

Axl receptor tyrosine kinase is a novel target of apigenin for the inhibition of cell proliferation

KYUNG-CHAN $\mathrm{KIM}^1, \, \mathrm{EUN}\text{-}\mathrm{HA}\,\mathrm{CHOI}^2\,$ and $\,\mathrm{CHUHEE}\,\mathrm{LEE}^2$

¹Department of Internal Medicine, College of Medicine, Catholic University of Daegu, Daegu 705-718; ²Department of Biochemistry and Molecular Biology, School of Medicine, Yeungnam University, Daegu 705-717, Republic of Korea

Received February 6, 2014; Accepted June 3, 2014

DOI: 10.3892/ijmm.2014.1804

Abstract. The Axl receptor tyrosine kinase (RTK), along with Tyro 3 and Mer, belongs to the TAM subfamily that promotes survival, stimulates proliferation and/or inhibits apoptosis. In various types of human cancer, including breast, lung and prostate cancer, Axl expression is increased and correlates with an advanced clinical stage. In this study, we examined whether apigenin has an effect on Axl expression, which in turn can affect cell proliferation. The treatment of the non-small cell lung cancer (NSCLC) cells, A549 and H460, with apigenin decreased Axl mRNA and protein expression in a dose-dependent manner. Axl promoter activity was also inhibited by apigenin, indicating that apigenin suppressed Axl expression at the transcriptional level. Upon treatment with apigenin, the viability of both the A549 and H460 cells was gradually decreased and the anti-proliferative effects were further confirmed by the dose-dependent decrease in the clonogenic ability of the apigenin-treated cells. Subsequently, we found that the viability and clonogenic ability of the cells treated with apigenin was less or more affected by transfection of the cells with a Axl-expressing plasmid or Axl targeting siRNA, compared to transfection with the empty vector or control siRNA, respectively. In addition, apigenin increased the expression of p21, a cyclin-dependent kinase inhibitor, but reduced the expression of X-linked inhibitor of apoptosis protein (XIAP). These cell cycle arrest and pro-apoptotic effects of apigenin were also attenuated or augmented by the up- or downregulation of Axl expression, respectively, which suggests that Axl is a novel target of apigenin through which it exerts its inhibitory effects on cell proliferation. Taken together, our data indicate that apigenin downregulates Axl expression, which subsequently results in the inhibition of NSCLC cell proliferation through the increase and decrease of p21 and XIAP expression, respectively.

Introduction

Apigenin (4',5,7,-tirhydoxyflavone) is a dietary flavone that is naturally found as a dimer, diapigenin (1), apigenin-7-O-glucoside or acetylated derivative (2). It is abundantly present in various fruits, plants, vegetables and some herbs (3-6). Plantderived beverages, including herbal tea and red wine, are good sources of apigenin (7). Apigenin has received significant attention as a preventive, as well as a therapeutic agent due to its efficacy and low intrinsic toxicity.

The effects of apigenin on growth inhibition, cell cycle arrest and the induction of apoptosis have been demonstrated in various types of cancer, such as breast cancer with high levels of HER2/neu (8), as well as cervical (8), lung (9), colon (10), hematologic (11) and ovarian cancer (12). In mouse models of prostate cancer, orally delivered apigenin has been shown to reduce tumor volume and suppress metastasis (13). The antiangiogenic property of apigenin has also been reported in lung (14), ovarian (15) and prostate cancer (16), which is the result of the transcriptional repression of vascular endonthe-lial growth factor expression through the downregulation of hypoxia-inducible factor (HIF)1- α expression levels.

In the human genome, 58 receptor tyrosine kinases (RTKs) are known to date; these are catagorized into 20 subfamilies and integrated into one large RTK family. Axl RTK, also known as Ark or Ufo, belongs to the TAM subfamily, which also contains Tyro 3 and Mer. It was initially identified in 1988 as an unidentified transforming gene (17) from chronic myelogenous leukemia patients and cloned in 1991 from primary human leukemia cells (18).

Axl has been shown to be overexpressed in various types of cancer, including acute leukemia (19), as well as breast (20), colon (21), thyroid (22), esophageal (23) and lung cancer (24). Enhanced Axl expression and activation through binding with its ligand, growth arrest-specific 6 (a vitamin K-dependent protein), has been shown to promote downstream signaling for cell proliferation and survival (25-27). Recently, Axl activation has also been identified as a novel mechanism that renders nonsmall cell lung cancer (NSCLC) cells and HER2/neu-positive breast cancer cells resistant to epidermal growth factor (EGF)

Correspondence to: Dr Chuhee Lee, Department of Biochemistry and Molecular Biology, School of Medicine, Yeungnam University, 317-1 Daemyung-Dong, Daegu 705-717, Republic of Korea E-mail: chlee2@ynu.ac.kr

Key words: apigenin, Axl, lung cancer, p21, X-linked inhibitor of apoptosis

RTK inhibitor (TKI) (28) and dual TKIs, such as lapatinib (29), respectively. Therefore, the inhibition of Axl expression and Axl-mediated signaling may be a potential therapeutic target for cancer treatment.

In this study, we examined the effects of apigenin on Axl expression and the subsequent effects on cell proliferation in human lung cancer cells. Apigenin was found to downregulate Axl expression, which resulted in the inhibition of cell proliferation through the induction of p21 protein expression and the suppression of the expression of X-linked inhibitor of apoptosis protein (XIAP).

Materials and methods

Reagents and antibodies. A549 and H460 cells were purchased from the American Type Culture Collection (Manassas, VA, USA). Apigenin was obtained from Sigma (St. Louis, MO, USA). Primers for Axl 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 Axl, p21, XIAP and GAPDH, as well as secondary antibodies were obtained from Santa Cruz Biotechnology (Dallas, TX, USA).

Cell culture. The A549 and H460 cells were grown in RPMI-1640 (Gibco BRL) containing 10% 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.

Reverse transcription PCR (RT-PCR) and quantitative PCR (qPCR). The A549 and H460 cells $(1x10^6)$ were seeded in a 100-mm culture dish and grown overnight. The cells were then treated with the indicated concentrations $(0, 10, 20 \text{ and } 40 \,\mu\text{M})$ of apigenin for 24 h. Total RNA was extracted using TRI reagent and subjected to cDNA synthesis and PCR. The specific primers were as follows: Axl sense, 5'-AACCTTCAACTCC TGCCTTCTCG-3' and antisense, 5'-CAGCTTCTCCTTCAGC TCTTCAC-3'; GAPDH sense, 5'-GGAGCCAAAAGGGTCAT CAT-3' and antisense, 5'-GTGATGGCATGGACTGTGGT-3'. To quantify the Axl mRNA levels in the A549 and H460 cells before and after treatment with apigenin, qPCR was performed using SYBR-Green PCR Master mix and the ABI PRISM 7900HT system (Applied Biosystems, Foster City, CA, USA). The mRNA level of Axl was normalized to that of GAPDH, using the $2^{-\Delta\Delta CT}$ method.

Western blot analysis. Total cell lysates were prepared from the A549 or H460 cells treated with the indicated concentrations (0, 10, 20 and 40 μ M) of apigenin 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 μ g) 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).

Clonogenic assay. The A549 or H460 cells were seeded in 35-mm culture dishes ($2x10^3$ cells/dish) and cultured for the following 7-10 days under the indicated concentrations (0, 10, 20 and 40 μ M) of apigenin to form colonies. Colonies of >50 cells were stained with Crystal violet (in 60% methanol; Junsei Chemical Co., Ltd., Tokyo, Japan) and images were acqired using the RAS-3000 Image Analysis System (FujiFilm, Tokyo, Japan).

Cell viability assay. The Cell Counting kit-8 (Dojindo Laboratories, Kumamoto, Japan) was used to measure the viability of the cells. Briefly, $1x10^3$ cells were seeded in each well of 96-well plates and grown overnight at 37°C and then treated with the indicated concentrations (0, 10, 20 and 40 μ M) of apigenin for 24 h. At the end of treatment, 10 μ l of CCK-8 solution were added followed by incubation for a further 4 h. The absorbance at 450 nm was measured using a microplate reader (Model 680 microplate reader; Bio-Rad Laboratories, Inc., Hercules, CA, USA). Values are the means \pm SD for triplicate wells and were normalized to those of the control group to determine the percentage viability.

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 or H460 cells (3x10⁵ cells in a 100-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.

Promoter activity test. To construct pGL3-AXL, the *Axl* 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 100 mm dish) with renilla luciferase vectors, pRL-SV40, as an internal control. Luciferase activity was measured using a Dual-Glo luciferase assay system.

siRNA trasnfection. To inhibit Axl expression, RNA interference (RNAi)-induced gene silencing was performed. The H460 cells (1x10⁶) were seeded in a 100-mm culture dish, grown overnight and then transfected with 50 nM siRNA targeting Axl(sense,5'-AAGAUUUGGAGAACACACUGA-3';





Figure 1. Apigenin reduces Axl expression in human lung cancer cells. The A549 and H460 cells were treated with the indicated concentrations of apigenin for 24 h. (A and B) The protein level of Axl was determined by western blot analysis in order to assess the effects of apigenin on its expression. GAPDH was used as a loading control. Results are from 3 independent experiments. P <0.05, apigenin-treated vs. untreated cells; ** P<0.001, apigenin-treated vs. untreated cells. (C) For RT-PCR, total RNA from the cells was isolated and used for the analysis of Axl mRNA expression. The level of Axl mRNA was normalized to that of GAPDH. The data shown are representative of 3 independent experiments. (D) qPCR (SYBR-Green) was performed to measure Axl mRNA expression in the A549 and H460 cells treated with apigenin for 24 h. Each sample was performed in triplicate, and the Axl mRNA expression was calculated relative to GAPDH. The asterisks indicate the significant difference compared to the control value (P <0.05, apigenin-treated vs. untreated cells). (E) To determine *Axl* promoter activity, the A549 cells (3x10³ cells/dish) were transfected with pGL3 or pGL3-Axl, *Axl* promoter-luciferase plasmid, using Lipofectamine 2000. The cells were then incubated with apigenin and total cell lysates were used to measure luciferase activity. The data shown are representative of at least 3 independent experiments. Data are expressed as the means \pm SD of triplicate samples conducted in 3 independent experiments. The asterisks indicate the significant difference compared to the control value (P <0.05, pGL3-Axl/apigenin and total cell lysates were used to measure luciferase activity. The data shown are representative of at least 3 independent experiments. Data are expressed as the means \pm SD of triplicate samples conducted in 3 independent experiments. The asterisks indicate the significant difference compared to the control value (P <0.05, pGL3-Axl/apigenin vs. pGL3; and *P <0.001, pGL3-Axl vs. pGL3).

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, as well as in cell proliferation and colony formation assays.

Statistical analysis. Data are expressed as the means \pm SD of triplicate samples or at least 3 independent experiments. To determine statistical significance, the Student's t-test was used with a p-value threshold of <0.05.

Results

Apigenin suppresses Axl expression in human lung cancer cells. To determine the effects of apigenin on Axl expression, the human lung cancer cells, A549 and H460, were incubated with 10, 20, or 40 μ M of apigenin for 24 h and subsequently, Axl protein expression was measured. As shown in Fig. 1A and B, the results from western blot analysis revealed that treatment with apigenin induced a dose-dependent decrease in the protein expression of Axl in both cell lines. More specifically, when the cells were exposed to 40 μ M apigenin for 24 h, the Axl protein levels in the A549 and H460 cells were diminished to 57 and 35% compared with the untreated cells, respectively.

The downregulation of Axl expression in the apigenintreated cells was further confirmed by RT-PCR and a promoter activity test. The results from RT-PCR revealed that in both cells lines, the mRNA levels of Axl were decreased following treatment with apigenin (Fig. 1C); these results were consistent with those from western blot analysis. Additionally, the results from qPCR revealed that the Axl mRNA levels in the A549 and H460 cells treated with 40 μ M apigenin for 24 h were reduced to 37 and 48% compared with the untreated cells, respectively (Fig. 1D).

To determine the effects of apigenin on the transcription of the Axl gene, Axl promoter activity was measured using luciferase reporter plasmid under the control of the human Axl promoter plasmid, pGL3-Axl. A549 cells were transfected with pGL3-Axl and then incubated with 40 μ M apigenin for 24 h. As illustrated in Fig. 1E, luciferase activity significantly declined following treatment with apigenin. The results of RT-PCR as well as those from the promoter activity test indicated that apigenin suppressed Axl expression in the lung cancer cells at the transcriptional level.

Apigenin-mediated downregulation of Axl is responsible for its anti-proliferative effects on human lung cancer cells. Since Axl has been known to transduce cell survival, growth and proliferation (19,20,25,27,31), we wished to determine whether the downregulation of Axl by apigenin affects lung cancer cell viability. The cells were incubated with 10, 20, 40 μ M of apigenin for 24 h, and the number of viable cells was then counted. As shown in Fig. 2A, treatment with apigenin reduced cell viability in a dose-dependent manner. Following



Figure 2. The apigenin-mediated downregulation of Axl is responsible for its anti-proliferative effects on human lung cancer cells. (A) Both A549 and H460 cells ($1x10^3$ cells/well) were seeded into 96-well plates, and the indicated concentrations of apigenin were then added to each well followed by incubation for 24 h. Cell proliferation was determined by CCK-8 assay. Data are expressed as the means \pm SD from 3 independent experiments. Data are expressed as the means \pm SD of triplicate samples conducted in 3 independent experiments. The asterisks indicate the significant difference compared to the control value (*P<0.05, 10 or 20 μ M apigenin-treated H460 cells vs. untreated H460 cells; and **P<0.001, 40 μ M apigenin-treated cells vs. untreated cells). (B) Cells ($2x10^3$ cells) were seeded into 35-mm dishes and allowed to grow for 7-10 days until colonies were formed. The colonies were visualized by Crystal violet staining. The data shown are representative of at least 3 independent experiments. (C) A549 cells were transfected with pcDNA3 or pcDNA3-Axl plasmid and then incubated with the indicated concentrations of apigenin for 24 h. Cell proliferation was determined by CCK-8 assay. Data are expressed as the means \pm SD from 3 independent experiments. The asterisks indicate the significant difference compared to the control value (*P<0.05, 40 μ M apigenin-treated pC-DNA3-Axl cells vs. untreated pC-DNA3-Axl cells; and **P<0.001, 40 μ M apigenin-treated pC-DNA3 cells vs. untreated pC-DNA3-Axl cells vs. untreated pC-DNA3-Axl cells; and **P<0.001, 40 μ M apigenin-treated pC-DNA3 cells vs. untreated pC-DNA3 cells vs. untreated pC-DNA3-Axl cells were prepared and the Axl protein level was determined by western blot analysis. (E) Colony formation assay was performed with the A549/Axl and H460/Axl cells in the presence or absence of apigenin. Colonies formed during 10 days of culture were visualized by Crystal violet staining.

treatment of the A549 and H460 cells with 40 μ M apigenin, only 49 and 37% of the cells survived, respectively.

The anti-proliferative effects of apigenin on the lung cancer cells were further confirmed by colony formation assay. The cells were cultured for 10 days in the presence of 10, 20, 40 μ M of apigenin. The exposure of the cells to apigenin resulted in a dose-dependent inhibition of colony formation (Fig. 2B). More specifically, the H460 cells failed to grow into a colony under 40 μ M apigenin, demonstrating the cytotoxic effects of apigenin on lung cancer cells.

To verify the involvement of Axl in the apigenin-mediated inhibition of cell proliferation, we examined whether the overexpression of Axl has an impact on the anti-proliferative effects of apigenin. The Axl-expressing construct, pcDNA3-Axl, was transiently transfected into the A549 cells and the cells were then cultured with or without apigenin for 24 h. Compared to the control cells transfected with a pcDNA3 empty vector, the cells transfected with the pcDNA3-Axl plasmid were found to be slightly less sensitive to apigenin treatment (Fig. 2C), indicating that the overexpression of Axl protein diminisehd







Figure 4. The apigenin-induced downregulation of Axl expression facilitates the induction of p21 and the reduction of XIAP expression. (A) A549 and H460 cells(3x10³ cells) were treated with apigenin for 24 h. Following treatment, total cell lysates were prepared, and western blot analysis was conducted to determine the p21 and XIAP protein levels. GAPDH was also detected as a loading control. The data shown are representative of 3 independent experiments. (B) H460 cells (3x10³ cells) were transfected with siCtrl or siAxl. Subsequently, 24 h post-transfection, the cells were treated with apigenin for 24 h. Following treatment, total cell lysates were prepared and western blot analysis was conducted to determine the p21 and XIAP protein levels. GAPDH was also detected as a loading control.

Figure 3. Inhibition of Axl augments the anti-proliferative effects of apigenin on human lung cancer cells. H460 cells (3x10³ cells) were transfected with control siRNA (siCtrl) or Axl-specific siRNA (siAxl). (A) Cells were harvested 48 h post-transfection, total cell lysates were prepared, and the Axl protein level was determined by western blot analysis. (B) H460, H460/siCtrl and H460/siAxl cells (1x10³ cells/well) were plated onto 96-well plates and the indicated concentrations of apigenin were added to each well followed by incubation for 24 h. Cell proliferation was determined by CCK-8 assay. Data are expressed as the means \pm SD from 3 independent experiments. The asterisks indicate the significant difference compared to the control value (^{*}P<0.05, 40 μ M apigenin-treated cells vs. untreated cells) (C) H460/siCtrl and H460/siAxl cells (2x10³ cells) were grown for 7-10 days until they formed colonies. The colonies were visualized by Crystal violet staining.

the anti-proliferative effects of apigenin in the cells transfected with the pcDNA3-Axl plasmid.

Subsequently, we established Axl-overexpressing stable cell lines, A549/Axl and H460/Axl, and observed the effects of apigenin on cell proliferation. Western blot analysis revealed that the Axl protein level in the Axl-overexpressing cells was higher than in their parental cells, even following treatment with apigenin (Fig. 2D). Colony formation assay also revealed that both the A549/Axl and H460/Axl cells formed more colonies and were less affected by treatment with apigenin (Fig. 2E); these results are consistent with those from western blot analysis.

To further demonstrate the role of Axl in the anti-proliferative effects of apigenin, its expression was inhibited by specific siRNA. In contrast to the control siRNA, siCtrl, the Axl targetingsiRNA, siAxl, induced a significant decrease in Axl expression (Fig. 3A), resulting in a reduction in cell proliferation (Fig. 3B), as well as in colony formation (Fig. 3C). Taken together, these results suggest that Axl protein expression correlates with cell proliferation and that the anti-proliferative effects of apigenin are mediated by the modulation of Axl expression.

Apigenin-induced downregulation of Axl expression facilitates the induction of p21 and reduction of XIAP expression. To elucidate the molecular mechanisms by which the apigeninmediated downregulation of Axl expression results in the inhibition of cell proliferation, we examined the effects of apigenin on several molecules related to cell cycle regulation and apoptosis. The A549 and H460 cells were treated with $40 \,\mu\text{M}$ apigenin for 24 h. The results from western bolt analysis revealed that treatment with apigenin increased the levels of the cyclin-dependent kinase inhibitor, p21, which induces cell cycle arrest, but decreased the levels of XIAP, which inhibits apoptosis (Fig. 4A and B). These effects (the induction of p21 and the decrease in XIAP expressino) upon treatment with apigenin were attenuated or augmented in the Axl-overexpressing A549 and H460 cells (Fig. 4A) or in the siAxl-transfected H460 cells (Fig. 4B), respectively. These data indicate that the apigenin-induced downregulation of Axl expression is a prerequisite for the subsequent increase or reduction in p21 and XIAP expression.

Discussion

The first observation by Birt et al demonstrated that apigenin is an anti-mutagenic and anti-promotion bioflavonoid based on its inhibitory effects on ornithine decarboxylase, which plays an important role in tumor promotion (32). Subsequently, a number of other studies further confirmed its antioxidant, anti-inflammatory, anti-angiogenic and anti-proliferative activities, which revealed various targets of apigenin simultaneously. For instance, apigenin has been shown to suppress lipopolysaccharide (LPS)-induced cyclooxygenase-2 and nitric oxide synthase-2 expression (33) and TNF- α induced nuclear factor- κ B activation (34), which are major mediators for eliciting inflammation. In addition, apigenin has been verified as an effective inhibitor of the p34 (cdc2) kinase, which increases p53 stability (35), and a good inducer of the p21 and Apaf-1 proteins, which are the main players for cell cycle arrest and the induction of apoptosis (36,37). Based on these data, apigenin

has evoked particular interest in its potential as a chemopreventive and chemotherapeutic agent.

In this study, we found a new target of apigenin, Axl RTK, which is a member of the TAM RTK subfamily and has been reported to be overexpressed in a various types of cancer and to have an oncogenic potential (18,23,24). The exposure of NSCLC cells (A549 and H460) to apigenin resulted in a decrease in Axl mRNA and protein expression (Fig. 1A-D). Furthermore, *Axl* promoter activity was also reduced following treatment with apigenin (Fig. 1E), indicating that apigenin suppresses Axl expression at the transcriptional level.

Specific targeting of Axl with RNAi or monoclonal antibodies, which causes the downregulation of Axl expression, has been reported to reduce the proliferation of NSCLC cells in vitro and in vivo (using tumor xenografts) (38,39). In addition, the overexpression and/or activation of Axl has been shown to be a mechanism that confers acquired resistance to various chemotherapeutic drugs, including gefitinib or erlotinib that are EGF receptor (EGFR) inhibitors in NSCLC (40-42) and head and neck cancer (31), imatinib that is a TKI in chronic myelogenous leukemia (43) and gastrointestinal stromal tumors (44), and tumor necrosis factor-related apoptosisinducing ligand in esophageal adenocarcinoma (45). Therefore, we examined whether the inhibition of Axl expression by apigenin affects the proliferation of NSCLC cells. In agreement with previous studies, we also found that treatment with apigenin inhibited the proliferation and clonogenic ability of the A549 and H460 cells in a dose-dependent manner (Fig. 2A and B). Furthermore, the anti-proliferative effects of apigenin were decreased or augmented by the induction (Fig. 2C and E) or inhibition of Axl expression (Fig. 3B), respectively. We also found that the ectopic expression of Axl diminished the apigenin-induced increase in p21 protein expression and the reduction in XIAP expression (Fig. 4A), which facilitates cell cycle arrest and apoptosis, respectively. By contrast, the inhibition of Axl expression using specific siRNA was found to augment both the induction of p21 and the reduction of XIAP upon treatment with apigenin (Fig. 4B). These results suggest that the Axl protein level correlates with cell proliferation, and the anti-proliferative effects of apigenin are mediated by the modulation of Axl expression.

In conclusion, our observations indicate that the treatment of NSCLC cells with apigenin suppresses Axl expression at the transcriptional level, which subsequently induces the inhibition of cell proliferation through the induction of cell cycle arrest and/or apoptosis. These results suggest that Axl is a novel target of apigenin through which it exerts its anti-proliferative effects on cancer cells.

Acknowledgements

This study was supported by the Basic Science Research Program through the, National Research Foundation of Korea (NRF) funded by the Ministry of Education, Science and Technology (grant no. 2006-2005303).

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