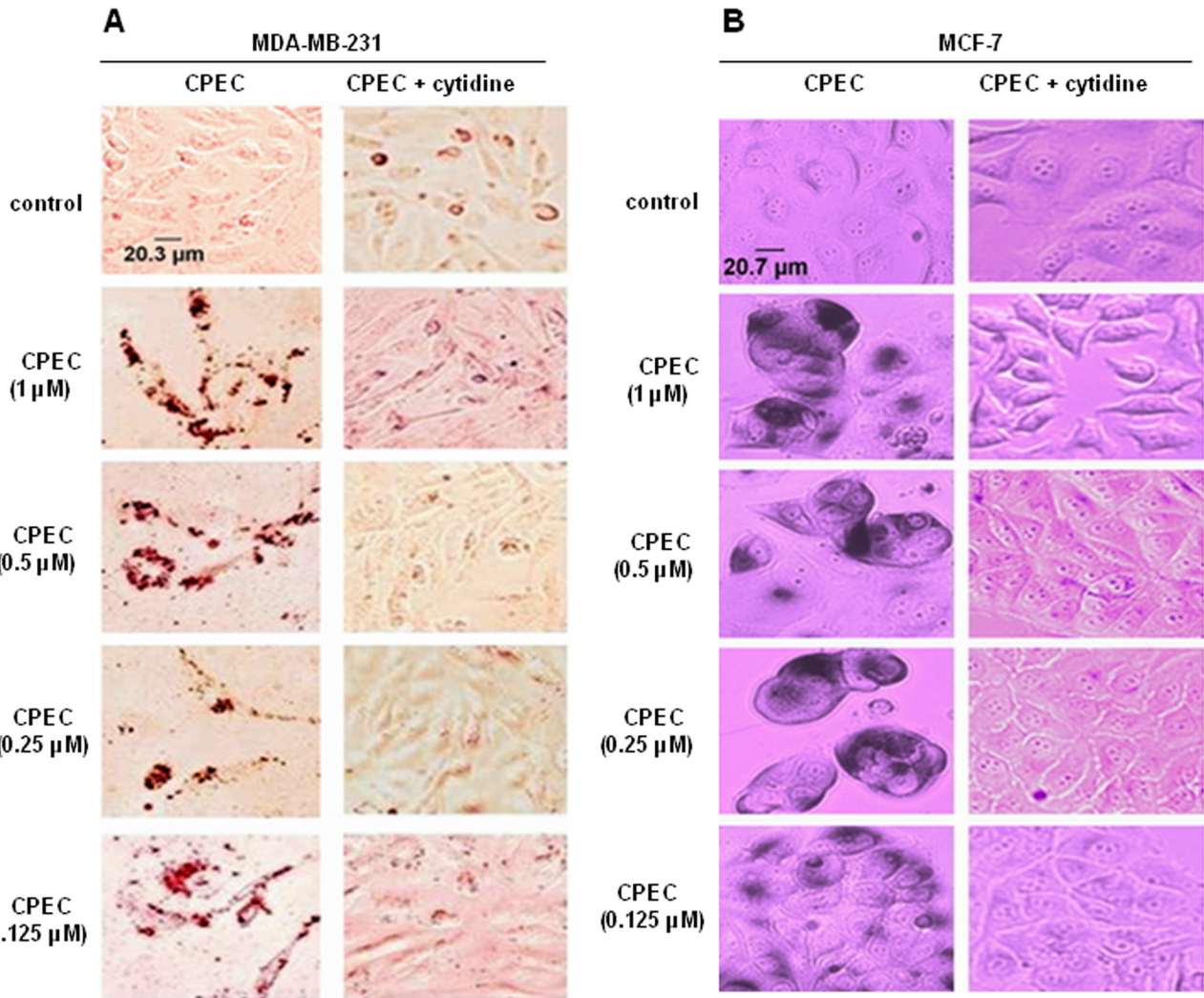


increase in the presence of lipid droplets after 5 days of CPEC treatments, whereas MCF-7 cells did not stain with oil-red (Fig. 1). The addition of cytidine, shown previously to replenish intracellular CTP pools through its phosphorylation to CMP via uridine/cytidine kinase (Huang et al., 2004), almost completely prevented both the differentiation of MDA-MB-231 cells (Fig. 2A) and the senescence of MCF-7 cells (Fig. 2B) because of CPEC treatment. Together, these results indicate that the induction of senescence or differentiation in MCF-7 and MDA-MB-231 cells, respectively, is dependent on CTP depletion.

**CPEC Treatment Differentially Affects Cell Cycle Progression in MCF-7 and MDA-MB-231 Cells.** To further analyze the mechanisms underlying growth arrest induced by CTP depletion, the effects of CPEC on tumor cell proliferation and cell cycle progression were examined using BrdU incorporation and flow cytometric analysis. Treatment with CPEC for 48 h initially increased the number of S-phase cells by approximately 2-fold for both MCF-7 (35.4–44.2.0 versus 24.6% for the control sample) (Supplemental Fig. 2A) and MDA-MB-231 cells (40.7–49.5 versus 18.4% for the con-

trol sample) (Supplemental Fig. 3A), decreasing the extent of BrdU incorporation into DNA (Supplemental Figs. 2B and 3B). Despite the induction of an increase in the S-phase cell population in both cell lines at 48 h, 5 days of CPEC treatment resulted in cell cycle arrest in both the S and G<sub>2</sub> phases in MCF-7 cells (Fig. 3, A and B) but only in G<sub>1</sub> in MDA-MB-231 cells (Fig. 4, A and B). Nutlin-3a induced both a G<sub>1</sub> and a G<sub>2</sub> arrest in MCF-7 cells (Fig. 3, A and B). In both cell lines, a sustained reduction in BrdU incorporation (from 37.5 to 12.8–23.7% in MCF-7 cells and from 20.2 to 2.9–11.1% in MDA-MB-231 cells) was observed 5 days after CPEC treatment (Figs. 3B and 4B), correlating with the lack of proliferation at this time point.

**Effect of CPEC on Activation of p53, p21, and  $\gamma$ -H2AX.** It has been well established that downstream targets of p53 are able to trigger replicative/premature senescence (Collado et al., 2007). p21 (Waf1/Cip1) is a potent cyclin-dependent kinase inhibitor that is transcriptionally regulated by p53 and seems to be central to both p53-dependent (Collado et al., 2007) and p53-independent (Fang et al., 1999) senescence. In MCF-7 cells, CPEC treatment resulted



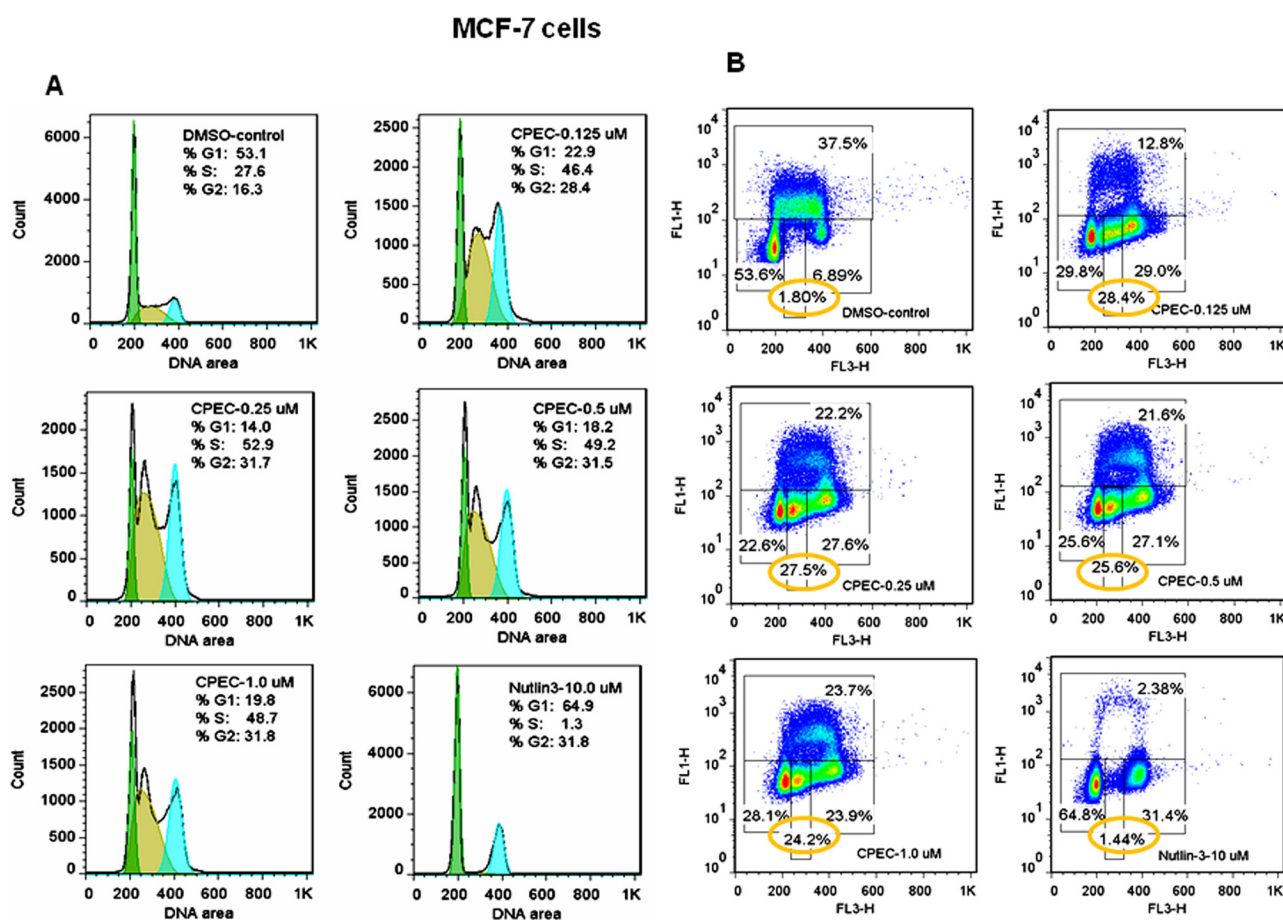
**Fig. 2.** Effect of cytidine on CPEC-induced senescence and differentiation. MDA-MB-231 (A) and MCF-7 (B) cells seeded in 12-well plates at a density of  $10^5$  cells/well were preincubated with 100  $\mu$ M cytidine for 30 min and exposed continuously to increasing concentrations of CPEC or vehicle control for 5 days in the absence or presence of cytidine. Control cells were allowed to grow for three days and passaged. Cells were washed, fixed, and stained for Oil-Red O (MDA-MB-231) or SA- $\beta$ -gal (MCF-7) as described under *Materials and Methods* and photographed at 20 $\times$  magnification. Scale bar represents 20.3  $\mu$ m for the MDA-MB-231 control cells and 20.7  $\mu$ m for the MCF-7 cells.

in a sustained increase in both p53 and p21 expression. In addition, the expression of Mdm2, another p53 target, was induced by CPEC treatment in MCF-7 cells (Fig. 5A), whereas  $\beta$ -actin expression remained constant. A moderate increase in the Mdm2 protein level was also detected in p53-mutant MDA-MB-231 cells, suggesting that there may be a p53-independent regulation of Mdm2 (Fig. 5B). The levels of p21 were unchanged in these cells (Fig. 5B). As expected, a similar pattern of p53 and Mdm2 accumulation was observed in nutlin3-treated MCF-7 cells (Supplemental Fig. 4). Phosphorylated  $\gamma$ -H2AX, an indicator of DNA damage response, increased dramatically in MDA-MB-231 cells after 24 h of CPEC exposure but did not change significantly compared with control in MCF-7 cells (Fig. 5, A and B).

**Role of p53 in CPEC-Induced Senescence.** To further confirm the role of p53 in mediating CPEC-induced senescence, we used a SMARTpool p53 siRNA that specifically depletes p53 protein expression and asked whether the knockdown of p53 would prevent CPEC-induced senescence. As shown in Fig. 6A, transfection of 200 pmol SMARTpool siRNA of p53 reduced the level of endogenous p53 protein by more than 80%. Depletion of p53 using the siRNA approach completely prevented CPEC-induced elevation of Mdm2 (Fig. 6A) and significantly reduced the degree of senescence ( $P < 0.001$ ) (Fig. 6, B and C) and increased the percentage of Annexin V and PI double-positive staining cells from 21.7 to

36.0% (Fig. 6D). An increased percentage of Annexin V and PI double-positive staining cells (73.5–77.9% of cells) suggestive of apoptosis was also observed in p53-null HL60 cells after exposure to CPEC at concentrations as low as 0.125  $\mu$ M for 32 h (Fig. 6E). To further determine whether cell death was due to apoptosis, DNA was isolated from control or CPEC-treated cells and separated by agarose gel electrophoresis. As shown in Supplemental Fig. 5A, DNA fragmentation characteristic of apoptosis occurred in CPEC-treated p53-deficient HL60 cells. Likewise, enhanced DNA ladder formation was observed in CPEC-treated p53-depleted MCF-7 cells compared with control MCF-7 cells (Supplemental Fig. 5B). These results indicate that p53 is central to CPEC-induced senescence of MCF-7 cells and that depletion of CTP by CPEC induces apoptotic cell death in the absence of p53.

**Effect of CPEC on the Localization of Nucleolar Proteins.** Nucleolar disruption caused by inducers of nucleolar stress, including inhibition of RNA polymerase I (Pol I), is implicated in the stabilization of p53 (Yuan et al., 2005). We have found previously that depletion of the cellular GTP pool by AVN-944 resulted in the inhibition of preribosomal RNA synthesis, disruption of nucleolar integrity, and stabilization of p53 (Huang et al., 2008). We therefore asked whether CTP depletion induced similar disruption of nucleolar integrity that might serve as an upstream signal resulting in p53



**Fig. 3.** Differential effects of CPEC on cell cycle progression and BrdU incorporation in MCF-7 cells. Analyses of cell cycle progression (A) and BrdU incorporation (B) in MCF-7 were performed as described under *Materials and Methods* after 5 days of culture in the absence or presence of CPEC at the concentrations indicated. The percentage of cells in each phases of the cell cycle is indicated for each sample in the histogram. Yellow circles indicate S phase-arrested cells lacking active BrdU incorporation. Data shown are representative of two experiments.

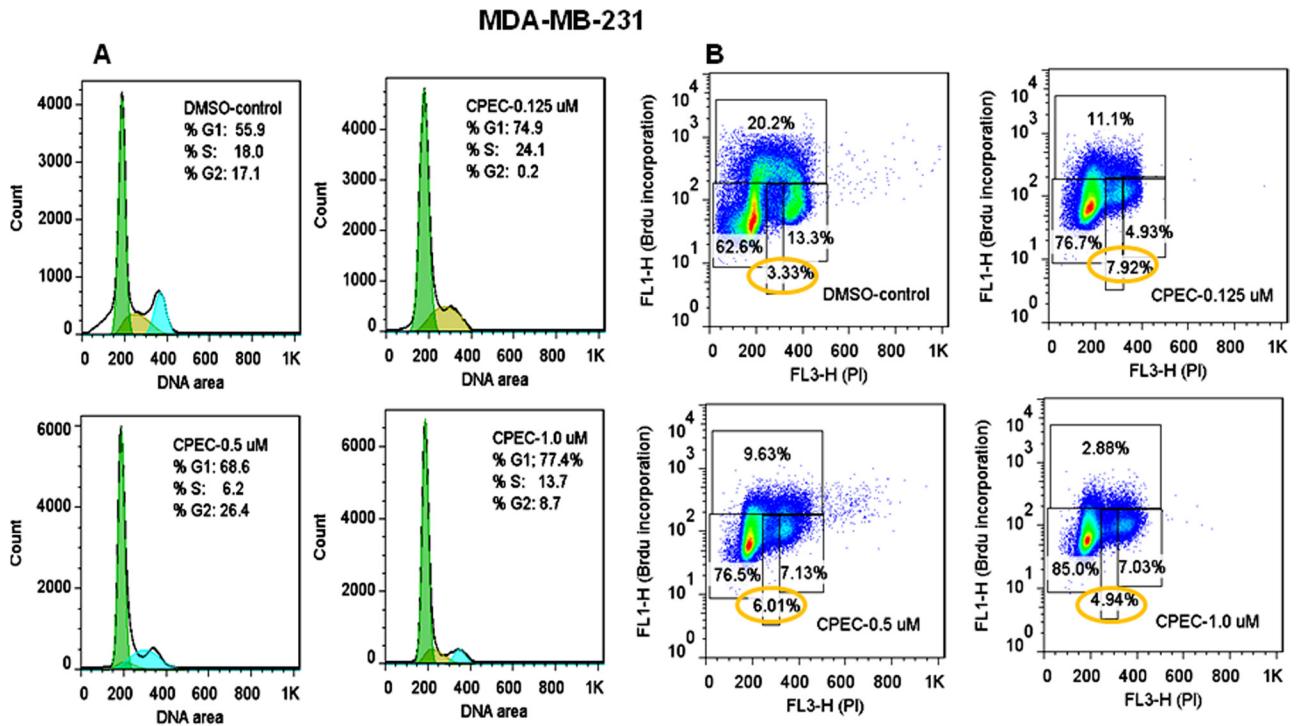


stabilization. Indeed, CTP depletion caused the rapid translocation of the nucleolar shuttle protein NS from the nucleolus to the nucleus and a moderate increase in Mdm2 protein expression in MCF-7 (Fig. 7A) and MDA-MB-231 cells (Fig. 7B). Because of their superior nucleolar morphology, we also used U2OS cells to evaluate the localization of nucleolar proteins. CPEC induced nucleoplasmic translocation of NS and nucleolin in this cell line as well (Fig. 7C; Supplemental Fig. 6). Previous data have demonstrated that inhibition of Pol I by actinomycin D results in translocation of nucleolar PAF53 and TIF-1A, two components of the Pol I complex, to the nucleolar caps (Huang et al., 2008), a phenomenon reflecting "nucleolar stress." CTP depletion also induces similar translocation of TIF-1A and PAF53 from the nucleolus to

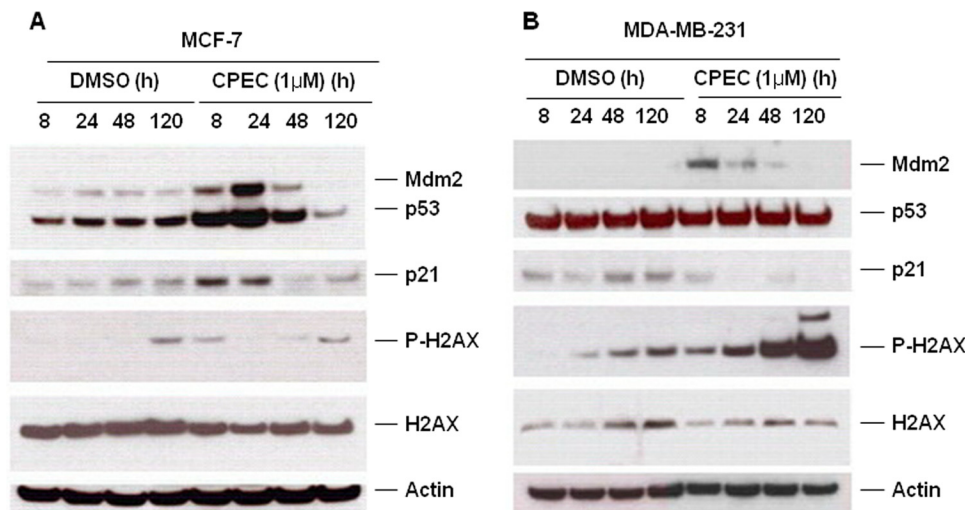
the nucleolar caps (Supplemental Fig. 6), further supporting its inhibitory effect on Pol I activity through CTP depletion.

## Discussion

p53 is a central mediator of senescence in response to a variety of stress stimuli, including DNA damage, oncogene overexpression, telomere dysfunction, and oxidative damage (Collado et al., 2007). Our finding that CPEC induces senescence in MCF-7 cells expressing wild-type p53 in conjunction with sustained increases in p53 and p53-dependent Mdm2 and p21 expression whereas senescence is prevented by p53 depletion strongly implicates p53 accumulation as a prerequisite for the senescent phenotype.



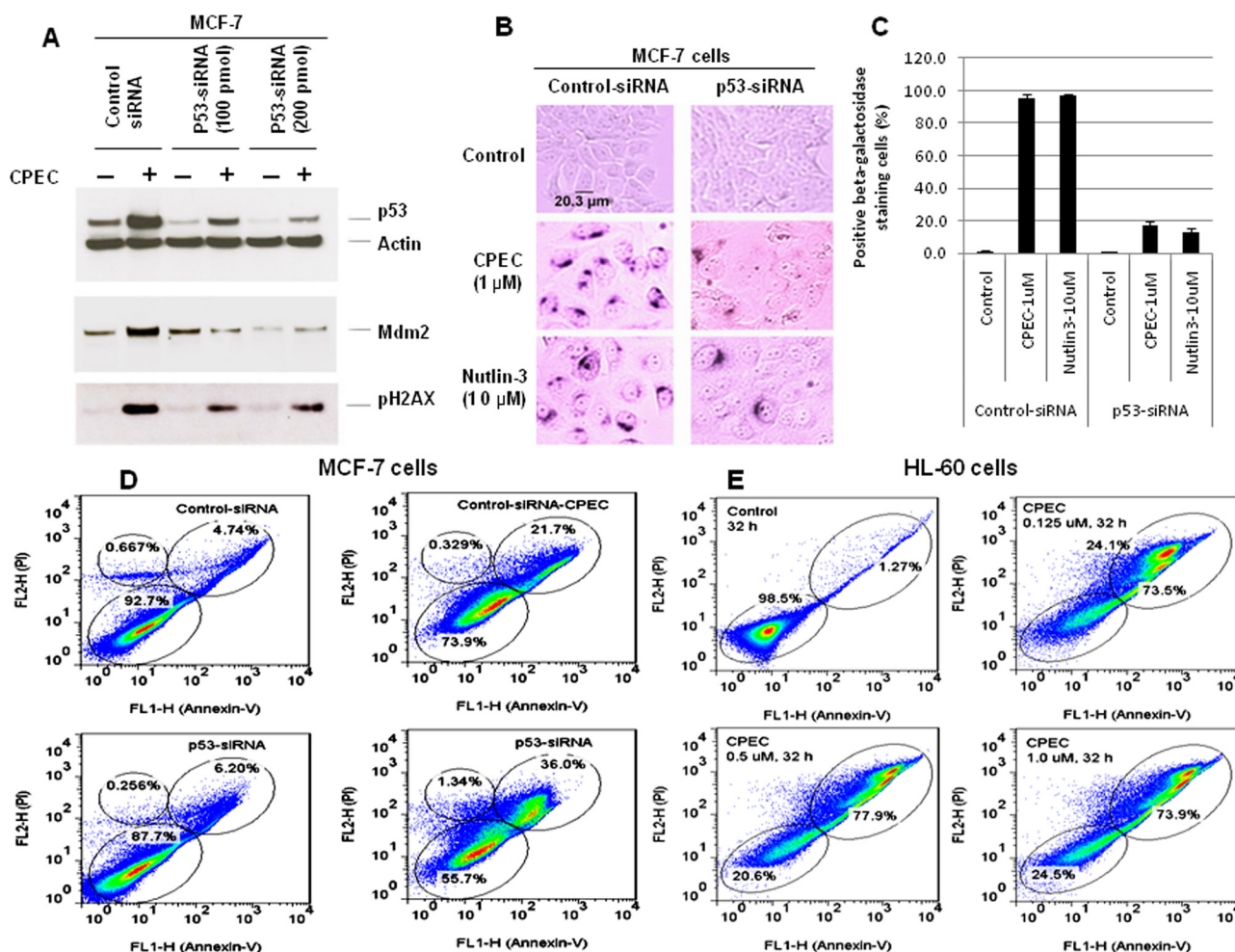
**Fig. 4.** Differential effects of CPEC on cell cycle progression and BrdU incorporation in MDA-MB-231 cells. Analyses of cell cycle progression (A) and BrdU incorporation (B) in MDA-MB-231 cells were performed as described under *Materials and Methods* after 5 days of culture in the absence or presence of CPEC at the concentrations indicated. The percentage of cells in each phase of the cell cycle is indicated for each sample in the histogram. Data are representative of two experiments.



**Fig. 5.** Differential effects of CPEC on the p53/p21 pathway. Immunoblot analysis of p53, nucleostemin, Mdm2, p21,  $\gamma$ -H2A, and p- $\gamma$ H2AX in MCF-7 (A) and MDA-MB-231 cells (B) in the absence or presence of 1  $\mu$ M CPEC.  $\beta$ -Actin serves as a loading control.

Mdm2 maintains p53 at low levels by actively controlling its nuclear export (Freedman and Levine, 1998), transactivation (Momand et al., 1992), ubiquitination and proteasome-dependent degradation (Honda et al., 1997), and translation (Ofir-Rosenfeld et al., 2008). The Mdm2 gene is also a transcriptional target of p53, and an increase in p53 expression upon stress leads to activation of Mdm2 transcription (Barak et al., 1993). This Mdm2-p53 autoregulatory loop normally maintains p53 at low levels in nonstressed cells. Our evidence that depletion of cellular CTP leads to stabilization of wild-type p53 in the presence of excessive Mdm2 suggests that the inhibitory regulation of p53 by Mdm2 might be disrupted by CTP depletion. Several Mdm2-binding proteins have been shown to interfere with this autoregulatory loop through direct binding to the central acidic domain of Mdm2, thus inhibiting its E3 ubiquitin ligase activity and increasing p53 levels. These Mdm2-binding proteins include ARF (Bothner et al., 2001), the ribosomal large-subunit proteins L5, L11, and L23 (Lindström et al., 2007), the nucleolar protein nucleostemin (Dai et al., 2008; Meng et al., 2008), and the ribosomal small subunit protein 7 (Zhu et al., 2009). Of

interest, most of those Mdm2 binding proteins are localized predominantly in the nucleolus under physiological conditions but are released from nucleolus into nucleoplasm in response to various stressors, including agents such as low-dose actinomycin D that inhibit Pol I. We have shown previously that depletion of GTP induced by inhibition of its de novo synthetic pathway likewise induces a nucleolar stress response (Huang et al., 2008). In the present study, CPEC-mediated CTP depletion also results in nucleolar stress, causing nucleoplasmic translocation of nucleostemin and other nucleolar proteins, including nucleolin and nucleophosmin 1 and a simultaneous translocation of TIF-IA and PAF53, two components of the Pol I complex, to the nucleolar caps. Segregation of Pol I components into these fibrillar caps is a characteristic feature of inhibition of RNA polymerase I-mediated transcription (Shav-Tal et al., 2005). It is reasonable to conclude that the release of nucleolar Mdm2-binding proteins triggered by CTP depletion is, at least in part, responsible for the increased stability of p53. The lack of ARF expression in MCF-7 cells rules out a role for ARF in the observed p53 activation (Della Ragione et al., 1995). How-



**Fig. 6.** Effect of p53 siRNA on CPEC-induced senescence in MCF-7 cells. MCF-7 cells were transfected with 100 nM/200 nM p53-siRNA or control-siRNA by Amaxa electroporation. A, 24 hours after transfection, the cells were treated with 1  $\mu$ M CPEC for 20 h, and 40  $\mu$ g of cell lysate was subjected to Western blot analysis using antibodies against p53, Mdm2, p- $\gamma$ H2AX, and  $\gamma$ -H2A.  $\beta$ -Actin served as loading control. B, 5 days after CPEC treatment, the cells were stained for  $\beta$ -gal activity. Cells were photographed at 20 $\times$  magnification. Scale bar represents 20.3  $\mu$ m for control MCF-7 cells. C, bar graph showing the percentage of  $\beta$ -galactosidase-positive cells (mean  $\pm$  S.D.) counted in five randomly selected fields containing at least 50 cells per field. D, 24 h after Amaxa electroporation of 100 nM p53-siRNA or control-siRNA, the cells were treated with 1  $\mu$ M CPEC for 5 days and then stained with Annexin-V and PI followed by flow cytometric analysis. E, HL60 cells were treated with CPEC at doses indicated for 32 h and then stained with Annexin-V and PI followed by flow cytometric analysis.



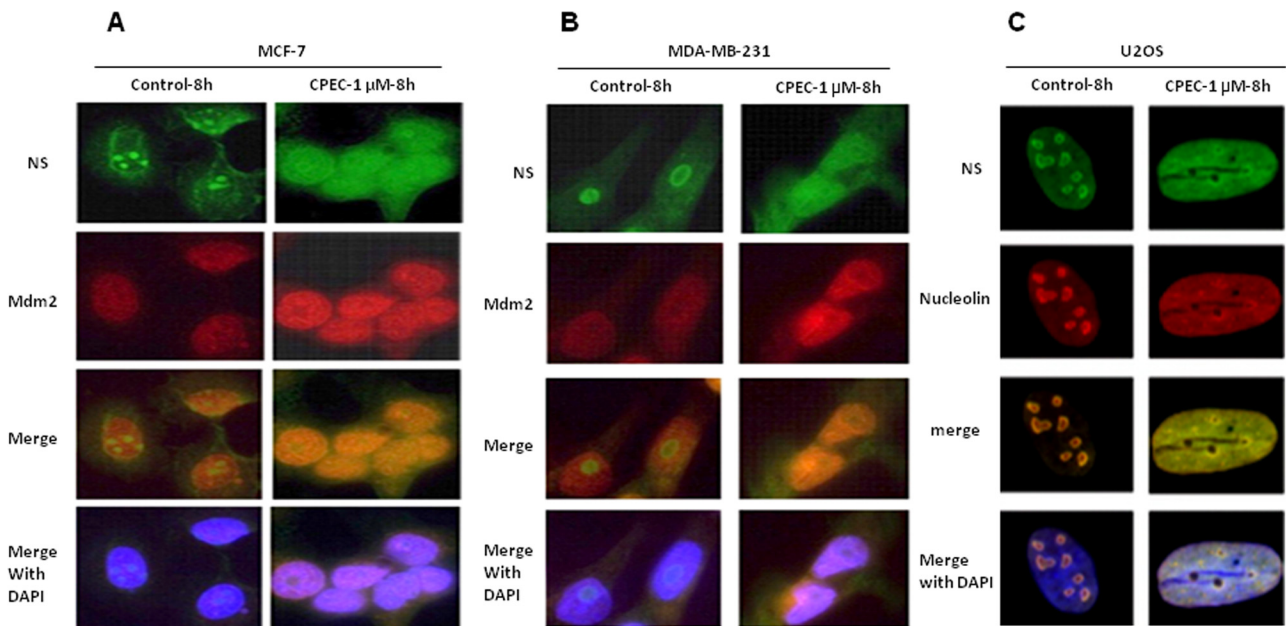
ever, the presumed release of other Mdm2-binding proteins, including a number of small ribosomal subunit proteins, is very likely to have played a role.

Cells entering senescence undergo irreversible cell cycle arrest, most often in the G<sub>1</sub>/S and/or G<sub>2</sub>/M phases of the cell cycle (Sugrue et al., 1997). Inducible expression of wild-type p53 in EJ bladder carcinoma cells, which have lost functional p53, was shown to trigger the rapid onset of G<sub>1</sub> and G<sub>2</sub>/M senescent growth arrest associated with p21 up-regulation (Sugrue et al., 1997). Supporting a role for p53 in mediating G<sub>2</sub> cell cycle arrest, nutlin-3a and CPEC both induce p53 accumulation and G<sub>2</sub> arrest in MCF-7 cells that undergo senescence (Fig. 3, A and B, and supplemental Fig. 4), but not in MDA-MB-231 cells expressing mutated p53 (Fig. 4, A and B). p16<sup>INK4a</sup> can also initiate senescence through inhibiting the cyclin-dependent kinases cdk4 and cdk6 (Uhrbom et al., 1997). However, MCF-7 and MDA-MB-231 cells both have homozygous deletions of p16<sup>INK4a</sup> (Della Ragione et al., 1995), ruling out involvement of this protein as an effector of CPEC-induced senescence or differentiation. The increase in p-H2AX caused by CPEC treatment in MDA-MB-231 cells reflects a DNA damage response, most likely due to concomitant depletion of the cytidine deoxyribonucleotide pool. Under similar conditions, other deoxyribonucleotides are misincorporated into DNA, resulting in a similar DNA damage response (Schimmel et al., 2007).

The observation that depletion of endogenous p53 switched the cell fate of MCF-7 cells from a senescent to an apoptotic fate indicates that activation of the p53 pathway plays an essential role in inducing the senescence phenotype. It has been well documented that p53 mutations such as R175H, R273H, and V135A mediate resistance to drug-induced apoptosis through suppression of procaspase-3 and caspase-3 (Lotem and Sachs, 1995; Tsang et al., 2005; Wong et al., 2007). These data raise the possibility that the expression of mutated p53 primes MDA-MB-231 cells for differentiation in

response to nucleolar stress induced by CTP depletion, whereas p53 deficiency primes cells for apoptosis.

Although CPEC alone has established antitumor effects in a variety of cancer model systems, both in vitro and in vivo (Schimmel et al., 2007; Van Bree et al., 2009), severe cardiovascular toxicity was observed in 5 of 26 patients with solid tumors treated in a phase I clinical trial. However, this toxicity occurred only at the highest doses administered with corresponding steady-state plasma CPEC concentrations varying between 1.8 and 2.75  $\mu$ M (Politi et al., 1995). Our data, in conjunction with those of others (Huang et al., 2004; Schimmel et al., 2007), indicate that CPEC induces senescence, differentiation, or apoptosis at nanomolar concentrations in tumor cell lines in vitro. In theory, the effects of CPEC against tumors could be mitigated by cytidine present in human plasma. However, the physiological concentrations of uridine (5  $\mu$ M) or cytidine (0.5  $\mu$ M) in human plasma are not sufficient to prevent cell cycle arrest induced by nanomolar concentration of CPEC (50 nM) (Huang et al., 2004). In contrast to transformed cells, primary mouse embryonic fibroblast cells expressing wild-type p53 are relatively resistant to CPEC treatment at concentrations up to 1  $\mu$ M (data not shown). It is quite possible that CPEC is selectively toxic to transformed cells due to increased uptake as the result of the increase in nucleoside transporters identified in tumor cells (Rauchwerger et al., 2000). Whether the intracellular metabolism of CPEC to its triphosphate form is also enhanced in some tumor cells through an increase in uridine-cytidine kinase activity (Kang et al., 1989) or a decrease in nucleotidase activity has not been investigated. If CPEC is to be reconsidered as an antineoplastic agent, especially given evidence of activity in p53-deficient cells, it will be important to investigate its relative uptake and metabolism in both tumor and normal tissue. A clinical resurgence of this drug would also require in vitro and in vivo studies to determine



**Fig. 7.** Effect of CPEC on the localization of the nucleolar proteins nucleostemin, Mdm2, and nucleolin in MCF-7, MDA-MB-231 cells, and U2OS cells. MCF (A) and MDA-MB-231 (B) cells were treated with 1  $\mu$ M CPEC for 8 h. Cells were then fixed, permeabilized, and stained for NS and Mdm2 as described under *Materials and Methods* and photographed using fluorescent microscopy at 100 $\times$  magnification. C, U2OS cells were treated with 1  $\mu$ M CPEC for 8 h, fixed, permeabilized, stained for NS and nucleolin, and photographed using fluorescent microscopy at 100 $\times$  magnification.



the optimal plasma concentration of CPEC that results in selective toxicity for tumor cells.

#### Authorship Contributions

*Participated in research design:* Huang and Mitchell.  
*Conducted experiments:* Huang, Whang, and Lewicki.  
*Performed data analysis:* Huang, Whang, and Lewicki.

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**Address correspondence to:** Dr. Beverly S. Mitchell, Stanford Cancer Center, LLSC Research Building (SIM1), 265 Campus Drive, Room 2167, Stanford, CA 94305-5458. E-mail: bmmitchell@stanford.edu