

The enhanced tumor inhibitory effects of gefitinib and L-ascorbic acid combination therapy in non-small cell lung cancer cells

KYOUNG EUN LEE^{1*}, EUNSIL HAHM^{2*}, SEYEON BAE², JAE SEUNG KANG² and WANG JAE LEE²

¹Division of Hematology-Oncology, Department of Internal Medicine, School of Medicine, Ewha Womans University, Seoul 07985; ²Department of Anatomy and Tumor Immunity Medical Research Center, College of Medicine, Seoul National University, Seoul 03080, Republic of Korea

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Abstract. Despite documentation of successful therapy with epidermal growth factor receptor (EGFR) tyrosine kinase inhibitors in patients with lung cancer, the response rate of patients treated with this therapy remains low. The present study investigated whether L-ascorbic acid serves an adjuvant role *in vitro* when combined with the EGFR tyrosine kinase inhibitor gefitinib (Iressa®) in lung cancer cell lines. A total of three human lung cancer cell lines were used. The antiproliferative effects and changes in the cell cycle and expression of intracellular signaling molecules, including extracellular signal-regulated kinases (Erk), signal transducer and activator of transcription 3 (Stat3) and protein kinase B (Akt), were measured in cells treated with gefitinib and/or L-ascorbic acid at various concentrations. When combined with gefitinib, L-ascorbic acid exhibited an additive effect on cell proliferation in all gefitinib-sensitive and gefitinib-resistant cell lines. A decrement of ~40% was observed with a low dose 0.5 mM L-ascorbic acid and gefitinib in the relatively gefitinib-resistant A549 cell line (85.6±5.4% with gefitinib alone vs. 52.7±7.3% with combination therapy; P=0.046). The downregulation of intracellular signaling cascades, including EGFR, Akt, Erk and Stat3, was also observed. L-Ascorbic acid serves an adjuvant role when administered in combination with gefitinib; however, the degree of inhibition of cell proliferation differs between lung cancer cell lines.

Introduction

Lung cancer is the leading cause of cancer-associated mortality worldwide (1,2). A total of one-third of patients with lung cancer initially present at an advanced stage, when only palliative therapies are typically available (1,2). Although complete resection of the lung is the only known curative therapy in stages I-III of this disease, a number of patients do experience relapse following surgery (3). A total of one-third of patients with lung cancer survive for a number of years following diagnosis (3). Recently, there has been a paradigm shift in lung cancer therapeutics, from the use of conventional cytotoxic drugs to the use of variable molecular-targeted therapeutics, including gefitinib, erlotinib and crizotinib (4).

The epidermal growth factor receptor (EGFR) is a member of the transmembrane tyrosine kinase receptor family and its expression is elevated in a number of solid tumors. Initially, the overexpression of EGFR was demonstrated to be a strong prognostic indicator in head and neck, ovarian, cervical, bladder and esophageal cancer (5). Numerous trials aimed at identifying the benefits of administering anti-EGFR therapies to patients with cancer have been conducted (6,7). There are currently two types of therapeutics available for inhibiting EGFR: One is a monoclonal antibody against EGFR and the other includes small-molecule inhibitors of EGFR tyrosine kinase (4). Gefitinib, trade name Iressa®, is a small molecule that inhibits EGFR tyrosine kinase activity, and is the first molecular-targeted agent registered to treat advanced non-small cell lung cancer (NSCLC). Gefitinib trials have demonstrated that there is an overall response rate of 10-20% in patients with advanced NSCLC, which could be as high as 70% in patients with EGFR gene mutations when used as a first-line treatment (8,9). There are two biomarkers, the presence of the EGFR gene mutations and an increase in gene copy number, known to be potent predictors of the response to gefitinib (10,11). The presence of mutations in the EGFR gene is a major determinant of the gefitinib response; these mutations are small deletions affecting amino acids 747-750, or are point mutations (12-15).

There are four obstacles involved in the resistance to gefitinib that have recently been identified (16,17). The first is the activation of alternative tyrosine kinase receptors that bypass the EGFR pathway. The second is the constitutive activation

Correspondence to: Dr Wang Jae Lee, Department of Anatomy and Tumor Immunity Medical Research Center, College of Medicine, Seoul National University, 103 Daehang-ro, Chongno-gu, Seoul 03080, Republic of Korea
E-mail: kinglee@snu.ac.kr

*Contributed equally

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of signaling pathways downstream of the EGFR pathway, including the phosphoinositide 3-kinase/protein kinase B (Akt) pathway or proto-oncogene tyrosine-protein kinase Src family kinases, which induce resistance. The third is the ligand-independent activation of EGFR and the associated resistance to therapy. The fourth is a secondary EGFR mutation which changes the sensitivity to EGFR inhibitors. Efforts to overcome these complications involved in the resistance to gefitinib include the use of tyrosine kinase inhibitors that target other molecules; however, many limitations remain (18).

In the case of other molecular-targeted therapeutics, such as imatinib mesylate, changing the redox status with L-ascorbic acid restores drug sensitivity in imatinib-resistant cell lines. Previous studies have revealed that oxidative stress increases phosphorylation of the EGFR in keratinocytes (19-21). The antioxidant activity of L-ascorbic acid (vitamin C, water-soluble L-ascorbic acid) is achieved primarily via its ability to donate electrons and therefore function as a reducing agent (22). Cell signaling activities and the expression of associated molecules are sensitive to exogenous and intracellular redox status, and L-ascorbic acid serves a role in downregulating the expression of certain signaling molecules, including nuclear factor- κ B, activator protein-1, Fos proto-oncogene and Jun proto-oncogene, and in regulating apoptosis (23).

There is considerable debate about whether L-ascorbic acid serves a therapeutic role in cancer. Recent reports have established that L-ascorbic acid also acts as a pro-oxidant and, depending on the dose, cell type and ability to stimulate apoptosis, may kill cancer cells (24). In human colon cancer cell lines, L-ascorbic acid functions as a potent antioxidant and blocks the chemotherapy-mediated induction of apoptosis (25). These results support the hypothesis that antioxidants may protect cancer cells from the free radical damage induced by chemotherapy. In addition to oxidation, L-ascorbic acid may serve a modulatory role in cellular phosphorylation-dephosphorylation events (20,26-28). However, the mechanisms underlying these effects are currently unclear. Clinical trials of antioxidant plus chemotherapy regimens in patients with cancer are uncommon, due to concerns over inhibiting the effect of chemotherapy via decreasing the production of reactive oxygen species (ROS) (25). However, EGFR tyrosine kinase inhibitors exhibit a different mechanism of action when administered with cytotoxic chemotherapy to kill cancer cells (29).

In the present study, the EGFR tyrosine kinase inhibitor gefitinib was used as an apoptotic stimulus in lung cancer cell lines, and L-ascorbic acid was used as an adjuvant, in order to kill cancer cells. The aims of the present study were to evaluate whether L-ascorbic acid when co-administered with gefitinib therapy serves an additive or synergistic tumor-inhibitory role, and to elucidate the mechanisms underlying its effects in NSCLC cell lines.

Materials and methods

Cell lines. Three human NSCLC cell lines, A549, Calu-3 and HCC827, were obtained from the American Type Culture Collection (Manassas, VA, USA). HCC827 possesses an acquired mutation in the EGFR tyrosine kinase domain

(E746-A750 deletion), and the other two cell lines contain a wild-type EGFR tyrosine kinase domain. The cells were cultured in complete RPMI-1640 (Thermo Fisher Scientific, Inc., Waltham, MA USA) supplemented with 2 mM L-glutamine, 100 U/ml penicillin, 100 μ g/ml streptomycin and 10% heat-inactivated fetal bovine serum (Thermo Fisher Scientific, Inc.), and were used for experiments whilst in the log phase of growth.

Reagents. Gefitinib (ZD1839; trade name IressaTM) was kindly provided by AstraZeneca Pharmaceuticals (Macclesfield, UK) and was dissolved in dimethyl sulfoxide at a stock concentration 10 mM and stored at -20°C for *in vitro* experiments. L-ascorbic acid (sodium salt) was purchased from Sigma-Aldrich (St. Louis, MO, USA). For the flow cytometric analysis, a phycoerythrin-conjugated mouse anti-human EGFR antibody was purchased from BD Pharmingen (San Diego, CA, USA). For the western blot analysis, antibodies against EGFR (cat. no. 4267S; dilution, 1:1,000), phosphorylated (p)-EGFR [Tyr845 (cat. no. 6963S; dilution, 1:1,000), Tyr992 (cat. no. 2235S; dilution, 1:1,000) and Tyr1068 (cat. no. 2234S; dilution, 1:1,000)], Akt (cat. no. 9272S; dilution, 1:1,000), p-Akt (cat. no. 4060S; dilution, 1:1,000), signal transducer and activator of transcription 3 (Stat3; cat. no. 9139S; dilution, 1:1,000) and p-Stat3 (cat. no. 9145S; dilution, 1:1,000) were purchased from Cell Signaling Technology, Inc. (Danvers, MA, USA). Antibodies against extracellular signal-related kinase (Erk; cat. no. sc-94; dilution, 1:1,000), p-Erk (cat. no. sc-7383; dilution, 1:1,000) and β -actin (cat. no. sc-47778; dilution, 1:2,000) were purchased from Santa Cruz Biotechnology, Inc. (Dallas, TX, USA). Trypan blue and 7-aminoactinomycin D were purchased from Sigma-Aldrich (Merck KGaA, Darmstadt, Germany).

Cell proliferation and viability assays. A total of 8×10^5 cells from each cell line were cultured in T25 culture flasks at 37°C in a 5% CO₂ humidified incubator, with or without dissolved L-ascorbic acid and/or gefitinib in PBS at 0, 20 and 40 mM gefitinib in A549 cells, 0.05 and 1.0 mM in Calu-3 cells, 0, 2.5 and 5.0 μ M in HCC827 cells, and 0 and 0.5 mM L-ascorbic acid in A549 cells, 0, 2.5 and 5.0 mM in Calu-3 cells, and 0, 0.5 and 1.0 mM in HCC827 cells. The cells were pretreated with L-ascorbic acid for 1 h and then treated with gefitinib for 48 h at room temperature. The number of cells and the viability of the cells were determined using a Trypan blue dye exclusion assay. An alternative method to monitor cell proliferation, the AlamarBlue[®] assay (cat. no. BUF012A; Bio-Rad Antibodies, Oxford, UK), was also used. AlamarBlue[®] is a redox indicator that is reduced by reactions innate to cellular metabolism (30). Thus, it provides an indirect measure of the number of viable cells. The cells (5×10^3) were seeded onto a 96-well plate and then treated with the aforementioned doses of L-ascorbic acid and/or gefitinib for 48 h at 37°C. AlamarBlue[®] (10% v/v in medium) was subsequently added to the cells, the cells were incubated for 6 h at 37°C and fluorescence was measured at 530 nm excitation and 590 nm emission wavelengths in a spectrofluorometer (Fluoroskan AscentTM FL; Labsystems Diagnostics Oy, Vantaa, Finland). The results are expressed as a percentage relative to the total cell number, and the groups

that were not treated with L-ascorbic acid and gefitinib were used as control group.

Detection of intracellular ROS and cell cycle analysis. Cells from the three lung cancer cell lines were seeded into a 96-well plate at a density of 1×10^4 cells/well and incubated with 50 mM 2',7'-dichlorodihydrofluorescein diacetate (Sigma-Aldrich; Merck KGaA) for 5, 10, 15, 20, 25 and 30 min at 37°C in a 5% CO₂ humidified incubator. The cells were then analyzed using a CytoFluor 2350 plate reader (EMD Millipore, Billerica, MA, USA) with the excitation wavelength set at 485 nm and the emission wavelength at 530 nm. For cell cycle analysis, cells were pretreated with L-ascorbic acid in complete medium for 1 h and then treated with gefitinib for 48 h at 37°C. The cells were then trypsinized, washed twice with cold PBS and centrifuged for 5 min with 200 x g at room temperature. The pellet was resuspended in 1 ml cold PBS and 4 ml cold ethanol for 30 min at 4°C. The cells were centrifuged at 200 x g for 5 min, and the pellet was washed twice with cold PBS, suspended in 500 µl propidium iodide staining solution (BD Pharmingen) and analyzed by flow cytometry BD FACSCalibur (BD Biosciences, Franklin Lakes, NJ, USA).

Western blotting. The cells were lysed and the proteins were extracted in a buffer containing 50 mM Tris-HCl (pH 7.4), 1% NP-40, 0.25% sodium deoxycholate, 150 mM NaCl, 1 mM EDTA, 50 mM NaF, 1 mM sodium orthovanadate, 1 mM phenylmethylsulfonyl fluoride and a protease inhibitor cocktail (Sigma-Aldrich; Merck KGaA). The protein concentration was measured using a Bio-Rad Protein Assay kit (Bio-Rad Laboratories, Inc., Hercules, CA, USA). Equal amounts of protein (20–30 µg) were resolved using 8% (for EGFR and p-EGFR) or 12% [for Akt, p-Akt (Ser473), Erk, p-Erk, Stat3, p-Stat3 and β-actin] SDS-PAGE and transferred onto nitrocellulose membranes. The membranes were blocked with 5% nonfat milk and 0.1% Tween-20-PBS for 1 h at RT with gentle agitation, washed with 0.1% Tween 20-PBS, and then exposed to the relevant primary antibody for 1 h at room temperature. The primary antibodies were diluted 1:1,000 in 0.1% Tween-20-PBS; the antibodies included anti-EGFR (cat. no. 4267S), anti-p-EGFR (Tyr845; cat. no. 6963S), anti-p-EGFR (Tyr992; cat. no. 2235S), anti-p-EGFR (Tyr1068; cat. no. 2234S), anti-Akt (cat. no. 9272S) and anti-p-Akt (Ser473; cat. no. 4060S) from Cell Signaling Technology, Inc.; anti-Erk (cat. no. sc-94) and anti-p-Erk (cat. no. sc-7383) from Santa Cruz Biotechnology, Inc. Subsequent to washing, the blots were exposed to a biotin-conjugated rabbit developed anti-mouse (cat. no. sc-358919; dilution, 1:1,000; Santa Cruz Biotechnology, or mouse developed anti-rabbit antibody (cat. no. sc-2491; dilution, 1:1,000; Santa Cruz Biotechnology,) for 1 h at room temperature. The membranes were then washed, incubated with a 1:5,000 dilution of streptavidin-horseradish peroxidase at room temperature, and the immunoreactive proteins were visualized using an ECL detection system (GE Healthcare Life Sciences, Chalfont, UK).

Statistical analysis. The results are presented as the mean, with error bars representing the standard deviation. Statistical analyses were performed using SPSS v.18 (SPSS, Inc., Chicago,

IL, USA). Non-parametric tests (Mann-Whitney U test) were used to evaluate significant differences among the continuous variables and the Kruskal-Wallis test was used to assess the statistical significance. For multiple comparisons, MedCalc® software for Windows (v.16.1) was used. (MedCalc® Software, Mariakerke, Belgium) P<0.05 was considered to indicate a statistically significant difference.

Results

Increased expression of EGFR is induced by L-ascorbic acid or gefitinib in NSCLC cell lines. Cell surface EGFR protein expression was observed in NSCLC cell lines treated with or without L-ascorbic acid and/or gefitinib. When L-ascorbic acid and gefitinib were administered together, a 91.8% of cells expressed surface EGFR proteins, as compared with 35.3% of untreated Calu-3 cells. Treatment with gefitinib alone increased surface EGFR expression in Calu-3 cells, compared with treatment with L-ascorbic acid alone (88.9 vs. 64.5% of cells; P<0.05). However, co-treatment with gefitinib and L-ascorbic acid resulted in no significant increase in EGFR expression, compared with treatment with gefitinib alone (91.8 vs. 88.9% of cells). In untreated A549 cells, 60.4% of cells expressed surface EGFR protein; this percentage was higher compared with that in untreated Calu-3 cells. The two agents demonstrated a tendency to increase EGFR expression in A549 cells; however, L-ascorbic acid exhibited no synergistic or additional effect on EGFR expression when co-administered with gefitinib. By contrast, L-ascorbic acid and gefitinib administered in combination decreased EGFR expression levels in A549 cells: 63.3% of cells co-treated with the two agents expressed the EGFR protein, compared with 80.5% of cells treated with L-ascorbic acid alone and 72.4% treated with gefitinib alone, although no significance was observed.

Effects of gefitinib and L-ascorbic acid on cancer cell growth and proliferation. A549 is a lung cancer cell line that is resistant to EGFR tyrosine kinase inhibitors. Proliferation of these cells was inhibited with high doses of gefitinib; however, relatively low doses of L-ascorbic acid (0.5 mM) administered with gefitinib inhibited cell proliferation to a greater extent (P=0.046; Fig. 1A). The addition of 0.5 mM L-ascorbic acid to 20 µM gefitinib inhibited cellular proliferation by >32% (85.6 ± 5.4 vs. $52.7 \pm 7.3\%$ of the total number of cells; P=0.046). The inhibitory effect of gefitinib on Calu-3 cells, which exhibit the wild-type EGFR but are sensitive to gefitinib, was ~60% of that observed in the control Calu-3 cells. The additional L-ascorbic acid significantly inhibited the proliferation of Calu-3 tumor cells at each concentration level of gefitinib (overall, P=0.027, no gefitinib; P=0.039, 0.5 µM gefitinib; and P=0.127, 1 µM gefitinib, respectively; Fig. 1B). The additional significant inhibitory effect of the L-ascorbic acid was also observed in gefitinib-sensitive HCC827 cells at each gefitinib concentration (overall, P=0.066, P=0.039 and P=0.027, respectively; Fig. 1C). The AlamarBlue assay used to measure growth inhibition and the Trypan blue dye exclusion assay demonstrated a similar pattern of inhibition (data not presented). Microscopic examination of Calu-3 cells revealed that they experienced

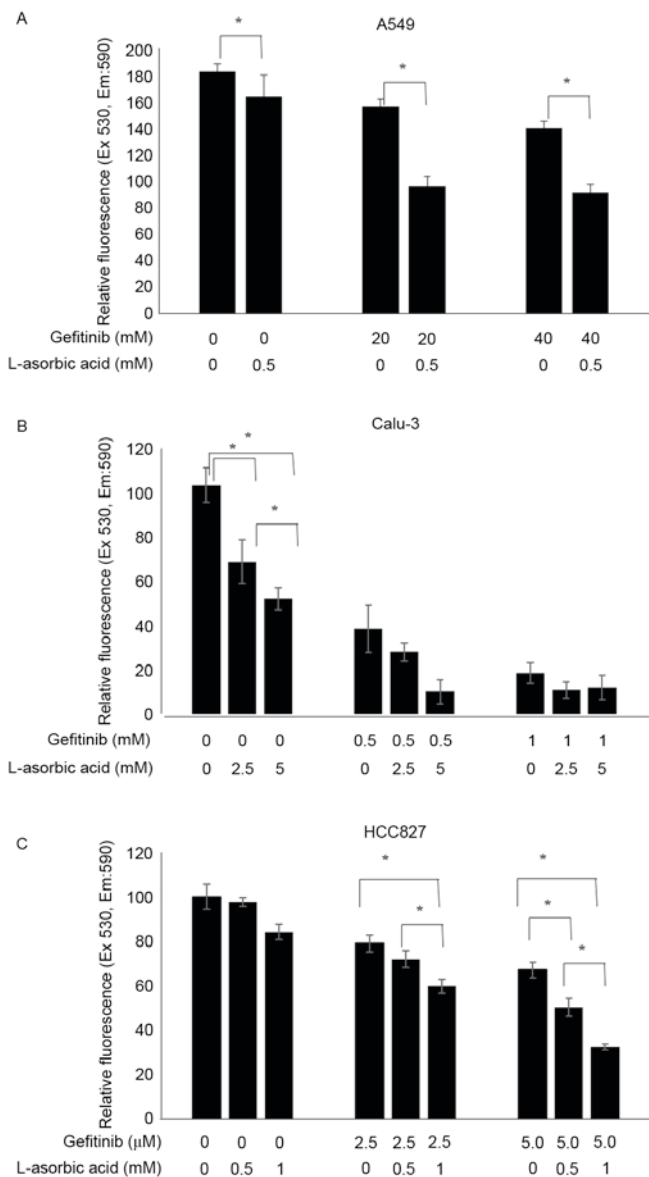


Figure 1. Increased antiproliferative activity of gefitinib with L-ascorbic acid in non-small cell lung cancer cell lines. A total of 1×10^6 cells were incubated with various concentrations of L-ascorbic acid and/or gefitinib, and the cytotoxic effects were assessed using an AlamarBlue assay. Cells were pretreated with L-ascorbic acid for 1 h, followed by gefitinib treatment for 48 h. Cell viabilities are expressed as the ratio of the absorbance of treated cells to control cells. Independent experiments were performed in triplicate and the data are presented as the mean \pm standard deviation. (A) A549 cells exposed to 0, 20 and 40 μ M gefitinib and 0 and 0.5 mM L-ascorbic acid. (B) Calu-3 cells exposed to 0, 0.5 and 1 μ M gefitinib and 0, 2.5 and 5 mM L-ascorbic acid. (C) HCC827 cells exposed to 0, 2.5 and 5.0 nM gefitinib and 0, 0.5 and 1 mM L-ascorbic acid. * $P < 0.05$, statistically significant difference between the groups. Ex, excitation; Em, emission.

growth inhibition (Fig. 2). Similar results were observed in all three cell lines examined.

Changes in the cell cycle and ROS production effected by gefitinib and L-ascorbic acid. The combination of L-ascorbic acid with gefitinib demonstrated a synergistic reduction in the number of cells in the S phase. In the Calu-3 cells, the percentage of cells in S phase of the cell cycle was reduced from 35.8% in the control cells to 17.9% in cells treated with gefitinib alone; this percentage was 23.1% in cells treated

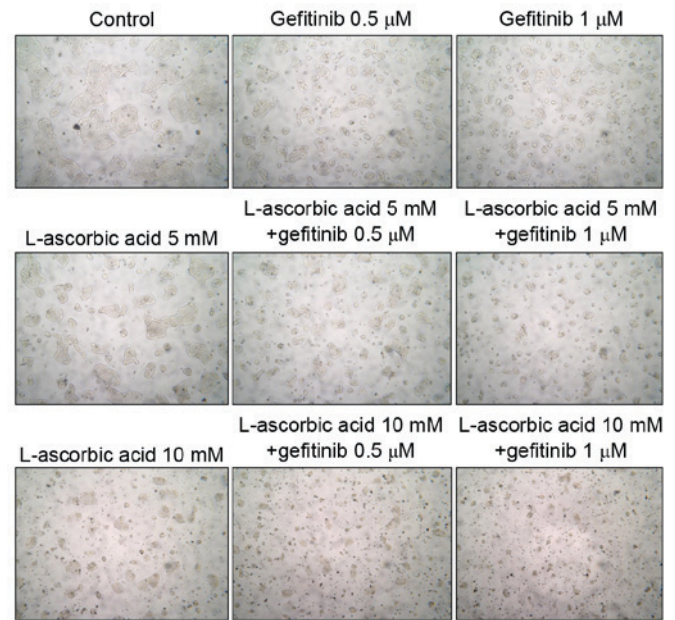


Figure 2. Microscopic images of Calu-3 cells treated with gefitinib and/or L-ascorbic acid. (Microscopic images were captured at magnification, $\times 100$, to illustrate the relative cellular growth patterns. A total of 1×10^6 cells were cultured in T25 culture flasks with/without L-ascorbic acid and gefitinib at various concentrations: 0.5 or 1 μ M of gefitinib and 5 or 10 mM L-ascorbic acid. Cells were pre-treated with L-ascorbic acid for 1 h and then treated with gefitinib for 48 h. The figure is representative of three independent experiments.

with L-ascorbic acid alone and 11% in cells co-treated with gefitinib and L-ascorbic acid. The highest observed arrest in the G_2 phase was with L-ascorbic acid alone, and gefitinib did not induce G_2 arrest. The effect of L-ascorbic acid on G_2 arrest was neutralized when it was administered in combination with gefitinib. Also, the antioxidant activity of L-ascorbic acid was observed: In Calu-3 cells, blocking EGFR tyrosine kinase with gefitinib induced high intracellular levels of ROS. This action was attenuated by the antioxidant L-ascorbic acid, which reduced intracellular ROS levels in Calu-3 cells to approximately the level observed in the untreated cells (Fig. 3).

Changes in EGFR phosphorylation following treatment with gefitinib and L-ascorbic acid. A total of three antibodies, Tyr992, Tyr1068 and Tyr845, were used to identify different phosphorylation loci of the EGFR. The addition of L-ascorbic acid to gefitinib did not significantly modify the phosphorylation levels of Erk1/2, Stat3 or EGFR in A549 cells, but slightly reduced the phosphorylation levels of Akt (Fig. 4, left panel). In Calu-3 cells, treatment with L-ascorbic acid alone was not observed to have a significant effect on EGFR phosphorylation or the associated downstream signaling pathways. By contrast, L-ascorbic acid co-administered with gefitinib decreased the phosphorylation levels of Akt and Erk1/2. The phosphorylation level of Stat3 was not significantly altered following the co-administration of L-ascorbic acid with gefitinib (Fig. 4, middle panel). In HCC827 cells, which exhibit an EGFR mutation and are sensitive to gefitinib, L-ascorbic acid administered in combination with gefitinib was not observed to significantly affect the signaling cascades downstream of the EGFR pathway. Monotherapy with gefitinib decreased

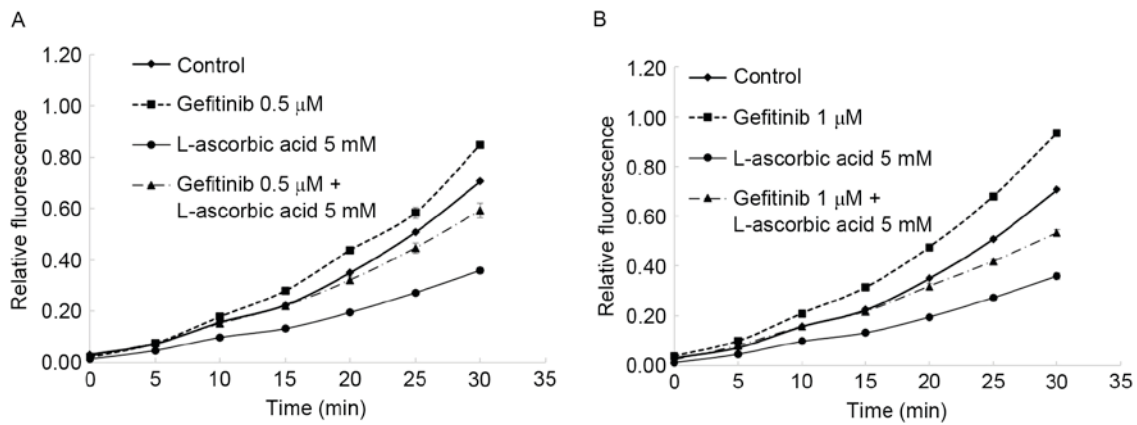


Figure 3. Changes in ROS in Calu-3 cells with L-ascorbic acid and/or gefitinib. A total of 1×10^4 cells in 96-well plates were treated with 5 mM L-ascorbic acid and/or 0.5-1 μ M gefitinib. (A and B) Using the DCFH-DA method (50 μ M DCFH-DA) to detect ROS, L-ascorbic acid reduced intracellular ROS levels in Calu-3 cells to approximately control cell levels. Relative fluorescence was measured for 30 min. The figure is representative of three independent experiments. ROS, reactive oxygen species; DCFH-DA, 2',7'-dichlorodihydrofluorescein diacetate.

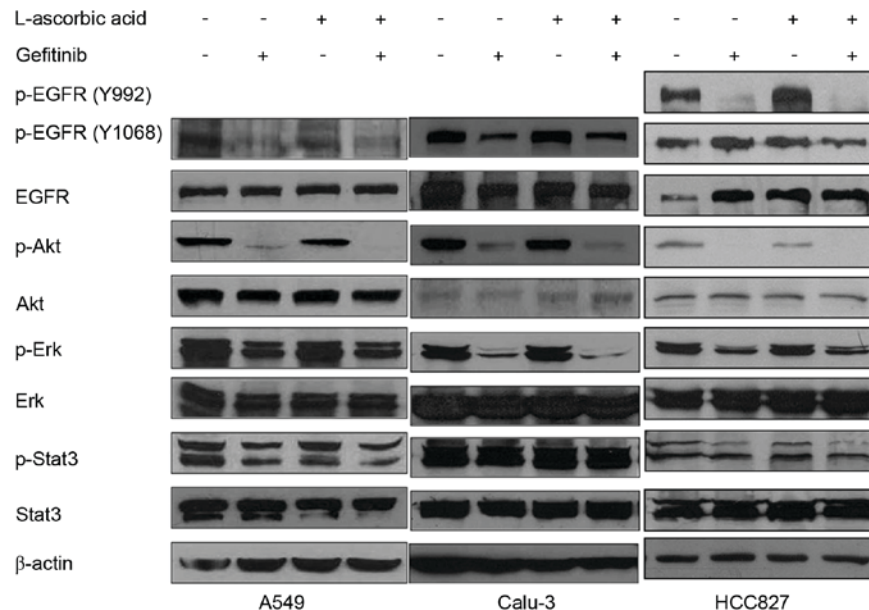


Figure 4. Immunoblotting analysis of gefitinib and/or L-ascorbic acid therapy in cancer cell lines. A total of 2×10^6 cells were cultured in serum-free medium with L-ascorbic acid or gefitinib. Calu-3 cells were pretreated with L-ascorbic acid for 30 min prior to gefitinib treatment, which was maintained for 2 h. A total of three antibodies were used to detect the various phosphorylation sites of EGFR. A549 cells were treated with 0.5 mM L-ascorbic acid and 20 μ M gefitinib; Calu-3 cells were treated with 2.5 mM L-ascorbic acid and 0.5 μ M gefitinib; HCC827 cells were treated with 1 mM L-ascorbic acid and 0.005 μ M gefitinib. The figure is representative of three independent experiments. EGF, epidermal growth factor; EGFR, EGF receptor; p, phosphorylated; Akt, protein kinase B; Erk, extracellular signal-regulated kinases; Stat3, signal transducer and activator of transcription 3.

the phosphorylation levels of the downstream molecules of the EGFR pathway, including Akt, in the HCC827 cell line (Fig. 4, right panel), but no significance was observed. It was also examined whether changes in ROS affected the signaling cascades downstream of the EGFR signaling pathway.

Discussion

Numerous small molecules have been developed to replace cytotoxic chemotherapy agents in cancer therapeutics; however, the development of novel resistance to these drugs, along with resistance mechanisms, remains problematic. Mutations in the receptor kinase domains of cancer cells are the principal cause of resistance to tyrosine kinase inhibitor

therapy (12,17,29). Imatinib mesylate, which is applied clinically in the treatment of chronic myelogenous leukemia (CML), is the first small-molecule drug for which resistance to tyrosine kinase inhibitors was recognized; the mechanism for resistance was identified to originate from a mutation in the kinase domain (7,13,31). The increase in intracellular ROS production induced by imatinib mesylate in patients with CML causes genetic instability and gene mutations, leading to mutation of the kinase domain and subsequent drug resistance. Antioxidants may decrease the levels of ROS present in cells and, therefore, decrease the risk of gene mutations and associated drug resistance (32).

Various previous *in vitro* and *in vivo* studies have suggested that certain antioxidants may increase the effects

of cytotoxic therapy. However, clinical studies using antioxidants as an adjuvant to chemotherapy have so far been unsuccessful (10,19,25,28,33,34). L-ascorbic acid may function as either a pro-oxidant or an antioxidant at physiological micromolar concentrations (normal range, <0.1 mM) in the presence of trace amounts of transition metals including copper and iron (20,34-40). There is controversy regarding the potential role and dose of L-ascorbic acid that should be recommended for use in cancer therapy (40). Gefitinib is the first small-molecule drug developed to inhibit tyrosine kinase in human NSCLC (41). Mutant forms of EGFR and increased EGFR copy numbers are considered to be successful predictive biomarkers for therapeutic responses.

Similar to imatinib mesylate, ROS may induce changes in cell signaling, genetic instability and other various steps in intracellular pathways. One of the functions of intracellular ROS is to decrease phosphatase activity and to upregulate tyrosine kinase once at a certain concentration; thus, ROS may serve a role in carcinogenesis (31). At present, there have been few studies on the effects of ROS on the phosphorylation of EGFR (42).

The present study hypothesized that ROS affects EGFR kinase activity and the sensitivity of cancer cells to EGFR tyrosine kinase inhibitor therapy, and that L-ascorbic acid may have an additive or synergistic effect when co-administered with an EGFR tyrosine kinase inhibitor. As the EGFR mutation is an important biomarker for the response to gefitinib, three human NSCLC cell lines were used, including Calu-3 and A549 (wild-type EGFR) and HCC827, which has a mutation in the EGFR tyrosine kinase domain (E746-A750 deletion) and is known to respond to gefitinib. The effects of L-ascorbic acid and gefitinib were synergistic in all cell lines when administered at various dosages. In the present study, gefitinib acted as a pro-oxidant and L-ascorbic acid attenuated the elevation in ROS levels in lung cancer cell lines. It was also revealed that L-ascorbic acid monotherapy reduced the percentage of Calu-3 cells in the S phase, and that this effect was augmented when administered in combination with gefitinib. Thomas *et al* demonstrated that L-ascorbic acid induces transient cell cycle arrest by delaying the accumulation and activation of *Cdc25C* (42).

The antiproliferative and anticancer mechanisms of L-ascorbic acid and gefitinib differ. L-ascorbic acid may kill or inhibit the growth of numerous tumor cell lines and also increase the potency of certain radiosensitizing drugs (43). Beyond its antioxidant role, the exact mechanisms underlying the effects L-ascorbic acid have yet to be elucidated. Although different concentrations of L-ascorbic acid were chosen for each cell line, none of the selected concentrations killed all the cells. In the present study, L-ascorbic acid administered alone inhibited cancer cell growth through its antioxidant function and by downregulating EGFR phosphorylation. Gefitinib alone generated ROS, inhibited kinase activity and downregulated Akt and Erk expression. In order to identify the optimal redox environment for EGFR tyrosine kinase inhibitors to obtain a maximum anticancer effect, the effect of ROS in EGFR tyrosine kinase inhibitor therapy was examined using nonspecific ROS blocking agents. It is not likely that ROS were the leading cause of the antitumor effects in the present study. These results indicate that L-ascorbic acid acts via its own antitumor

mechanisms that are separate to ROS activity, and that it is involved in the inhibition of the EGFR signaling cascade.

In conclusion, the present study has demonstrated that, when co-administered with gefitinib, L-ascorbic acid functions as an antioxidant and is associated with the downregulation of EGFR phosphorylation in three NSCLC cell lines. However, several issues remain to be resolved, including the practical delivery of a clinically relevant therapeutic dose of L-ascorbic acid, the optimal redox status that exhibits an antitumor effect and the *in vivo* responses. It is suggested that L-ascorbic acid may overcome the wild-type EGFR tyrosine kinase inhibitor resistance and induce a synergistic response when co-administered with EGFR tyrosine kinase inhibitors in NSCLC. These results also suggest that improved clinical outcomes in lung cancer therapy may be obtained through L-ascorbic acid therapy combined with an EGFR tyrosine kinase inhibitor therapy such as gefitinib or erlotinib.

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