

# The Cdc2/Cdk1 inhibitor, purvalanol A, enhances the cytotoxic effects of taxol through Op18/stathmin in non-small cell lung cancer cells *in vitro*

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**Abstract.** Purvalanol A is a highly selective inhibitor of Cdc2 [also known as cyclin-dependent kinase 1 (CDK1)]. Taxol is an anti-tumor chemotherapeutic drug which is widely used clinically. In this study, the CDK1 inhibitor, purvalanol A was applied to explore the relevance of Cdc2 signaling and taxol sensitivity through analyses, such as cellular proliferation and apoptosis assays, ELISA, western blot analysis and immunoprecipitation. We demonstrated that purvalanol A effectively enhanced the taxol-induced apoptosis of NCI-H1299 cells, as well as its inhibitory effects on cellular proliferation and colony formation. In combination, purvalanol A and taxol mainly decreased the expression of oncoprotein 18 (Op18)/stathmin and phosphorylation at Ser16 and Ser38, while purvalanol A alone inhibited the phosphorylation of Op18/stathmin at all 4 serine sites. Co-treatment with purvalanol A and taxol weakened the expression of Bcl-2 and activated the extrinsic cell death pathway through the activation of caspase-3 and caspase-8. Further experiments indicated that Cdc2 kinase activities, including the expression of Cdc2 and the level of phospho-Cdc2 (Thr161) were significantly higher in taxol-resistant NCI-H1299 cells compared with the relatively sensitive CNE1 cells before and following treatment with taxol. These findings suggest that Cdc2 is positively associated with the development of taxol resistance. The Cdc2 inhibitor, purvalanol A, enhanced the cytotoxic effects of taxol through Op18/stathmin. Our findings may prove to be useful in clinical practice, as they may provide a treatment strategy with which to reduce the doses of taxol applied clinically, thus alleviating the side-effects.

## Introduction

Cdc2, also known as cyclin dependent kinase 1 (CDK1), controls the cell cycle entry from the G2 to the M phase and promotes the commencement of mitotic phase events (1), the abnormal activation of which directly causes aberrant cell proliferation, and malignant transformation and tumorigenesis in prostate cancer cells (2-4). Cdc2 activation also depends on the phosphorylation of Thr161, and CDC25-mediated dephosphorylation at Thr14 and Tyr-15, which exhibits enzymatic activity when only phospho-Thr161 remains (5).

Purvalanol A is a selective inhibitor of Cdc2, which strongly inhibits Cdc2 kinase activity at a low concentration of 2  $\mu$ M (6,7). Investigators have identified that purvalanol A effectively suppresses Cdc2 activity and the progression from the G2 phase to mitosis, which leads to the loss of clonogenicity and cellular apoptosis in both MKN45 and MKN28 X-irradiated gastric cancer cells (8).

Oncoprotein 18 (Op18)/stathmin is a small molecule weight phosphoprotein which is highly expressed in cancer cells. Its main functions are to regulate the equilibrium of microtubule (MT) dynamics and control cell cycle progression, which is closely associated with the maintenance of tumor malignant phenotypes (9-12). Op18/stathmin has 4 phosphoserine sites (p-Ser16, p-Ser25, p-Ser38 and p-Ser63), which integrates and relays various signals from intra- or extracellular stimuli through phosphorylated inactivation and dephosphorylated activation (13-15).

In a previous study, we found that Epstein-Barr virus-specific protein-latent membrane protein 1 (LMP1) regulates the Op18/stathmin signaling pathway by mediating Cdc2, which accelerates cell cycle progression and promotes cell proliferation (16). In another recent study of ours, we confirmed that human NCI-H1299 non-small cell lung cancer cells highly expressing Op18/stathmin were the most highly resistant to taxol among 5 different cancer cells originating from epithelia, including CNE1, Hep3B-2, MGC, MCF-7 and NCI-H1299 (17).

In this study, NCI-H1299 cells were employed to clarify the association between Cdc2 signaling and taxol resistance, and to elucidate the related molecular mechanisms.

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## Materials and methods

**Cells and cell growth conditions.** Both the CNE1 cells and NCI-H1299 cells were a kind gift from Professor Ya Cao from the Cancer Research Institute of Central South University and cultured in RPMI-1640 (Gibco-BRL, Grand Island, NY, USA) with 10% fetal bovine serum (FBS; HyClone, Logan, UT, USA), 100 IU/ml penicillin and 100 µg/ml streptomycin at 37°C, 5% CO<sub>2</sub>.

CNE1 human cancer cells were testified to be the most sensitive to taxol among 5 various epithelial-deriving tumor cells in our previous study (17). Thus, they were used in this study for a comparison to the NCI-H1299 cells.

**Antibodies and reagents.** The primary antibodies used were anti-stathmin (Cat. no. 569391; Calbiochem, Billerica, CA, USA), anti-phosphoserine (Cat. no. 618100; Zymed, San Francisco, CA, USA), anti-Cdc2 (sc-954; Santa Cruz Biotechnology, Inc., Santa Cruz, CA, USA), anti-phospho-Thr161-Cdc2 (Cat. no. 9114; Cell Signaling Technology, Danvers, MA, USA), anti-cyclin B1 (sc-7393), anti-caspase-3 (sc-7272) (both from Santa Cruz Biotechnology, Inc.), anti-caspase-8 (Cat. no. 9746), rabbit monoclonal anti-caspase-9 (Cat. no. 9502) (both from Cell Signaling Technology), anti-phospho-stathmin [phospho-S16 (ab47328), S25 (ab194752), S38 (ab194757), S63 (ab76583); Abcam, Cambridge, MA, USA], anti-Bcl-2 (sc-492; Santa Cruz Biotechnology, Inc.), anti-extracellular signal-regulated kinase 1 (ERK1; sc-93), anti-phospho-ERK1 (sc-7383) (both from Santa Cruz Biotechnology, Inc.), anti-β-actin (Cat. no. A2228; Sigma, St. Louis, MO, USA). The secondary antibodies used were horseradish peroxidase (HRP)-conjugated goat anti-rabbit IgG (sc-2004) and rabbit anti-mouse IgG (sc-358914) (both from Santa Cruz Biotechnology, Inc.).

Taxol (Santa Cruz Biotechnology, Inc.) and purvalanol A (Calbiochem) were both purchased and dissolved in dimethyl sulfoxide (DMSO) as mother solutions at -20°C for use.

**Flow cytometric analysis.** The cells were plated in 6-well plates, and pre-treated with 0, 1 and 5 µM purvalanol A for 2 h when they reached 80% confluency. This was followed by the addition of 100 nM taxol for 12 h. DMSO was used as a solvent control. The cells were then rinsed with phosphate-buffered saline (PBS) and digested with 0.25% trypsin.

The cells were cropped by centrifugation and stained with a mixture of 5 µl propidium iodide (PI) and 10 µl Annexin V-FITC according to the protocol provided with the Annexin V-FITC Apoptosis Detection kit (Nanjing Biobio Tech. Co, Ltd, Nanjing, China).

Cellular apoptosis was assessed by flow cytometry (FCM) by a specialized agency (the Second Xiangya Hospital Affiliated Central South University). All experiments were performed in triplicate.

**MTT assay.** The cells at the logarithmic phase were seeded at 5,000 cells/well in 96-well plates and pre-treated with various concentrations of purvalanol A (0, 1 and 5 µM) for 2 h, followed by the addition of 100 nM taxol. Finally, 10 µl of 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl (MTT) were added at the specified time points of 24, 48 and 72 h. The values

of optical density (OD) at a wavelength 570 nm were then detected using a microplate reader (BioTek, Winooski, VT, USA). Cell proliferation was calculated using the following formula: relative proliferation rates (%) = (OD treatment/OD control) x 100%. Six parallel wells were set in each group.

**Colony formation assay.** A total of 2,500 cells in a single suspension were plated per well in a 6-well plate and divided into 6 groups, including 3 co-treatment groups and 3 gradient groups of purvalanol A alone. Cell growth was terminated when colonies were observed by the naked eye after 2 weeks. Colonies containing over 50 cells were counted using an inverted microscope, and images were acquired using a DMCI microscope (Leica, Wetzlar, German) at x200 magnification. All experiments were performed in triplicate.

**Assessment of Cdc2 kinase activity.** Assays to determine Cdc2 kinase activity were performed according to the instructions provided with the MESACUP Cdc2 kinase assay kit (code no. 5235; MBL, Nagoya, Japan). Cells reaching 80% confluence were treated with various concentrations of taxol (0, 10 and 100 nM) for 12 h, then lysed in a sample buffer. The supernatants were then collected for the detection of Cdc2 kinase activity as described in our previous study (16). ELISA was performed according to the instructions provided with the MESACUP Cdc2 Kinase Assay kit. Briefly, 100 µl of cell extracts were transferred to a microwell strip coated with monoclonal antibody 4A4 (4 parallel wells/sample), incubated at 25°C for 60 min and washed 5 times. This was followed by the addition of 100 µl peroxidase (POD) conjugated streptavidin, and incubation at 25°C for 30 min and washing 5 times. The cells were then treated with 100 µl POD substrate solution for 5 min, and the reaction was then terminated with 100 µl Stop Solution (20% phosphoric acid). The OD value was read at a 490 nm wavelength using a microplate reader (BioTek). All experiments were carried out in triplicate.

**Western blot analysis.** Following the removal of the supernatant, the cells were lysed in cell lysis buffer (50 mM Tris-HCl pH 8.0, 1 mM EDTA, 2% SDS, 5 mM DTT and 10 mM PMSF). Total proteins (50 µg) were then separated by 10% sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) and transferred onto nitrocellulose membranes and incubated with the specific primary antibodies at 4°C overnight, followed by the addition of HRP conjugated secondary antibodies for 2 h at room temperature. An enhanced chemiluminescence detection kit (Pierce, Rockford, IL, USA) was applied for immunoblotting.

**Immunoprecipitation (IP) - western blot analysis.** In brief, cell extracts were collected in IP lysis buffer, as described in our previous studies (16,18). Immunoprecipitates of Op18/stathmin were pulled down by excessive anti-stathmin antibody binding magnetic immunobeads and separated by 10% SDS-PAGE. anti-phosphoserine antibody was used to detect the total level of phosphorylated Op18/stathmin by western blot analysis.

**Statistical analysis.** Statistical analysis was carried out using SPSS 17.0 statistical software. The specific statistical method

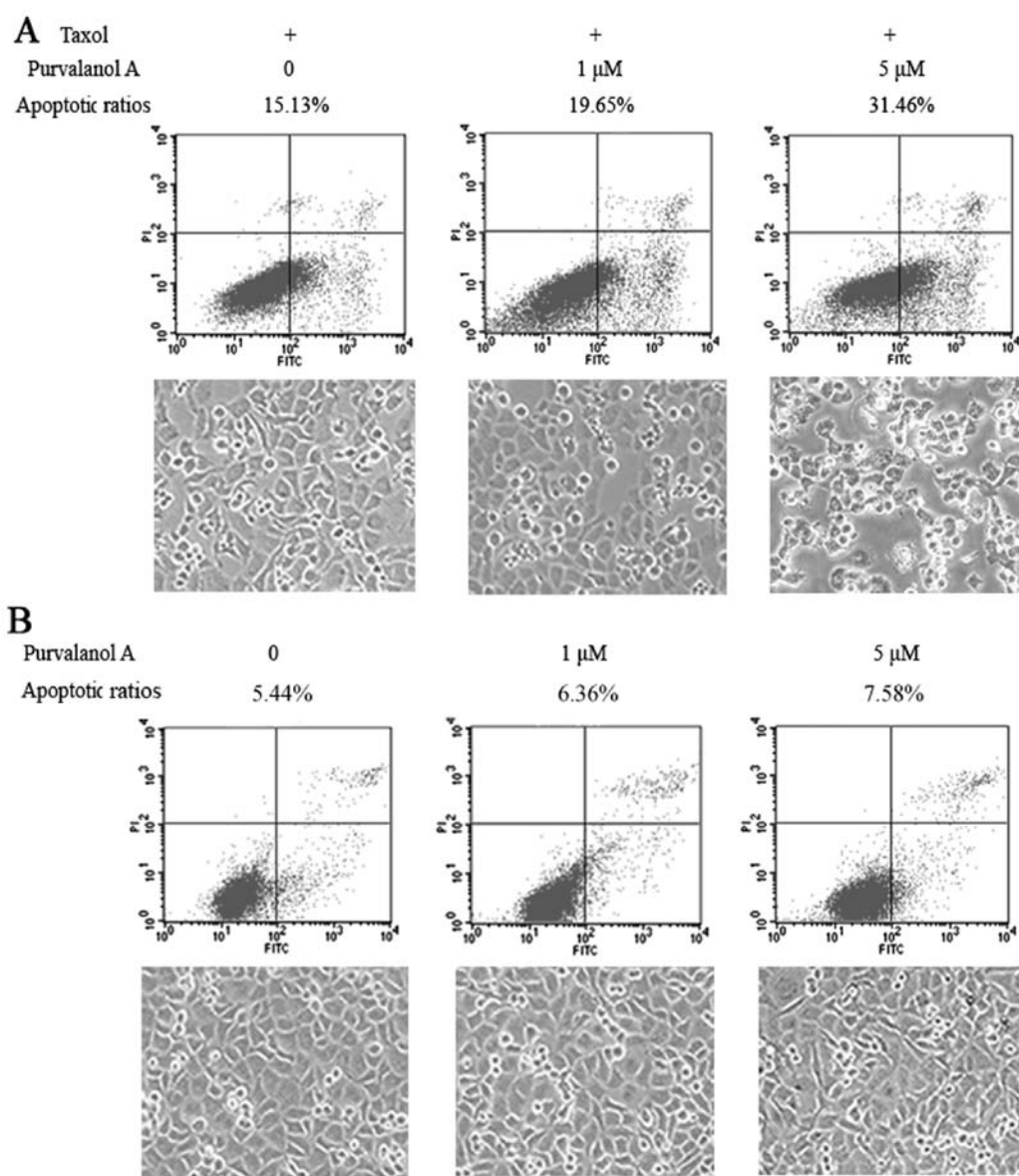


Figure 1. Purvalanol A promotes taxol-induced cellular apoptosis and growth inhibition. (A) Upper panel shows cellular apoptotic ratios, and the images in the lower panel represent the results of cellular growth. (B) Status of cellular apoptosis and growth in cells treated with purvalanol A alone at increasing concentrations.

applied was the t-test, and a value of  $p < 0.05$  was considered to indicate a statistically significant difference. All data are presented as the means  $\pm$  SD.

## Results

*Purvalanol A enhances the taxol-induced apoptosis of NCI-H1299 cells.* The cellular apoptotic ratios were 15.13, 19.65 and 31.46% in the NCI-H1299 cells treated with taxol in presence of 0, 1 and 5  $\mu$ M purvalanol A, respectively. Purvalanol A augmented taxol-induced cellular apoptosis in a concentration dependent manner. The images of cellular growth revealed that a greater number of translucent floating cells appeared in the medium of the cells treated with purvalanol A (Fig. 1A).

In the cells treated with purvalanol A alone at concentrations of 0, 1 and 5  $\mu$ M, the cellular apoptotic ratios were 5.44, 6.36

and 7.58%, respectively. These cells almost reached complete confluence and there were only a few translucent floating cells (Fig. 1B).

*Purvalanol A enhances the inhibitory effects of taxol on cellular proliferation and colony formation in NCI-H1299 cells.*

Purvalanol A enhanced the inhibitory effects of taxol on cellular proliferation in NCI-H1299 cells. In the cells co-treated with 5  $\mu$ M purvalanol A and taxol, the representative curve steeply descended at the time points of 24, 48 and 72 h; the relative proliferation ratios were 80.53, 59.01 and 41.12%, respectively. In the cells co-treated with 1  $\mu$ M purvalanol A and taxol, the representative curve descended slowly, with proliferation ratios of 94.98, 89.6 and 78.32% at 24, 48 and 72 h, respectively (Fig. 2A).

The histograms indicated that a statistically significant difference existed between the cells co-treated with 5  $\mu$ M purvalanol A and taxol, and the other 2 groups (taxol only,

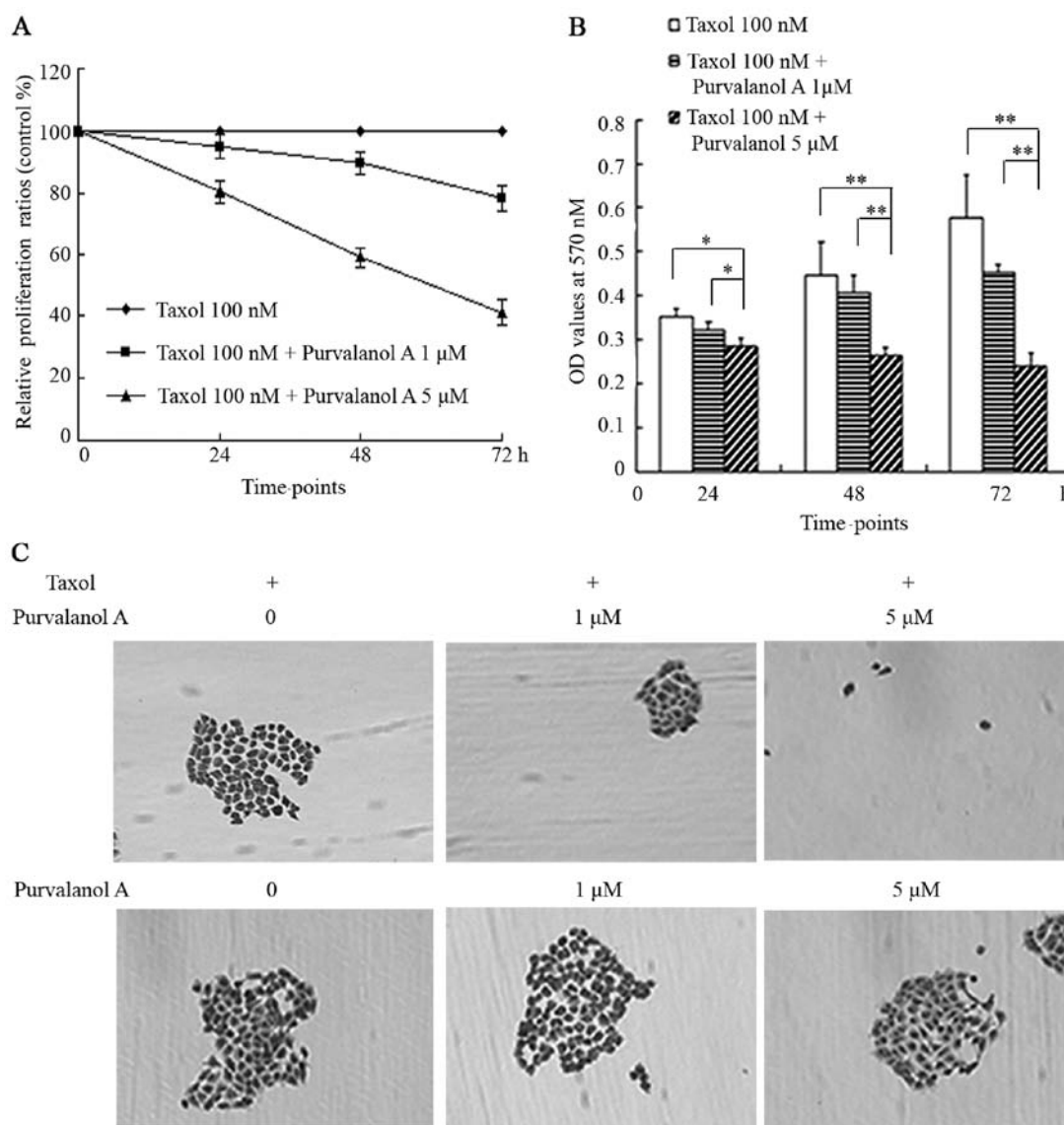


Figure 2. Combination of purvalanol A and taxol suppresses cellular proliferation and colony formation. (A) Curves represent proliferation trends at 3 time points. (B) Histograms show statistically significant differences. \* $p < 0.05$  and \*\* $p < 0.01$ . (C) Images show the capacities of colony formation. Original magnification,  $\times 200$ .

and taxol + 1  $\mu$ M purvalanol A) at 24 h ( $p < 0.05$ ), while the differences became more and more significant with the passing of time ( $p < 0.01$ ) (Fig. 2B).

Only a few and sparse small colonies appeared in the group treated with 1  $\mu$ M purvalanol A combined with taxol; no visibly typical colonies were formed in the group co-treated with 5  $\mu$ M purvalanol A and taxol, while a large number of colonies appeared among the group treated with taxol only and in the 3 control groups treated with purvalanol A only at increasing concentrations, which usually merged into a large colony with ambiguous borders (Fig. 2C).

*Co-treatment with purvalanol A and taxol inhibits the expression of Op18/stathmin and phosphorylation.* The expression of Op18/stathmin was weakened in the NCI-H1299 cells treated with a combination of purvalanol A and taxol; the synergistic inhibitory effects were the most evident in the cells treated with a combination of 5  $\mu$ M purvalanol A

and taxol in comparison with the other 2 groups (Fig. 3A). Treatment with taxol alone slightly decreased the expression of Op18/stathmin at the concentration of 100 nM; however, no notable alternation in the expression of Op18/stathmin was observed in the cells treated with purvalanol A at increasing concentrations (Fig. 3B).

IP analysis revealed that the total levels of phospho-Op18/stathmin were notably decreased in the cells co-treated with purvalanol A and taxol, compared with the controls or the cells treated with taxol or purvalanol A alone (Fig. 3C).

Treatment with purvalanol A alone inhibited the phosphorylation of Op18/stathmin at all 4 serine sites, including Ser16, Ser25, Ser38 and Ser63 compared with the cells treated with taxol alone. Treatment with both purvalanol A and taxol mainly inhibited the phosphorylation of Op18/stathmin at Ser16 and Ser38 compared with the cells treated with purvalanol A alone; however, the phosphorylation levels of Ser25 and Ser63 sites were similar between the 2 groups (Fig. 3D).

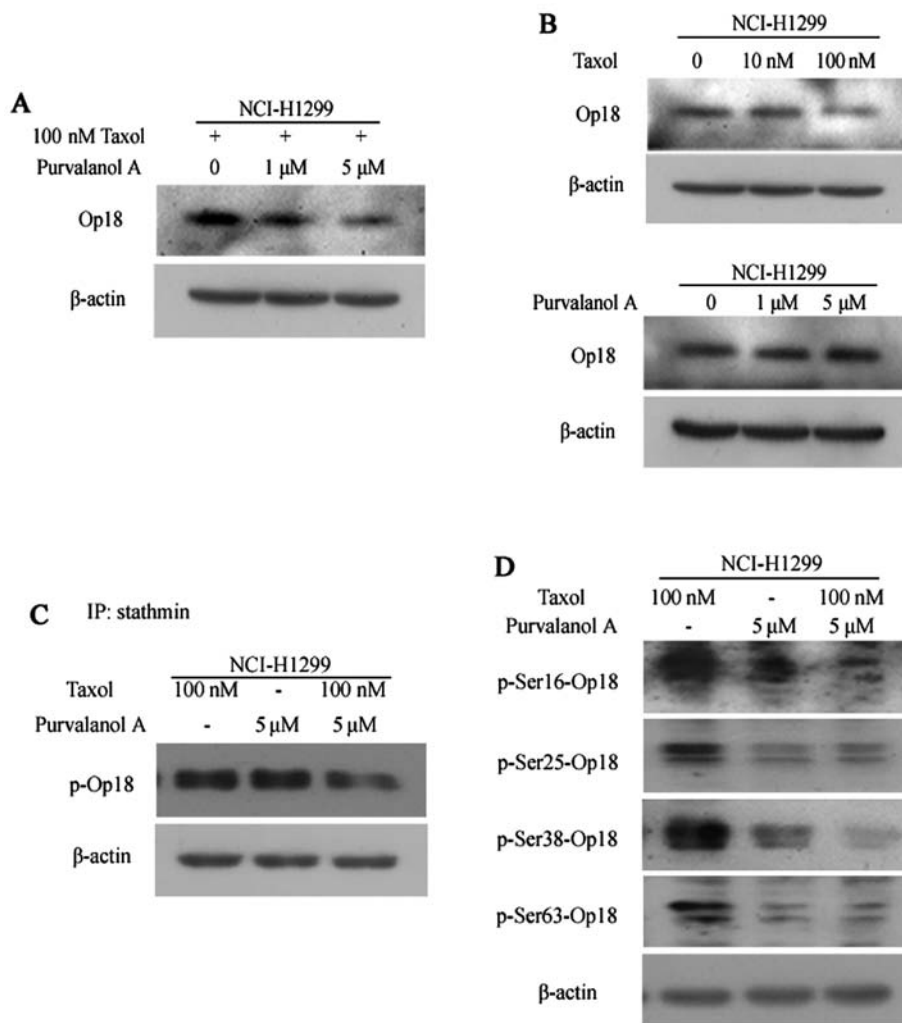


Figure 3. Purvalanol A combined with taxol jointly inhibits the expression of oncoprotein 18 (Op18)/stathmin and phosphorylation. (A and B) The expression of Op18/stathmin was detected by western blot analysis. (C) Immunoprecipitation (IP) - western blot was used to detect the total levels of phospho-Op18/stathmin. (D) Purvalanol A combined with taxol mainly inhibited the phosphorylation of Op18/stathmin at the Ser16 and Ser38 sites.

Treatment with both purvalanol A and taxol decreases the expression of *Bcl-2* and initiates the activation of *caspase-3* and *caspase-8*. The expression and phosphorylation of ERK was not altered in the cells treated with both purvalanol A and taxol. However, combination treatment markedly inhibited the expression of *Bcl-2*, with a greater decrease observed with increasing concentrations of purvalanol A (Fig. 4A). No differences were observed in the expression of these molecules in the cells treated with purvalanol A alone at increasing concentrations (Fig. 4B).

Purvalanol A and taxol collectively increased the expression levels of *caspase-3* and *caspase-8* with a greater increase observed with increasing concentrations of purvalanol A. No obvious changes were observed in the levels of *caspase-9* (Fig. 4C). Similarly, treatment with purvalanol A alone at increasing concentrations did not lead to any alternations in the levels of *caspase-3*, *caspase-8* and *caspase-9* in the NCI-H1299 cells (Fig. 4D).

*Cdc2* is positively associated with the development of taxol resistance in different epithelia-derived tumor cells. The histograms indicated that *Cdc2* kinase activity was markedly

higher in the NCI-H1299 cells compared with the CNE1 cells before and after treatment with taxol. Taxol also downregulated the activity of *Cdc2* to a certain extent in a concentration-dependant manner in both cell lines (NCI-H1299 and CNE1), although this decrease in the levels *Cdc2* was more evident in the CNE1 cells in comparison to the NCI-H1299 cells; the differences became more significant with the addition of Taxol ( $p < 0.01$ ) (Fig. 5A).

The results of western blot analysis revealed that the expression levels of *Cdc2* and phosphorylation at the Thr161 site were markedly higher in the NCI-H1299 cells than in the CNE1 cells before and after treatment with taxol. Treatment with taxol alone did not affect the expression of *Cdc2*, but slightly decreased the phosphorylation of *Cdc2*-Thr161 in both cell lines, which was coincident with the analysis of *Cdc2* kinase activity by enzyme-linked immunosorbent assay (ELISA) (Fig. 5B).

FCM analysis revealed that the cellular apoptotic ratios were 12.53 and 34.94% in the NCI-H1299 cells and CNE1 cells treated with taxol, while the cellular apoptotic ratios were 5.02 and 5.49% in the controls treated with solvent DMSO; the status of cell growth shown in the images correlated with the findings of apoptotic analysis (Fig. 5C and D).



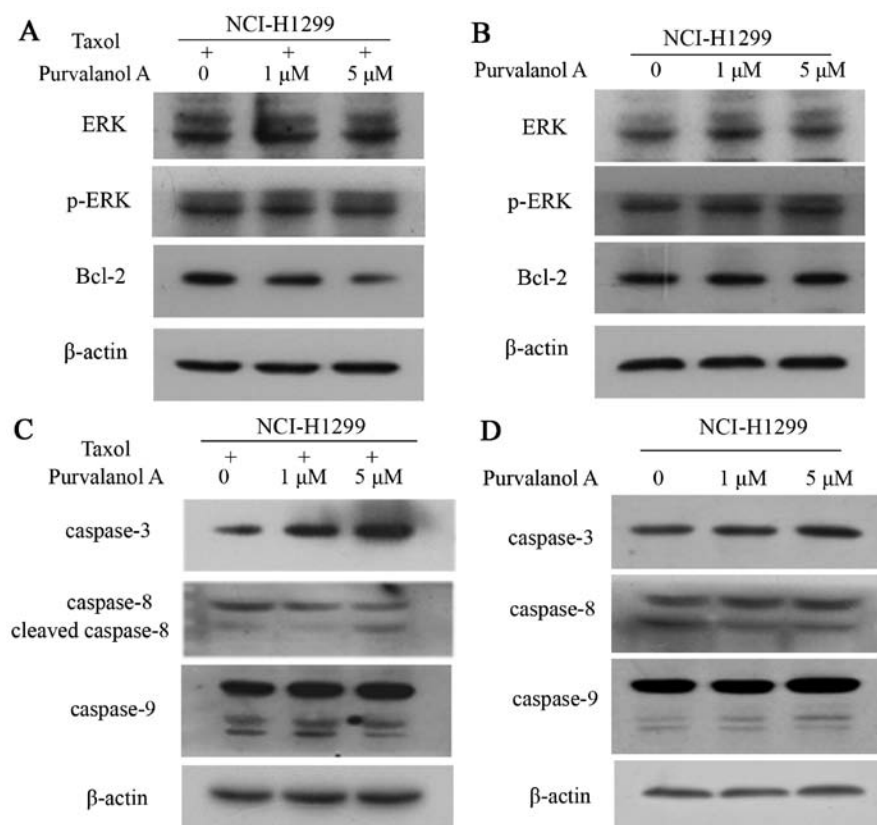


Figure 4. Combination of purvalanol A and taxol decreases the expression of Bcl-2 and initiates the activation of caspase-3 and caspase-8, while the expression of extracellular signal-regulated kinase (ERK) and p-ERK was not affected. (A) Cells were treated with a combination of purvalanol A and taxol. (B) Cells treated with purvalanol A at increasing concentrations were used as controls. (C) Status of caspase-3, caspase-8 and caspase-9 was detected in the co-treated groups. (D) No any obvious changes were observed in the cells treated with purvalanol A alone at increasing concentrations.

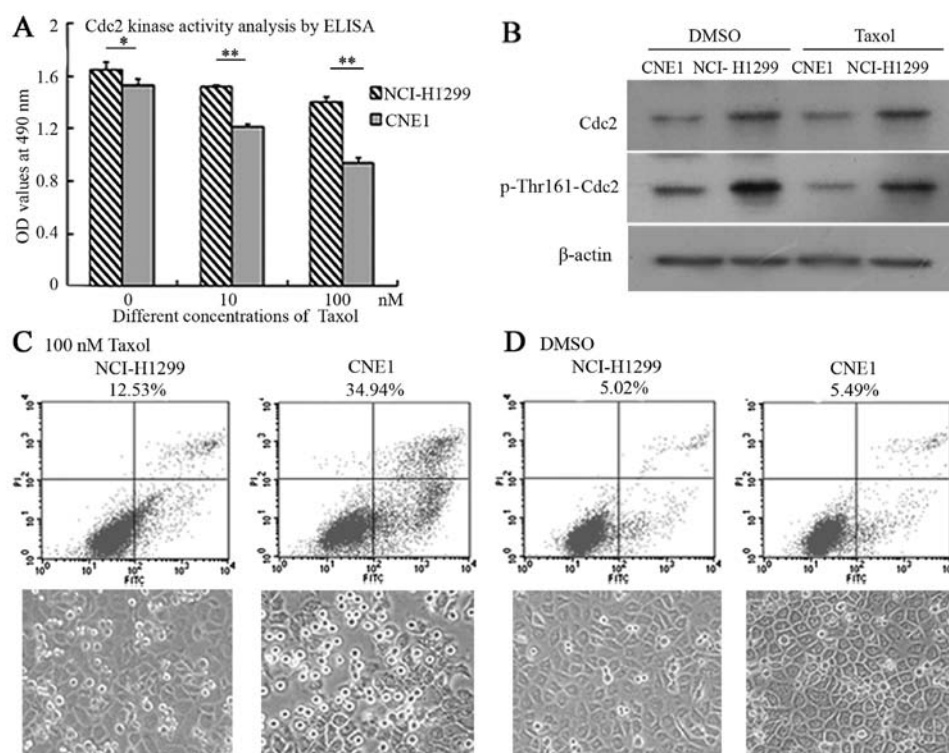


Figure 5. Cdc2 is positively associated with the development of taxol resistance in different epithelia-derived tumor cells. (A) Histograms show the differences in Cdc2 kinase activity by enzyme-linked immunosorbent assay (ELISA) between the NCI-H1299 cells and CNE1 cells. \* $p < 0.05$  and \*\* $p < 0.01$ . (B) Western blot analysis was used to detect the expression of Cdc2 and the level of p-Thr161-Cdc2. (C) Upper panel shows cellular apoptotic ratios in cells treated with taxol, and the images in the lower panel illustrate the status of cell growth. (D) Upper panel shows apoptotic ratios in different tumor cells treated with solvent dimethyl sulfoxide (DMSO), and the images in the lower panel indicate the growth situation of control cells.

## Discussion

Taxol is an effective antitumor chemotherapeutic agent derived from plants, which induces cell cycle arrest and cellular apoptosis by promoting MT polymerization (19-21). With the wide application of taxol in clinical practice, a large number of resistant tumors have emerged; therefore, high doses of taxol are increasingly prescribed, which results in severe side-effects. To date, no antagonizing agents have been identified (22). To minimize the dosage of taxol, the combined use of taxol with other traditional drugs has been used in an attempt to improve taxol sensitivity (23,24). These strategies have achieved some success, but also lead to new risks of acquiring multidrug resistance.

In this study, we demonstrated that the Cdc2/CDK1 inhibitor, purvalanol A, effectively enhanced the sensitivity of NCI-H1299 cells to taxol and inhibited cellular proliferation and colony formation. Other studies have also confirmed that CDK1 inhibition significantly enhances drug-induced colony formation and apoptosis in breast cancer cells and colon carcinoma cells and glioma cells (25-27).

Op18/stathmin is a downstream molecular target of Cdc2, and tumor cells which highly express Op18/Stathmin are resistant to taxol (17,28). This study certified that the combination of purvalanol A and taxol mainly inhibited the expression of Op18/stathmin and phosphorylation at the Ser16 and Ser38 sites, while purvalanol A alone uniformly inhibited the phosphorylation of Op18/stathmin at all 4 serine sites. Other studies have shown that Cdc2 predominantly phosphorylates Op18/stathmin at Ser25 and Ser38 sites, which is partly different from our results, which may implicate other inhibitory activities of purvalanol A (6,29).

Bcl-2 is an anti-apoptotic factor involved in the resistance of conventional drugs. The overexpression of Bcl-2 potentially induces the development of taxol resistance in lung cancer cells (30). ERK is also an upstream kinase of Op18/stathmin, and ERK-mediated Op18/stathmin signaling also complicates to cell cycle progression and taxol sensitivity (18,31). Cellular apoptosis is closely associated with the activation of caspase-3, caspase-8 and caspase-9. As previously demonstrated, the blocking of ERK and taxol jointly induced exogenous cellular apoptosis via the upregulation of the expression of caspase-3 and caspase-9 in NCI-H1299 cells; the activation of caspase-3 and caspase-8 promoted cellular apoptosis and autophagy in A549 human lung cancer cells exposed to *Paris polyphylla* steroidal saponins (PPSS); paclitaxel and Op18 silencing collectively induced cell death through initiating caspase-3 and caspase-9 activations in nasopharyngeal carcinoma CNE1 cells (18,32,33). This study indicated that co-treatment with purvalanol A and taxol decreased the expression of Bcl-2 and initiated the activation of the extrinsic cell death pathway through the activation of caspase-3 and caspase-8, but did not markedly affect the expression of ERK and phosphorylation.

We further found that both the expression of Cdc2 and the level of p-Thr161-Cdc2 was markedly higher in the NCI-H1299 cells in contrast with the CNE1 cells, which was consistent with the detection of Cdc2 kinase activity, while the NCI-H1299 cells with a high Cdc2 kinase activity exhibited obvious resistance to taxol in comparison with the relatively sensitive CNE1 cells, which implied that Cdc2 kinase activity is positively associated with the development of taxol resistance in different epithelia-

derived cancer cells. Another study also demonstrated that a high expression of Cdc2 was negatively associated with the curative effects of chemotherapeutics and was a poor prognostic factor in epithelial-derived ovarian cancer and laryngeal squamous cell carcinoma (34,35).

In conclusion, the findings of our study suggest that purvalanol A enhances the cytotoxic effects of taxol through Op18/stathmin in non-small cell lung cancer. Our findings may prove to be helpful in reducing the doses of taxol applied clinically and to alleviate the side-effects.

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