

Curcumin-induced downregulation of Axl receptor tyrosine kinase inhibits cell proliferation and circumvents chemoresistance in non-small lung cancer cells

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Abstract. Lung cancer is still in the first place in terms of both incidence and mortality. In the present study, we demonstrated the effect of curcumin, a phytochemical of the plant *Curcuma longa*, on expression and activation of Axl receptor tyrosine kinase (RTK) which plays an important role in cell survival, proliferation and anti-apoptosis. Curcumin treatment of non-small cell lung cancer (NSCLC) A549 and H460 cells, was found to decrease Axl protein as well as mRNA levels in a dose- and time-dependent manner. Axl promoter activity was also reduced by curcumin, indicating that curcumin down-regulates Axl expression at the transcriptional level. Moreover, Axl phosphorylation in response to binding of its ligand, Gas6, was abrogated by curcumin, suggesting the inhibitory effect of curcumin on Gas6-induced Axl activation. We next found cytotoxic effect of curcumin on both the parental A549 and H460 cells, and their variants which are resistant to cisplatin (A549/CisR and H460/CisR) and paclitaxel (A549/TR and H460/TR). Exposure of these cells to curcumin resulted in dose-dependent decline of cell viability and clonogenic ability. It is further observed that the anti-proliferative effect of curcumin on A549 cells overexpressing Axl protein was reduced, while that on H460 cells transfected Axl specific siRNA was augmented, confirming that curcumin inhibits cell proliferation via downregulation of Axl expression. In addition, curcumin was found to cause the induction of p21, a

cyclin-dependent kinase inhibitor, and reduction of X-linked inhibitor of apoptosis protein (XIAP), an anti-apoptotic molecule, in parental H460 cells as well as chemoresistant cells, H460/CisR and H460/TR. Taken together, our data imply that Axl RTK is a novel target of curcumin through which it exerts anti-proliferative effect in both parental and chemoresistant NSCLC cells.

Introduction

Worldwide, lung cancer is the most frequently diagnosed cancer and the leading cause of cancer deaths (1). Approximately 85-90% of lung cancer is non-small cell lung cancer (NSCLC) which is characterized by relatively low growth rate and poor responsiveness upon chemotherapy, compared to small cell lung cancer (SCLC) (2). Although platinum or taxol-based chemotherapy has been the standard treatment for NSCLC patients, the intrinsic and acquired resistances to these drugs are the major obstacles to achieve the successful long-term outcomes. To overcome the resistance, the second line or combination chemotherapy regimens have been used (3-5), but the overall survival benefits of various chemotherapies in NSCLC is not yet satisfactory.

The large receptor tyrosine kinase (RTK) family in the human genome contains 58 RTKs and is divided into 20 subfamilies. One of the subfamilies is TAM family composed of three RTK members which are Tyro3 (also referred to Brt, Dtk, Rse, Sky or Tif), Axl (also referred to Ark, Tyro7 or Ufo) and Mer (also referred to Eyk, Nyk or Tyro12) (6-9). Each of them has similar structural features, which are extracellular domains, two immunoglobulin-like and two fibronectin type III domains and cytoplasmic kinase domain (10,11). Among several ligands including growth arrest-specific 6 (Gas6), protein S, tubby and tulip, Gas6 is the only one able to bind and activate all three TAM RTKs, which evokes transduction of many extracellular signals to cause cell growth, survival, proliferation, migration and inhibition of apoptosis (11,12).

Since the first identification of Axl in 1988, *Axl* cDNA was cloned in 1991 from chronic myelogenous leukemia patients as a novel RTK (7). Axl overexpression and its activation

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Abbreviations: EGFR, epidermal growth factor receptor; Gas6, growth arrest-specific 6; NSCLC, non-small cell lung cancer; RTK, receptor tyrosine kinase; XIAP, X-linked inhibitor of apoptosis

Key words: Axl, chemoresistance, curcumin, Gas6, non-small cell lung cancer, receptor tyrosine kinase, X-linked inhibitor of apoptosis

upon Gas6 stimulation have been reported in many types of cancer such as acute leukemia (13), breast (14), colon (15), esophageal, lung (16), ovarian (17), prostate (18) and thyroid cancer (19), which subsequently signals for cell survival and proliferation (20-22). For example, in more than half of non-small cell lung cancer (NSCLC) cell lines, the levels of Ax1, Mer and ligands, Gas6 and protein S, were elevated (23-25). In 48.3% of clinical samples of lung adenocarcinoma, Ax1 overexpression was observed, which was also associated with disease stages and lymph node metastasis (25). Therefore, Ax1 has been receiving increased attention as a potent therapeutic target for cancer treatment.

Curcumin, a polyphenolic compound extracted from *Curcuma longa*, is a well-known natural product with anti-oxidant, anti-inflammatory and anticancer activities (26,27). Recent clinical investigations demonstrated that curcumin has not only chemo-preventive, but also therapeutic potential in many types of cancer. Moreover, oral administration of up to 12 g/day of curcumin was reported to be safe enough, suggesting its low toxicity at effective dose. Curcumin has also been demonstrated to modulate many signal transduction pathways involved in survival, carcinogenesis and apoptosis (28-30).

In the present study, we examined the effect of curcumin on expression and activation of Ax1 RTK in NSCLC cells, which subsequently inhibits cell proliferation and overcomes chemo-resistance via both induction of p21 and reduction of X-linked inhibitor of apoptosis (XIAP), suggesting that Ax1 RTK is a novel target of curcumin to exert its anticancer activity.

Materials and methods

Reagents and antibodies. Curcumin was obtained from Sigma-Aldrich (St. Louis, MO, USA). A549 and H460 cells were purchased from the American Type Culture Collection (ATCC; Manassas, VA, USA). Primers for Ax1 were synthesized by the domestic company, Bioneer Corp. (Daejeon, Korea). TRI reagent was obtained from Solgent Co., Ltd. (Daejeon, Korea). AmpliTaq DNA polymerase and Lipofectamine 2000 were obtained from Roche Diagnostics Corp. (Indianapolis, IN, USA) and Invitrogen (Carlsbad, CA, USA), respectively. G418 was from Gibco-BRL (Gaithersburg, MD, USA). The plasmid, pGL3-basic vector and the Dual-Glo luciferase assay kit were purchased from Promega Corp. (Madison, WI, USA). For western blot analysis, specific antibodies against Ax1, cyclin D1, p21, XIAP and GAPDH, as well as secondary antibodies were obtained from Santa Cruz Biotechnology (Dallas, TX, USA).

Cell culture and establishment of cisplatin and paclitaxel-resistant cells. The A549 and H460 cells were grown in RPMI-1640 medium (Gibco-BRL) containing 10% fetal bovine serum (FBS), 2 mM L-glutamine, 10 U/ml penicillin and 10 g/ml streptomycin at 37°C in 5% CO₂ in a water-saturated atmosphere. The variants of A549 and H460 cells which are resistant to cisplatin (A549/CisR and H460/CisR) or paclitaxel (A549/TR and H460/TR cells) were established by stepwise exposure of the parental cells to escalating concentrations of cisplatin (ranging from 0.5 to 2 mM) and paclitaxel (ranging from 3 to 24 nM), respectively.

Reverse transcription PCR (RT-PCR). Cells (2x10⁵) were seeded in a 60-mm culture dish and grown overnight. They were then treated with the indicated concentrations (0, 5, 10 and 20 μM) of curcumin for 24 h. Total RNA was extracted using TRI reagent and subjected to cDNA synthesis and PCR. The specific primers were as follows: Ax1 sense, 5'-AACCTTCAACTCCTGCCTTCTCG-3' and antisense, 5'-CAGCTTCTCCTTCAGCTCTTCAC-3'; GAPDH sense, 5'-GGAGCCAAAGGGTCATCAT-3' and antisense, 5'-GTGATGGCATGACTGTGGT-3'. The mRNA level of Ax1 was normalized to that of GAPDH.

Promoter activity test. The promoter reporter plasmid, pGL3-Ax1, which contains the Ax1 promoter region ranging from -887 to +7 bp of the transcriptional start site was amplified by PCR and subcloned into the pGL3-basic vector, the luciferase reporter plasmid. The constructed promoter reporter plasmid was co-transfected into cells (3x10⁵ cells in a 60-mm dish) with *Renilla* luciferase vectors, pRL-SV40, as an internal control. Luciferase activity was measured using a Dual-Glo luciferase assay system.

Western blot analysis. Total cell lysates were prepared from the parental or chemoresistant cells treated with the indicated concentrations (0, 5, 10 and 20 μM) of curcumin using lysis buffer [1% Triton X-100, 50 mM Tris (pH 8.0), 150 mM NaCl, 1 mM PMSF, 1 mM Na₃VO₄ and protease inhibitor cocktail]. Untreated cells were used as controls. Protein concentrations were determined using Bio-Rad protein assays. Proteins from the cell lysates (20-40 μg) were separated by 12% SDS-PAGE, and electrotransferred onto nitrocellulose membranes. The membranes were blocked for 30 min at room temperature in Tris-buffered saline with 0.05% Tween-20 (TTBS) containing 5% non-fat dry milk, and then incubated with TTBS containing a primary antibody for 4 h at room temperature. After 3x10 min washes in TTBS, the membranes were incubated with peroxidase-conjugated secondary antibody for 1 h. Following 3 additional 10-min washes with TTBS, the protein bands of interest were visualized using an enhanced chemiluminescence detection system (Amersham™ ECL™ Prime Western Blotting Detection reagent; GE Healthcare, Piscataway, NJ, USA).

Cell viability measurement. To assess cell viability, the number of viable cells was counted using Trypan blue. Briefly, 3x10³ cells were seeded into 60-mm culture dish, grown overnight and then treated with the indicated concentrations (0, 5, 10 and 20 μM) of curcumin for 24 h. After curcumin treatment, cells were harvested and stained with 0.4% Trypan blue solution. Dye-excluding viable cells were counted under the microscope. Cell viability was also expressed as a percentage of the viable cells with respect to untreated control cells.

Clonogenic assay. Cells were seeded into 24-well plates (1x10² cells/well) and treated with the indicated concentrations (0, 5, 10 and 20 μM) of curcumin for 24 h. Curcumin-treated cells were then cultured for the next 7-10 days to form colonies. Colonies of >50 cells were stained with Crystal violet (in 60% methanol; Junsei Chemical Co., Ltd., Tokyo, Japan) and

images were acquired using the RAS-3000 Image Analysis System (FujiFilm, Tokyo, Japan).

Ectopic expression of *Axl*. To ectopically express *Axl*, the recombinant plasmid, pcDNA3-*Axl*, was constructed by cloning the *Axl* cDNA into the *Eco*RI and *Bam*HI sites of the pcDNA3 vector and 2 μ g of purified plasmids were transfected into the A549 cells (3×10^5 cells in a 60 mm dish) using Lipofectamine 2000 (Invitrogen). To establish stable cell lines, which constitutively express *Axl*, the transfected cells were cultured in the presence of 400 μ g/ml of G418. The RPMI-1640 medium containing G418 was refreshed every 3 days. After 3-4 weeks, the *Axl*-expressing cells were enriched and the *Axl* expression in these cells was analyzed by western blot analysis.

Transfection of siRNA. To reduce *Axl* expression, RNA interference-mediated gene silencing was performed. Cells (3×10^5) were seeded in 60-mm culture dishes, grown overnight and then transfected with 50 nM siRNA targeting *Axl* (sense, 5'-AAGAUUUGGAGAdACACACUGA-3' and antisense, 5'-UCAGUGUGUUCUCCAAAUCUU-3'), as previously described (30) or control siRNA. The cells were harvested for 24 and 48 h after transfection and used to evaluate protein expression and cell proliferation, respectively.

Statistical analysis. Data were expressed as the means \pm SD of triplicate samples or at least three independent experiments. To determine statistical significance, the Student's t-test was used with a P-value threshold of <0.05.

Results

Curcumin suppresses expression of *Axl* receptor tyrosine kinase at transcriptional level. We first examined if curcumin alters expression of *Axl* receptor tyrosine kinase (RTK) in the lung cancer A549 and H460 cells. After treatment of cells with 5, 10 and 20 μ M curcumin for 24 h, *Axl* protein level was determined by western blot analysis. The results of western blot analysis showed that *Axl* protein level in curcumin-treated cells was reduced in the dose-dependent manner (Fig. 1A). Additionally, this inhibitory effect of curcumin on *Axl* expression was also time-dependent, since *Axl* protein level was found to gradually decrease, when these cells were treated with 20 μ M curcumin for 6, 12 and 24 h (Fig. 1B).

Downregulation of *Axl* expression by curcumin was further demonstrated by RT-PCR. Consistent with western blot results, *Axl* mRNA levels of A549 and H460 cells were also markedly and dose-dependently diminished by the indicated concentrations of curcumin (Fig. 1C). Moreover, the effect of curcumin on transcription of the *Axl* gene was examined using the *Axl* promoter-luciferase reporter plasmid, pGL3-*Axl*. As shown in Fig. 1D, luciferase activities of A549 cells transfected with pGL3-*Axl* and treated with 5, 10 and 20 μ M curcumin for 6 h were significantly declined, indicating that curcumin inhibits *Axl* expression at the transcriptional level.

Curcumin inhibits activation of *Axl* upon the growth arrest-specific gene 6 stimulation. Binding of growth arrest-specific gene 6 (Gas6), a validated ligand, to *Axl* results in its activation,

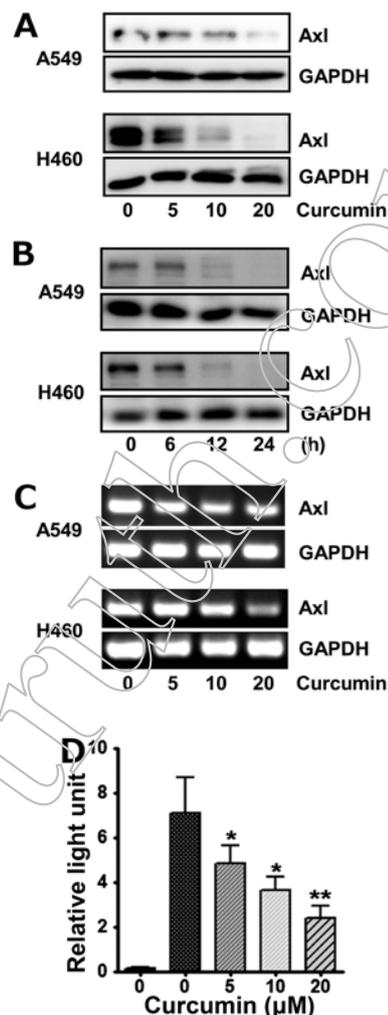


Figure 1. Curcumin decreases *Axl* expression at transcriptional level. Cells (3×10^3 cells/dish) were seeded onto 60-mm dishes, grown overnight and exposed to curcumin. (A) A549 or H460 cells were treated with 5, 10 and 20 μ M curcumin for 24 h and then cells were harvested. (B) A549 or H460 cells were treated with 20 μ M curcumin for 6, 12 and 24 h and then cells were harvested. *Axl* protein levels were determined by western blot analysis. GAPDH was used as a loading control. The result shown is a representative of three independent experiments. (C) For RT-PCR, total RNAs from the cells treated with the indicated concentrations of curcumin for 12 h were isolated and used to determine *Axl* mRNA levels. As an internal control, *GAPDH* mRNA was also amplified by RT-PCR. The data shown are a representative of three independent experiments. (D) To examine the effect of curcumin on *Axl* promoter activity, the H460 cells (3×10^4 cells) were transfected with pGL3 or pGL3 *Axl* using Lipofectamine 2000. The cells were then incubated with 20 μ M curcumin for 24 h and total cell lysates were used to measure luciferase activity. Data are expressed as the means \pm SD of triplicate samples conducted in three independent experiments. The asterisks indicate a significant difference compared to the control value (*P<0.05 and **P<0.01, vs. untreated group).

that is the phosphorylation of tyrosine residues at intracellular kinase domain (31,32). We next examined the effect of curcumin on *Axl* phosphorylation after Gas6 treatment.

Serum-starved A549 and H460 cells were treated with Gas6 for 15, 30, 60 and 120 min and phosphorylated *Axl* levels were determined by western blot analysis. As illustrated in Fig. 2A, *Axl* phosphorylation by Gas6 occurred within 15 min in both cells and returned back to each of their basal levels by 120 min. However, we found that pre-incubation of H460 cells with curcumin inhibited Gas6-induced *Axl* phosphorylation

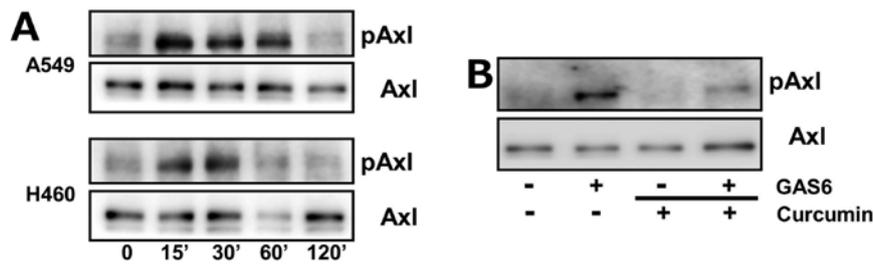


Figure 2. Curcumin inhibits Axl activation upon Gas6 stimulation. Cells (3×10^3 cells/dish) were seeded onto 60-mm dishes, serum-starved overnight, and treated with 250 ng/ml Gas6. (A) A549 or H460 cells were treated with Gas6 for 15, 30, 60 and 120 min and then harvested. (B) H460 cells were pre-incubated with 20 μ M curcumin for 60 min, followed with Gas6 treatment and then harvested. Phosphorylated Axl protein levels were determined by western blot analysis. Total Axl protein level was used as a loading control. The result shown is a representative of three independent experiments.

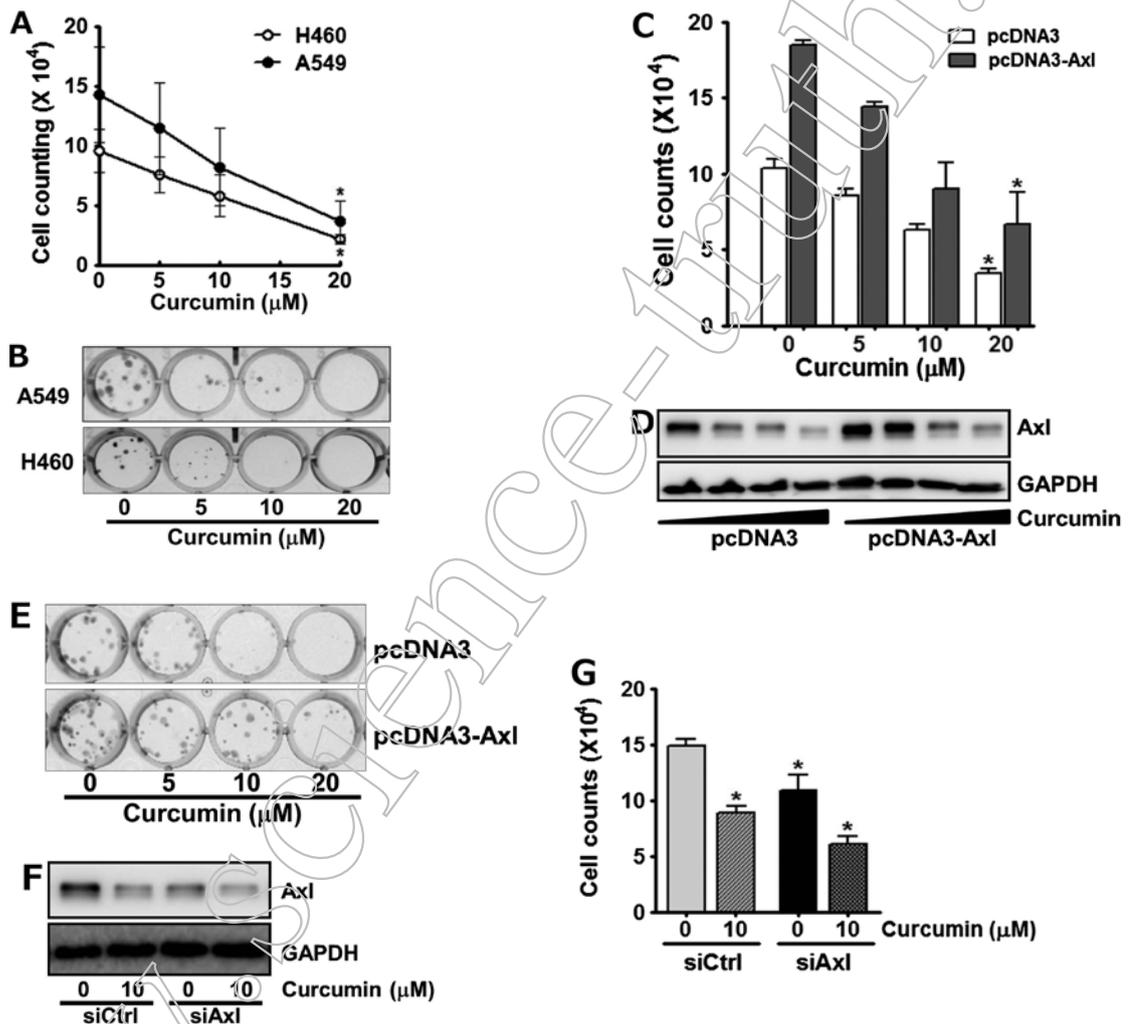
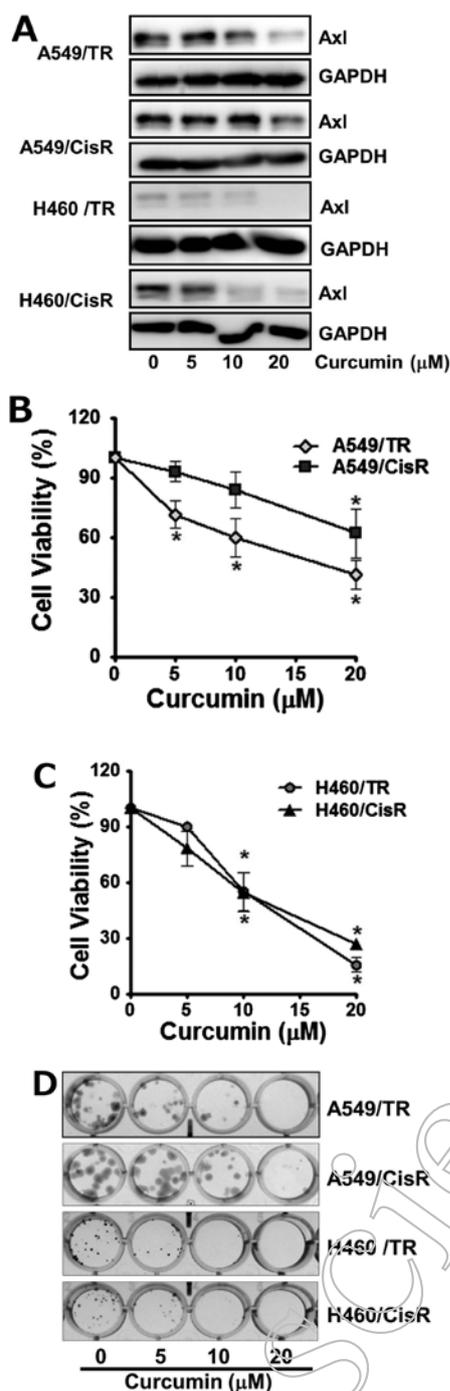


Figure 3. Curcumin inhibits cell proliferation and its anti-proliferative effect is reduced or augmented by Axl protein level. A549 or H460 cells (3×10^3 cells/dish) were seeded onto 60-mm dishes, grown overnight. (A) Cells were treated with 5, 10 and 20 μ M curcumin for 24 h and then harvested. The number of viable cells were counted using trypan blue exclusion assay. Data are represented as mean \pm SD of at least three independent experiments. The asterisks indicate a significant difference compared to the control value ($P < 0.05$ vs. untreated group). (B) Cells (2×10^3 cells/dish) were seeded onto 35-mm dishes, treated with the indicated dose of curcumin for 24 h and allowed to grow for next 7-10 days. Colonies were visualized by crystal violet staining. The data shown are representative of three independent experiments. (C) A549 cells were transfected with pcDNA3 or pcDNA3-Axl plasmid using Lipofectamine 2000. The transfected cells were treated with the indicated doses of curcumin for 24 h, then harvested, and stained with trypan blue to count the viable cells. Data are expressed as the means \pm SD from three independent experiments. The asterisks indicate the significant difference compared to the control value ($P < 0.05$ vs. untreated group). (D) A549/pc-DNA3 and A549/pc-DNA3-Axl cells were treated with the indicated concentrations of curcumin for 24 h. The total cell lysates were prepared and Axl protein level was determined by western blot analysis. (E) Colonogenic assay was also conducted with A549/pc-DNA3 and A549/pc-DNA3-Axl cells exposed to the indicated concentrations of curcumin for 24 h. Colonies formed during 7-10 days of culture were visualized by crystal violet staining. (F) H460 cells (3×10^4 cells) were transfected with Axl specific siRNA or control siRNA, respectively. Cells were harvested 24 h post-transfection, divided into two groups, and grown for 24 h in the presence or absence of 10 μ M curcumin. The total cell lysates were prepared and Axl protein levels were determined by western blot analysis. GAPDH was used as a loading control and results shown are representative of at least three independent experiments. (G) After curcumin treatment, cells were harvested and stained with trypan blue to count viable cells. Data are represented as mean \pm SD of at least three independent experiments. The asterisks indicate a significant difference compared to the control value ($P < 0.05$ vs. untreated group).



(Fig. 2B), suggesting the inhibitory effect of curcumin on Axl activation upon Gas6 stimulation.

Curcumin targets Axl to inhibit cell proliferation. Since overexpression and activation of Axl had been reported to be involved in oncogenesis, cell survival, proliferation and anti-apoptosis (13,14,20,22), we next assessed if the down-regulation of Axl by curcumin affects cell viability. The cells were incubated with 0, 5, 10 and 20 μM of curcumin for 24 h, and the number of viable cells was then counted. As shown in Fig. 3A, treatment of cells with curcumin reduced cell viability in a dose-dependent manner. Of note, following the incubation of A549 and H460 cells with 20 μM curcumin, only 30% and 22% of the cells survived, respectively.

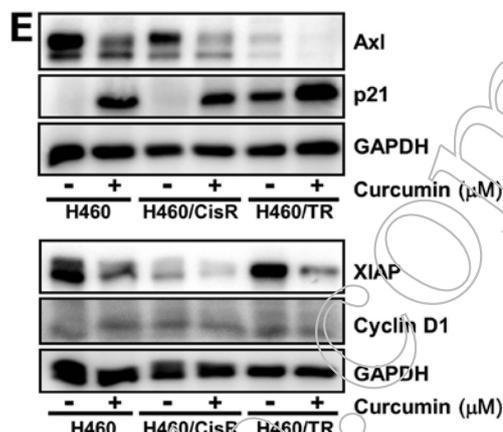


Figure 4. Curcumin inhibits proliferation of cisplatin- and paclitaxel-resistant lung cancer cells and causes induction of p21 as well as reduction of XIAP. Each variants of A549 and H460 cells, which are cisplatin- (A549/CisR and H460/CisR) and paclitaxel-resistant (A549/TR and H460/TR), respectively, were seeded onto 60-mm dishes, and grown overnight. (A) Cells (3×10^3 cells/dish) were treated with 5, 10 and 20 μM curcumin for 24 h and then harvested. Axl protein levels were examined by western blot analysis. GAPDH was used as a loading control. The result shown is a representative of three independent experiments. (B and C) Cells (1×10^3 cells/dish) were seeded onto 96-well plates and treated with the indicated concentrations of curcumin for 24 h. Cell proliferation was determined by CCK-8 assay. The data shown are representative of at least three independent experiments. Data are represented as mean \pm SD. The asterisks indicate a significant difference compared to the control value ($P < 0.05$ vs. untreated group). (D) Cells (2×10^3 cells/dish) were seeded onto 35-mm dishes and treated with 5, 10 and 20 μM curcumin for 24 h, and allowed to grow for 7-10 days. The colonies were visualized by crystal violet staining. The data shown are representative of at least three independent experiments. (E) H460, H460/CisR and H460/TR cells (3×10^3 cells/dish) were treated with 10 μM curcumin for 24 h and then harvested. The levels of p21, cyclin D1 and XIAP protein were determined by western blot analysis. GAPDH was used as a loading control. The result shown is a representative of three independent experiments.

The anti-proliferative effects of curcumin on the A549 and H460 cells were also observed by clonogenic assay. The cells were exposed to the indicated concentrations of curcumin for 24 h and then allowed to grow for the next 10 days. Curcumin treatment was found to result in the dose-dependent decline of colony formation (Fig. 3B). Specifically, H460 cells as well as A549 cells failed to form visible colonies at 10 and 20 μM curcumin, respectively, indicating that curcumin seems to be more cytotoxic toward H460 cells than A549 cells.

To further confirm the involvement of Axl in the antiproliferative effects of curcumin, we examined the cytotoxic effect of curcumin on cells manipulated to enhance or reduce Axl expression. As shown in Fig. 3C, A549 cells transfected with

pcDNA3-Axl, a recombinant plasmid containing Axl cDNA for its overexpression, were less sensitive to curcumin treatment compared to the control cells transfected with pcDNA3 vector, indicating that Axl overexpression reduced the anti-proliferative effects of curcumin. Western blot analysis consistently showed that the Axl level of the pcDNA3-Axl transfected A549 cells was higher than that of their control cells even after curcumin treatment (Fig. 3D). Colony formation assay also manifested that in contrast to the control cells, Axl-overexpressing A549 cells formed more colonies and were relatively less affected by curcumin treatment (Fig. 3E). Subsequently, H460 cells were transfected with Axl specific siRNA, siAxl, or control siRNA, siCtrl and then treated with curcumin for 24 h. We found that Axl targeting by siAxl significantly decreased Axl expression (Fig. 3F), which resulted in the augmentation of anti-proliferative effect of curcumin (Fig. 3G). Taken together, these results demonstrated that Axl protein level tightly correlates with cell proliferation and verified that curcumin inhibits cell proliferation via down-regulation of Axl expression.

Curcumin suppresses proliferation of both cisplatin- and paclitaxel-resistant lung cancer cells and results in the elevation of p21 as well as reduction of XIAP expression. Next, we asked if curcumin could be cytotoxic in the cisplatin- and paclitaxel-resistant NSCLC cells which are the variants of A549 and H460 cells. Each of the variants was established by stepwise exposure of parental cells to increasing concentrations of cisplatin (A549/CisR and H460/CisR cells) or paclitaxel (A549/TR and H460/TR cells), respectively. As shown in Fig. 4A, curcumin reduced the Axl protein levels of A549/TR, A549/CisR, H460/TR and H460/CisR cells in a dose-dependent manner. Notably, in H460/TR cells, the Axl protein level was found to be fairly low and still decreased by curcumin treatment.

In accordance with the western blot results, the viability of both paclitaxel- and cisplatin-resistant cells were dose-dependently declined by curcumin treatment. Especially, exposure of cells with 20 μ M curcumin for 24 h was found to result in only 41.3% (A549/TR), 62.8% (A549/CisR), 15.6% (H460/TR), 27% (H460/CisR) survival of these cells, respectively (Fig. 4B and C). Colonogenic activity of curcumin-treated the variants of A549 and H460 cells further showed cytotoxicity of curcumin on these chemoresistant cells. As shown in Fig. 4D, treatment of these cells with curcumin reduced the number of colonies as well as the size of each colony. In contrast to chemoresistant A549 cells, H460/TR and H460/CisR cells were more profoundly affected by curcumin treatment, which are consistent with the result from cell viability measurement.

To demonstrate the intracellular effectors which are involved in the curcumin-mediated downregulation of Axl expression and result in the inhibition of cell proliferation, we assessed the levels of cell cycle regulator, p21 and apoptosis related protein, the X-linked inhibitor of apoptosis protein (XIAP). The H460, H460/TR and H460/CisR cells were treated with 10 μ M curcumin for 24 h and western blot analysis showed that curcumin induced the expression of the cyclin dependent kinase inhibitor p21, which causes cell cycle arrest, but reduced that of XIAP, which inhibits apoptosis

(Fig. 4E). Collectively, these results indicate that curcumin downregulates Axl expression, subsequently increases p21 protein level and decreases XIAP protein level, which result in the inhibition of cell proliferation.

Discussion

Standard chemotherapy for NSCLC has been the combination of platinum-based agents (cisplatin or carboplatin) and a second drug (pemetrexed, gemtastine, paclitaxel or vinorelbine), but low response rate (20-35%) and eventual development of chemoresistance among initial responders have been the main causes of poor prognosis (33,34). Linger *et al* (24) have demonstrated that Mer or Axl inhibition enhanced sensitivity of NSCLC cells to various cytotoxic agents such as cisplatin, carboplatin, doxorubicin or etoposide by promoting apoptosis. Since TAM family members of RTKs have been reported to play important roles in cell survival, proliferation and apoptosis (35), targeting of these RTKs seems to be a potent strategy to improve standard chemotherapy regimens.

In the present study, we observed that curcumin had inhibitory effects on Axl expression, Gas6-dependent Axl phosphorylation, and Axl promoter activity in NSCLC cells (Figs. 1 and 2). Silencing of Axl expression by RNA interference or specific monoclonal antibodies against Axl have been demonstrated to inhibit cell proliferation, metastasis and xenograft tumor growth in NSCLC (36,37). Consistent with previous reports, we also observed that curcumin decreased the viability of NSCLC cells (A549 and H460). Moreover, anti-proliferative effect of curcumin was reduced by ectopic expression of Axl and augmented by Axl knockdown using siRNA, respectively, suggesting that curcumin abrogates these A549 and H460 cell proliferation via downregulation of Axl expression and further confirming that Axl is a new target of curcumin that contributes to its anticancer effects which have been known to be due to negative regulation of diverse intracellular molecules including transcription factors (38), growth factors, protein kinases (39) and oncogenic proteins (40), resulting in cell cycle arrest and/or apoptosis.

Several reports have shown that curcumin could affect each stage of cancer such as initiation, promotion and progression, ingestion of curcumin was even found to significantly inhibit the activity of lymphocytic glutathione S-transferase, a phase II detoxification enzyme, involved in the development of chemoresistance (41,42). The growing body of evidence also indicates that overexpression and/or activation of Axl is a novel mechanism to induce the acquired resistance to various anticancer drugs including cytotoxic agents and various tyrosine kinase inhibitors, especially EGF receptor inhibitors (gefitinib or erlotinib) (43,44). Consistently, our data also showed that the viabilities of cisplatin/taxol-resistant cells (A549/CisR, H460/CisR, A549/TR and H460/TR) were decreased by curcumin treatment (Figs. 3A and B and 4B-D) and Axl protein levels of each cell type were also declined by curcumin, implying again that Axl plays a critical role in proliferation of both parental and chemoresistant NSCLC cells and the acquisition of resistance against chemotherapeutic drugs.

In summary, our data indicate that curcumin has inhibitory effects on Axl expression and the activation in response to

Gas6 binding, which are associated with its anti-proliferative activity in parental as well as each type of cisplatin/paclitaxel-resistant NSCLC cells. Thus, Axl seems to be a potent therapeutic target of curcumin to inhibit cell proliferation and to overcome chemoresistance of NSCLC cells.

Acknowledgements

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