p27^{kip1} Overexpression Promotes Paclitaxel-Induced Apoptosis in pRb-Defective SaOs-2 Cells

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Abstract p27^{kip1} is a cyclin-dependent kinase (CDK) inhibitor, which controls several cellular processes in strict collaboration with pRb. We evaluated the role of p27^{kip1} in paclitaxel-induced apoptosis in the pRb-defective SaOs-2 cells. Following 48 h of exposure of SaOs-2 cells to 100 nM paclitaxel, we observed an increase in p27^{kip1} expression caused by the decrease of the ubiquitin-proteasome activity. Such increase was not observed in SaOs-2 cells treated with the caspase inhibitors Z-VAD-FMK, suggesting that p27^{kip1} enhancement at 48 h is strictly related to apoptosis. Finally, we demonstrated that SaOs-2 cells transiently overexpressing the p27^{kip1} protein are more susceptible to paclitaxel-induced apoptosis than SaOs-2 cells transiently transfected with the empty vector. Indeed, after 48 h of paclitaxel treatment, 41.8% of SaOs-2 cells transiently transfected with a pcDNA3-p27^{kip1} construct were Annexin V-positive compared to 30.6% of SaOs-2 cells transfected with the empty vector (P < 0.05). In conclusion, we demonstrated that transfection of the pRb-defective SaOs-2 cells with the p27^{kip1} gene via plasmid increases their susceptibility to paclitaxel-induced apoptosis. The promoting effect of p27^{kip1} overexpression on apoptosis makes p27^{kip1} and proteasomal inhibitors interesting tools for therapy in patients with pRb-defective cancers. J. Cell. Biochem. 98: 1645–1652, 2006. © 2006 Wiley-Liss, Inc.

Key words: p27^{kip1}; apoptosis; paclitaxel; Osteosarcoma

Apoptosis is an active process of programmed cell death that is preserved throughout evolution [Vaux and Korsmeyer, 1999]. The cell cycle

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is the highly conserved innate mechanism by which eukaryotic cells replicate themselves. In metazoans the processes of cell loss and cell gain are homeostatically balanced in order to maintain the normal development and architecture of tissues. Accumulating evidence shows that the manipulation of the cell cycle may either prevent or induce an apoptotic response [Pucci et al., 2000]. This ability has been recognized for several cell cycle proteins such as p53 and the retinoblastoma family members, pRb/ p105, p107, and pRb2/p130 [Morgenbesser et al., 1994; Paggi et al., 1996; Pucci et al., 2002; Tonini et al., 2004]. Data concerning the role of cyclin-dependent kinases (CDKs), cyclins, and CDK inhibitors (CKIs) also support a direct coupling between the proliferating machinery and the apoptotic process [Kasten and Giordano, 1998].

Abbreviations used: PTX, paclitaxel.

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Growing evidence suggests that apoptosis frequently occurs in cells in the G₁ phase of the cell cycle [Meikrantz and Schlegel, 1995], thus CKIs expressed during late G₁ like p27^{kip1} are logical targets for the apoptotic process. The cyclin-dependent kinase inhibitor p27^{kip1} is an important negative regulator in cell cycle identified as a 27 kDa protein that binds to and inhibits the activity of the cyclin E-CDK2 complexes in cells treated with transforming growth factor TGF- β and is regulated by growth inhibitory cytokines and by contact inhibition [Koff et al., 1993; Hengst et al., 1994; Slingerl et al., 1994; Polyak et al., 1994a,b].

In several cellular systems $p27^{kip1}$ is regulated primarily post-transcriptionally by the ubiquitin-proteasome pathway and phosphorylation of $p27^{kip1}$ at Thr187 by cyclin E-CDK2 is required to initiate the ubiquitination-proteasomal degradation [Sheaff et al., 1997; Montagnoli et al., 1999].

Although $p27^{kip1}$ has been shown to prevent drug-induced apoptosis in carcinoma cells [St Croix et al., 1996], other studies have pointed toward its pro-apoptotic role. Recombinant adenovirus overexpressing $p27^{kip1}$ triggers apoptosis in several cancer cell lines [Craig et al., 1997; Katayose et al., 1997] and cell death induction by $p27^{kip1}$ overexpression is mediated by the presence of pRb in lung cancer cell lines [Naruse et al., 2000]. Moreover, it has been recently shown that the adenoviral transduction of a proteolysis-resistant form of $p27^{kip1}$ is able to increase apoptosis in breast cancer cells [Zhang et al., 2005].

Paclitaxel is an antimitotic agent [Kumar, 1981] that inhibits the formation of mitotic spindle [Schiff and Horwitz, 1980], thus causing G_2/M arrest as well as cellular toxicity via apoptosis [Roberts et al., 1990; Bhalla et al., 1993]. Its cytotoxic effects have been shown in several tumor cell lines [Wani et al., 1971; Douros and Suffness, 1978; Pucci et al., 1999], in a variety of murine tumors, and human xenografts [Riondel et al., 1986, 1988], as well as in advanced human carcinomas resistant to traditional chemotherapy [Rowinsky and Donehower, 1995]. However, the underlying mechanism by which paclitaxel induces cell death remains poorly understood. Among apoptotic factors, few of the Bcl-2 family proteins have been shown to be involved in paclitaxel-induced apoptosis [Ferlini et al., 2003]. For example, Bcl-2 is phosphorylated

and consequently inactivated by Raf-1 during paclitaxel apoptosis [Ling et al., 1998]. Overexpression of another anti-apoptotic protein, Bcl-x_L, confers resistance to paclitaxel-induced apoptosis in HL-60 cells. On the other hand, the overexpression of the pro-apoptotic proteins Bax and Bcl-x_S respectively sensitize SW626 and MCF7 cells to paclitaxel treatment [Sumantran et al., 1995; Strobel et al., 1996; Wang et al., 2000]. Several cell cycle regulators also appear to be involved in paclitaxel-induced cell death. Among those, p34cdc2 is activated during paclitaxel-induced apoptosis and is responsible for G_2/M arrest and cell death [Yu et al., 1998]. Overexpression of the CKIs p21^{cip1} or p27^{kip1} produces a differential resistance to paclitaxel in RKO colon carcinoma cells, causing a G₁ but mostly G₂ arrest, which prevents cells from entering M phase and becoming paclitaxelsusceptible [Schmidt et al., 2000]. When transferred from monolayer to three-dimensional culture, A2780 and CAOV3 ovarian cancer cell lines show an increase of resistance to paclitaxel treatment that correlates with the upregulation of p27^{kip1} and multidrug resistance P-glycoprotein expression [Xing et al., 2005].

However, induction of p27^{kip1} conferred resistance to paclitaxel in Rb-positive but not Rb nonfunctional adenocarcinoma cell lines [Schmidt and Fan, 2001].

In the present study, we showed that p27^{kip1} overexpression increases paclitaxel-induced apoptotic rate in the pRb defective SaOs-2 osteosarcoma cell line.

MATERIALS AND METHODS

Cell Culture

SaOs-2 (p53 and pRb-null) human osteosarcoma cell line was purchased from ATCC (Rockville, MD), and grown in monolayer cultures in Dulbecco's modified Eagle's medium D-MEM (GIBCO, Grand Island, NY) supplemented with 10% fetal bovine serum (FBS; HyClone), 2 mM L-glutamin (GIBCO, Invitrogen) and gentamicin 50 μ g/ml (GIBCO, Invitrogen), at 37°C in a humidified atmosphere 5% CO₂ and 95% air.

Treatments

Paclitaxel stock solutions (10 mM) were dissolved in dimethyl sulfoxide (DMSO) and protected from light. Drug dilutions were obtained at the appropriate concentration with medium and were freshly prepared before each experiment. Broad-range caspase inhibitor Z-VAD-FMK (Bachem Bioscience, Inc., Philadelphia, PA) was prepared from 100 mM in DMSO stock solution. Twenty-four hours after plating, cells were treated with 100 nM paclitaxel from 16 to 48 h in the presence 100 μ M of the caspase inhibitor. Z-VAD-FMK was added fresh daily to prevent its degradation.

Degradation Assay

SaOs-2 cells untreated or treated with 100 nM paclitaxel for 16 and 48 h were collected and protein extracts were prepared by preserving ubiquitinating enzymes. Purified histidine-tagged p27^{kip1} (0.06 μ l) was incubated at 37°C for 72 h in 30 μ l of degradation mix containing 50 μ g of extracts, 50 mM Tris-HCl (pH 8), 5 mM MgCl₂, 1 mM DTT, 2 mM ATP, 10 mM creatine phosphate, and 5 μ M ubiquitin [Pagano et al., 1995]. The degradation of p27^{kip1} was analyzed by immunoblotting with an antihuman p27^{kip1} polyclonal antibody (clone C-19, Santa Cruz, CA).

Transfections

SaOs-2 cells were seeded on six-well plates $(1 \times 10^5$ cells/well) and after 24 h pcDNA3 empty vector and pcDNA3-p27^{kip1} [Sgambato et al., 1998] were transiently transfected with LipofectamineTM 2000 and PlusTM Reagent (Invitrogen, Carlsbad, CA), according to the manufacturer's guidelines. Briefly, SaOs-2 cell line was transfected with a ratio $1:2(0.5 \mu g DNA)$ and 1 µl LipofectamineTM 2000) in OptiMEM[®] I (GIBCO) for 16 h. 24 h after transfection SaOs-2 treated with 100 nM paclitaxel (Calbiochem, La Jolla, CA) for 16, 24 and 48 h. SaOs-2 cell line was also transfected with a plasmid vector expressing GFP and the transfection efficiency was assessed to be around 50% by flow cytometry analysis.

Western Blot Analysis

Cells were harvested, washed and lysed in lysis buffer (50 mmol/L Tris-Cl, pH 7.4; 5 mmol/L EDTA; 250 mmol/L NaCl; 50 mmol/L NaF; 0.1% Triton X-100; 0.1 mmol/L Na₃VO₄; 1 mmol/L phenylmethylsulfonyl fluoride, 10 μ g/mL leupeptin). Protein concentrations were measured using the BCA protein assay (Pierce, Rockford, IL) and 50 μ g of protein extracts from each sample were separated on denaturing SDS-PAGE gels and blotted on nitrocellulose

membrane. Immunodetections were performed using anti-p27^{kip1} (clone c-19) antibody, obtained from Santa Cruz Biotecnology (Santa Cruz, CA), and anti-pRb antibody (clone G3-245; BD PharMingen). To check the amount of proteins transferred to nitrocellulose membrane, β -actin was used as control and it was detected by anti- β -actin antibody (clone AC-15; Sigma, Saint Louis, MI). Protein detection was performed using the enhanced chemiluminescence detection reagents (Amersham Corp, Buckinghamshire, UK). Densitometric evaluation of Western blot analysis of p27^{kip1} expression was performed using Molecular Analyst Software (Bio-Rad) and normalized with relative β -actin expression.

Flow Cytometry Analysis

Cells were trypsinized, counted with Coulter Counter (model ZM, Kontron Instruments, Watford, Herts, UK) and washed with PBS. For cell cycle analysis, cells were fixed 1×10^6 cells/ml in ice-cold 70% EtOH while vortexing and stored at 4°C. After a wash in PBS, cells were stained in 10 µg/mL propidium iodide (Sigma, Saint Louis, MI), 250 µg/mL RNAase (Sigma) PBS solution and incubated for 30 min at 37°C in the dark. Apoptosis was assessed using VybrantTM Apoptosis Assay Kit (Molecular Probes) according to the manufacturer's instructions. All flow cytometry acquisitions were performed using a FACSCalibur instrument (Becton Dickinson Bioscences). Cell cycle distribution was evaluated using ModFit software and Annexin V-positive/PI negative cell percentage was quantified using CellQuest software.

RESULTS

Increase of p27^{kip1} Levels in Paclitaxel-Induced Apoptosis of SaOs-2 Cell Line

The role of p27^{kip1} was evaluated in SaOs-2 cells undergoing paclitaxel-induced apoptosis [Pucci et al., 1999]. Cells were grown in medium containing 100 nM paclitaxel, harvested after 16, 24, and 48 h of treatment and analyzed by Western blot analysis for p27^{kip1} levels. As shown in Figure 1, p27^{kip1} expression decreased at 16 h; however, starting from 24 to 48 h following paclitaxel treatment, p27^{kip1} was expressed again. β -actin was used as a loading control (Fig. 1).



Fig. 1. $p27^{kip1}$ expression during paclitaxel treatment of osteosarcoma Rb-null SaOs-2 cell line. Western blot analysis of $p27^{kip1}$ expression in SaOs-2 cell line, treated with 100 nM paclitaxel at indicated times. Untreated control cells were collected following 16 h of treatment. SaOs-2 cells were treated for 48 h with 100 nM paclitaxel together with 100 μ M of the broad-range caspase inhibitor Z-VAD-FMK.

Next step was to test if $p27^{kip1}$ increase at 48 h of paclitaxel treatment was specifically related to apoptosis. In a previous study [Pucci et al., 1999], we demonstrated that the broad-range caspase inhibitor Z-VAD-FMK is able to block the apoptotic process induced by paclitaxel. Thus, we treated SaOs-2 cells with 100 nM paclitaxel in the presence of Z-VAD-FMK for 48 h and the levels of $p27^{kip1}$ were determined by Western blot analysis. As Figure 1 shows, $p27^{kip1}$ levels did not increase following 48 h of paclitaxel treatment in presence of Z-VAD-FMK.

Inhibition of p27^{kip1} Protein Degradation During Paclitaxel-Induced Apoptosis

To verify if the proteasomal activity was involved in modulation of p27^{kip1} protein during paclitaxel treatment, SaOs-2 cells untreated or treated with 100 nM paclitaxel for 16 and 48 h were collected and a degradation assay was performed by adding purified recombinant p27^{kip1} to the protein extracts [Esposito et al., 1997]. This assay was designed for testing a specific proteasome-dependent degradation activity, indeed protein extracts were prepared by preserving ubiquitinating enzymes and the assay was conducted in presence of ubiquitin. Figure 2 shows that recombinant p27^{kip1} protein was degraded only in the extracts obtained following 16 h of treatment with paclitaxel. No degradation was observed in the extracts obtained following 48 h of treatment. As expected, p27^{kip1} protein was not degraded in either of the extracts from untreated cells nor in the reaction mix, which consisted of the buffer plus the recombinant protein.



Fig. 2. Degradation activity of SaOs-2 cells, untreated or treated with 100 nM paclitaxel at the indicated times. Purified histidine-tagged $p27^{kip1}$ was used as substrate of reaction.

p27^{kip1} Overexpression Enhances Paclitaxel-Induced Apoptosis in SaOs-2 Cell Line

Finally, to study the role of p27^{kip1} increase during paclitaxel-induced apoptosis, we examined the effect of a transient overexpression of p27^{kip1} on the cytotoxicity of paclitaxel. SaOs-2 cells were transiently transfected with the constructs pcDNA3 and pcDNA3-p27^{kip1} [Sgambato et al., 1998]. The day after transfection, the cells were treated with 100 nM paclitaxel for 16, 24, and 48 h. At the end of each treatment, the cells were collected for Western blot and flow cytometry analyses. SaOs-2 cells transfected with the empty vector or pcDNA3-p27^{kip1} were collected at 16 h and used as an internal control. Figure 3A shows that p27^{kip1} was overexpressed at each time point studied and β -actin expression was used as the loading control. SaOs-2 trasfection with pcDNA3-p27^{kip1} vector determined an increase of p27^{kip1} protein level of about 100 times compared to pcDNA3-trasfected cell line. Figure 3B shows the densitometric evaluation of Western blot analysis showed in panel A. p27^{kip1} levels following paclitaxel treatment of SaOs-2 cells transfected with empty vector (Fig. 3B, left panel) are comparable with those observed in SaOs-2 parental cell line in Figure 1. Also in p27^{kip1}-overexpressing SaOs-2 cell line (Fig. 3B, right panel), paclitaxel treatment induced a decrease of exogenous p27^{kip1} protein level at 16 h of treatment followed by an increase at 24 and 48 h of paclitaxel treatment. Flow cytometry analysis of Annexin V positivity was performed to assess the percentage of cells committed to apoptosis in the presence or absence of p27^{kip1} overexpression (Fig. 3C–D). Cell cycle distribution in transfected cells treated by paclitaxel was also evaluated: induction of p27^{kip1} expression did not cause p27^{kip1} Promotes Paclitaxel-Induced Apoptosis



Fig. 3. A: Western blot analysis of $p27^{kip1}$ expression in SaOs-2 cells transfected with pcDNA3- $p27^{kip1}$ expression vector or control vector and treated with 100 nM paclitaxel at the indicated times. All the samples were run on the same gel but protein expression in $p27^{kip1}$ -overexpressing samples was obtained from a less exposed film, to avoid the issue of a saturated signal of $p27^{kip1}$ protein. β -actin expression was used as the loading control. **B**: Densitometric evaluation of $p27^{kip1}$ protein level obtained by the Western blot analysis showed in **panel A**. The two groups of histograms represent the effect of paclitaxel treatment on $p27^{kip1}$ level, normalized to β -actin expression, on SaOs-2

 G_1 arrest or apoptosis compared to the parental cell line or to SaOs-2 cells transfected with empty vector only (Table I). As Figure 3C shows, after 48 h of paclitaxel treatment, 41.8% of

cell line trasfected with pcDNA3 empty vector (**left panel**, white bars) or with pcDNA3-p27^{kip1} vector (**right panel**, gray bars) compared to their untreated control samples. **C**: Annexin V-positive/PI negative apoptotic cell percentages in SaOs-2 cells transfected with pcDNA3-p27^{kip1} or pcDNA3 control vector and treated with 100 nM paclitaxel. The data are presented as the mean ± standard deviation of three different experiments. **P* < 0.05 (Student's *t*-test of p27^{kip1}-overexpressing cells vs. pcDNA3-transfected ones). **D**: A representative result of flow cytometric evaluation of apoptosis showed in **panel C**.

SaOs-2 cells transiently transfected with the pcDNA3-p27^{kip1} construct are committed to apoptosis compared to 30.6% of SaOs-2 cells transfected with the empty vector (P < 0.05).

TABLE I. Effect of 100nM Paclitaxel on Cell Cycle Phases of SaOs-2 Cell Line Overexpressing p27 ^{kip1}	;

			G_0/G_1	S	G_2	Apoptosis
Control	16 h	Parental pcDNA3	$56.1 \pm 5.7 \ 51.0 \pm 0.5$	$28.7 \pm 3.3 \\ 20.2 \pm 1.5$	$15.1 \pm 2.4 \\ 28.8 \pm 2.0$	$10.7 \pm 2.3 \\ 11.3 \pm 4.7$
		$p27^{kip1}$	49.4 ± 1.0	26.5 ± 6.7	24.1 ± 7.7	9.6 ± 1.4
Paclitaxel	16 h	pcDNA3 $p27^{kip1}$	$21.3 \pm 1.0 \ 21.4 \pm 0.7$	$32.7 \pm 2.5 \ 31.1 \pm 4.3$	$\begin{array}{c} 46.0 \pm 1.5 \\ 47.5 \pm 5.1 \end{array}$	$\begin{array}{c} 12.1\pm1.1 \\ 9.7\pm3.4 \end{array}$
	24 h	$pcDNA3 p27^{kip1}$	$4.6 \pm 0.1 \\ 5.6 \pm 2.3$	$25.4 \pm 7.1 \\ 29.4 \pm 1.7$	$\begin{array}{c} 70.0 \pm 7.1 \\ 65.0 \pm 4.0 \end{array}$	$12.1 \pm 1.1 \\ 20.8 \pm 5.3$
	48 h	$pcDNA3 p27^{kip1}$	$4.2\pm 0.8\ 3.7\pm 0.6$	$3.8 \pm 4.0 \\ 6.5 \pm 1.8$	$92.3 \pm 4.5 \\ 89.8 \pm 1.3$	$\begin{array}{c} 30.6 \pm 4.2 \\ 41.8 \pm 5.8 \end{array}$

Data are presented as the mean \pm standard deviation of three different experiments.

DISCUSSION

In a previous study, we demonstrated that 100 nM paclitaxel treatment induces apoptosis in the SaOs-2 osteosarcoma cell line [Pucci et al., 1999]. In this study, at 16 h of continuous paclitaxel treatment, we observed a decrease of p27^{kip1} protein levels. Following 24 to 48 h of treatment, times of paclitaxel exposure that induced apoptosis in SaOs-2 cell line [Pucci et al., 1999], we observed that p27^{kip1} is expressed again, suggesting that p27^{kip1} increase in paclitaxel-treated SaOs-2 cells is in conjunction with the activation of the apoptotic program. The broad range caspase inhibitor Z-VAD-FMK treatment is able to inhibit paclitaxel-induced apoptosis [Pucci et al., 1999] and p27^{kip1} increase following 48 h of paclitaxel treatment, suggesting that, in SaOs-2 cells, the increase of p27^{kip1} observed following drug exposure is strictly related to the apoptotic process.

It has been demonstrated that $p27^{kip1}$ is regulated primarily at the post-transcriptional level by the ubiquitin-dependent proteasome pathway [Pagano et al., 1995]. Paclitaxel treatment induces an activation of the ubiquitinproteasome complex, determining a degradation of p27^{kip1} during the first 16 h of treatment. In conjunction with the activation of apoptotic program, the proteasome activity is inhibited determining an increase of p27^{kip1} protein. These results suggest that the inhibition of the proteasome activity observed in SaOs-2 cells following 48 h of paclitaxel treatment may account for the abnormal accumulation of p27^{kip1}, which may in turn play an important role in sustaining the apoptosis of transformed cells. Although the mechanism responsible for the decrease of proteasome activity in SaOs-2 cells at 48 h of treatment is unclear, it is conceivable that several subunits of the proteasome undergo caspase-dependent proteolysis and that this results in diminished protein turnover, as recently reported [Adrain et al., 2004]. Moreover, the pro-apoptotic role of lack of proteasome activity in paclitaxelinduced cell death has been observed in a human lung adenocarcinoma cell line (LC-2-AD): inhibition of proteasome by the proteasome inhibitor 1 (PS1) in these cells treated with paclitaxel leads to an increase in the number of apoptotic cells [Oyaizu et al., 2001]. The involvement of p27^{kip1} accumulation in proteasome

inhibitor-induced apoptosis has already been reported in several human carcinoma cells [An et al., 1998; Kudo et al., 2000] and the adenoviral expression of a degradation-resistant form of $p27^{kip1}$ can induce growth arrest and apoptosis in a breast cancer cell line model [Zhang et al., 2005].

To understand the significance of $p27^{kip1}$ protein modulation during paclitaxel-induced apoptosis, we examined the effect of a transient overexpression of $p27^{kip1}$ on SaOs-2 cell line sensibility to paclitaxel. Unexpectedly, $p27^{kip1}$ did not cause any G₁ arrest of SaOs-2 cells in our study. The explanation may be found in the genotype of SaOs-2 cells. Indeed, this osteosarcoma cell line expresses a truncated form of pRb that is defective for E2F binding and cell cycle regulation [Shew et al., 1990]; therefore, even if cyclin D/cdk4 and cyclin E/cdk2 are inhibited by $p27^{kip1}$, the cell cycle is not arrested.

In our study, transient overexpression of $p27^{kip1}$ protein in SaOs-2 cell line determines an increase of paclitaxel-induced apoptosis. Several studies, mostly in normal cells and in vivo, have pointed toward a protective role in apoptosis for $p27^{kip1}$ [St Croix et al., 1996; Eymin et al., 1999]. However, additional research using adenoviral vectors to overexpress $p27^{kip1}$ in several cancer cells cell lines [Craig et al., 1997; Katayose et al., 1997; Wang et al., 1997] has shown that $p27^{kip1}$ may play a role in promoting apoptosis by inducing G₁ arrest in the presence of a functional pRb [Schreiber et al., 1999; Naruse et al., 2000].

Regarding the pro-apoptotic role of $p27^{kip1}$ following paclitaxel treatment, our data are in contrast with a previous study in RKO human colon adenocarcinoma cells, where overexpression of $p27^{kip1}$ slightly inhibited taxol-induced cell death [Schmidt et al., 2000]. The use of two different cell lines may account for this difference. Again, SaOs-2 cells are pRb-defective, thus $p27^{kip1}$ could be ineffective in inducing G_1 arrest, whereas in RKO cells, $p27^{kip1}$ can induce G_1 arrest by increasing the levels of pRb, thus preventing cells from arresting in the $G_2/$ M phase.

In this study, we have shown that an increase in $p27^{kip1}$ levels occurs in vivo during paclitaxel treatment of SaOs-2 cells due to the blockade of proteasomal activity. Moreover, $p27^{kip1}$ overexpression is able to inhibit cell proliferation leading to apoptosis in a G₁-independent manner. The mechanism underlying a proapoptotic role of $p27^{kip1}$ is still unclear. It is conceivable that $p27^{kip1}$ overexpression could act by sending an aberrant signal to those cells that are able to escape paclitaxel-induced apoptosis in G₂/M phase, or by downregulating anti-apoptotic molecules such as mcl-1 [Woltman et al., 2003].

In conclusion, we have demonstrated that transfection of pRb-null SaOs-2 cells with the $p27^{kip1}$ gene via plasmid leads to an increase of paclitaxel-induced apoptosis. The combined effect of $p27^{kip1}$ on proliferation and apoptosis in tumor cells makes $p27^{kip1}$ and proteasomal inhibitors interesting tools for therapy in patients with pRb-defective cancers.

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