Increased efficacy of gefitinib on cisplatin-resistant wild-type epidermal growth factor receptor non-small cell lung cancer cells

Amin Li1,2*, Weiya Cao1*, Xueke Liu1*, Yinci Zhang1*, Yongfang Ma1, Ruyue Xu1, Xiaolong Tang1

1Medical School, Anhui University of Science & Technology, Huainan, China; 2First Affiliated Hospital, Anhui University of Science & Technology (Huainan First People’s Hospital), Huainan, China

Contributions: (I) Conception and design: A Li, W Cao, X Tang; (II) Administrative support: X Tang; (III) Provision of study materials or patients: A Li, X Tang; (IV) Collection and assembly of data: Y Ma, R Xu; (V) Data analysis and interpretation: A Li, X Liu, Y Zhang; (VI) Manuscript writing: All authors; (VII) Final approval of manuscript: All authors.

*These authors contributed equally to this work.

Background: Gefitinib is a first-generation epidermal growth factor receptor-tyrosine kinase inhibitor (EGFR-TKI) that has become first-line treatment for patients with mutant EGFR non-small cell lung cancer (NSCLC). Despite its anti-tumor activity, the benefit of gefitinib in patients with wild-type EGFR NSCLC is debated. This work aimed to evaluate the effects of gefitinib on cisplatin-resistant wild-type EGFR NSCLC cells in in vitro and in vivo animal xenografts.

Methods: We established a cisplatin-resistant wild-type EGFR NSCLC cell line, H358R, to evaluate the cells’ sensitivity to gefitinib compared with that of parental cell line H358. We first tested the p-EGFR of gefitinib’s target in H358R and H358 cell line by western blot and immunofluorescence. 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT), clone formation assay, flow cytometry and annexin V-fluorescein/propidium iodide staining were used to investigate cellular proliferation and apoptosis of H358R/H358 cells treated with gefitinib, and the anti-tumor effect was evaluated in female BALB/c nude mice models of xenografts in vivo.

Results: EGFR and the downstream node molecules ERK and AKT were significantly more phosphorylated in H358R than in the parental cells and were inhibited by gefitinib. In H358R cells, gefitinib increased the inhibition of cell survival/proliferation, and the promotion of apoptosis in vitro. The increased anti-tumor effect was present also in H358R xenografts in vivo.

Conclusions: Abnormal activation of EGFR in H358R cells results in enhanced sensitivity to gefitinib. The improved efficacy of gefitinib on cisplatin-resistant wild-type EGFR NSCLC cells suggests that gefitinib as sequential therapy for patients with cisplatin-resistant wild-type EGFR NSCLC should be considered.

Keywords: Gefitinib; cisplatin; epidermal growth factor receptor (EGFR); non-small cell lung cancer (NSCLC); wild-type EGFR

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Introduction

Treatment with epidermal growth factor receptor-tyrosine kinase inhibitors (EGFR-TKIs) has had promising results in the treatment of patients with non-small cell lung cancer (NSCLC) that harbor a mutated, activated form of EGFR (1). Gefitinib (IRESSA, AstraZeneca Pharmaceuticals, Wilmington, DE, USA) was the first TKI approved by the European Medicines Agency for first-line therapy in these
patients (2). However, for patients with advanced wild-type EGFR NSCLC, the treatment effect of gefitinib, as well as that of other EGFR-TKIs, is not satisfactory. In recent years, immunotherapy has shown advantages in patients with wild-type EGFR NSCLC. According to the latest NSCLC treatment guidelines, the PD-L1 inhibitor, durvalumab is recommended first as consolidation therapy after concurrent chemoradiation (Type 1A evidence) for locally advanced and driver-negative NSCLCs. Pabolizumab monotherapy, limited to PD-L1 ≥50% (Type 1A evidence), PD-L1 1–49% (Type 2A evidence), and pabolizumab combined with pemetrexed and platinum (Type 1A evidence) are first-line recommended treatment in stage IV NSCLC without driver gene mutation (3,4).

Thus, immunotherapy likely plays a significant role in the treatment of NSCLC without driver mutations. However, for most driver-negative, especially wild-type EGFR NSCLC patients, even with PD-L1 ≥50%, platinum-containing chemotherapy remains the backbone of therapy (5).

A problem is that drug resistance inevitably accompanies long-term use of platinum-containing chemotherapy, which hampers its therapeutic efficacy and clinical outcomes (6). In such circumstances, clinicians often give replacement chemotherapy drugs or palliative therapy with EGFR-TKIs to patients who cannot tolerate chemotherapy and cannot benefit from immunotherapy. Some patients benefit from EGFR-TKIs palliative treatment, and clinical trials have found that wild-type EGFR patients benefit from EGFR-TKI as a maintenance treatment after platinum-based first-line therapy (7).

First-line chemotherapy often changes the genetic level or molecular biology phenotype of tumor cells, and these changes may affect the response to subsequent treatment (8,9). Therefore, we speculate that benefit from EGFR-TKI after platinum-containing chemotherapy may be related to changes in biological characteristics of the EGFR-TKI target. This target, EGFR, is the most important prosurvival signal receptor in wild-type EGFR NSCLC cells. The abnormal activation of EGFR pathways induces tumor-cell proliferation, anti-apoptosis, angiogenesis, and drug resistance (10).

In our study, we first investigated the activation of EGFR in a wild-type EGFR NSCLC parental cell line and a cisplatin-resistant cell line, and assessed the efficacy of gefitinib on these two cell lines in vitro. Further, we assessed the inhibitory effect of gefitinib on the two xenografts in female BALB/c nude mice that do not reject xenografts and can maintain their biological characteristics. We found that gefitinib enhanced inhibition of cisplatin-resistant wild-type EGFR NSCLC cells and supports the view that gefitinib is a sequential therapy for patients with cisplatin-resistant wild-type EGFR NSCLC.

We present the following article in accordance with the ARRIVE reporting checklist (available at: http://dx.doi.org/10.21037/tcr-20-1441).

**Methods**

**Cell lines, chemicals and antibodies**

Human wild-type EGFR NSCLC cell line H358 (ATCC Cat# CRL-5807, RRID: CVCL_1559) was obtained from the American Type Culture Collection, Rockville, MD, USA. Cisplatin-resistant cell line derived from H358, named H358<sup>R</sup>, was induced by constant exposure to cisplatin (2 μmol/L) to imitate acquired resistance. Both cell lines were cultured in 10% fetal bovine serum-containing medium (RPMI1640, Gibco, Thermo Fisher Scientific) and maintained in a 5% CO<sub>2</sub> incubator at 37 °C. Cisplatin (A8321) and gefitinib (ZD1839) were purchased from APEXBIO Technology LLC (Houston, Texas, USA) and Med Chem Express (Monmouth, Junction, USA), respectively. 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) and annexin V-FITC/Pi Kit were obtained from Sigma-Aldrich (St. Louis, MO, USA). Total EGFR (Cell Signaling Technology Cat# 4267, RRID: AB_2246311), phospho-EGFR (Cell Signaling Technology Cat# 9911, RRID: AB_10695905) antibodies were provided from Cell Signaling Technology (Danvers, MA, USA).

**IC<sub>50</sub> measurements**

H358 and H358<sup>R</sup> cells were separately plated in 96-well plates overnight and treated with cisplatin for 48 h. The cells were incubated with MTT (5 mg/mL) for 4 h at 37 °C. Formazan crystals were solubilized in dimethyl sulfoxide, and optical density (OD) was read at 570 nm with a microplate reader. IC<sub>50</sub> were calculated as described (11).

**Immunofluorescence analysis**

Phosphorylated EGFR in H358 and H358<sup>R</sup> cells was
detected by immunofluorescence. Cells were separately plated in 24-well plates overnight and fixed by 4% paraformaldehyde, permeabilized by 90% methanol, and saturated by 5% bovine serum albumin. The cells were incubated in the p-EGFR antibody diluted in 1% bovine serum albumin for 1 h at 4 °C and coupled to fluorescent (Alexa Fluor 488) secondary antibody at room temperature for 30 min. Fluorescence was visualized with a fluorescence microscope (Leica Microsystems Inc., Germany).

**Western blot analysis**

Total protein was extracted from cells lysed with radioimmunoprecipitation buffer (RIPA, Beyotime Biotechnology, Shanghai, China) containing a protease inhibitor cocktail (Beyotime Biotechnology, Shanghai, China) and centrifuged. Protein concentration of the supernatant was measured with BCA200 protein assay kit (Biosharp Life Science, Hefei, China) and equalized before loading to the gel. Target proteins were separated by sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) and electrotransferred onto a polyvinylidene fluoride membrane (Millipore, USA). After blocking with Tris-buffered saline/Tween 20 containing 5% fat-free milk, detecting antibodies were applied. Protein bands were visualized with a chemiluminescence detection kit (Thermo), and images were captured with a scanner using Quality One software (Bio-Rad).

**Viability assay**

Cell viability was measured with MTT assay in accordance with the manufacturer’s instructions. H358 and H358R cells were seeded into 96-well plates and cultured for 24 h in a 5% CO2 incubator at 37 °C. The cells were treated with gefitinib for 48 h. Carrier dimethyl sulfoxide was used as a control. Ten microliters of MTT (5 mg/mL) were added to each well and incubated for 4 h. After aspiration, 100 μL of dimethyl sulfoxide was added. Cell proliferation/inhibition rate was calculated with GraphPad Prism Version 5.0 software according to ODλ=570nm by use of a microplate reader (Bio-Rad Model 680, Bio-Rad Laboratories Inc., Hertfordshire, UK).

**Clone formation assay**

Cells were seeded in six-well culture plates at a density of 1,000 cells per well and cultured for 24 h. The cells were treated with gefitinib for another 48 h. The medium containing gefitinib was removed and replaced with fresh medium. The cells were cultured for about 10 days until colonies were visible. Colonies were fixed with 4% paraformaldehyde and stained with crystal violet. The cell colonies were counted and compared to those of the control group.

**Cell-cycle analysis**

H358 and H358R cells were planted in 12-well plates and incubated for 24 h followed by treatment with gefitinib for an additional 48 h. The cells were dissociated and fixed with cold 75% ethanol, then incubated with RNase (100 ng/mL) for 1 h and stained with propidium iodide (PI, 50 ng/mL) for 30 min. Dye was removed, and the cells were resuspended with PBS for cell-cycle analysis by flow cytometry (BD FACSCalibur, USA).

**Apoptosis test**

To detect cell apoptosis, cells were seeded on glass slides prepared in 24-well plates and incubated for 24 h, followed by gefitinib treatment for an additional 48 h, then incubated with annexin V-FITC/PI at room temperature for 15 min in the dark. The staining was visualized with a fluorescence microscope (Leica Microsystems Inc., Germany). Flow cytometry was performed after cells were dissociated and incubated with fluorescence staining.

**Animal experiments**

Twelve female BALB/c nude mice of about 5–6 weeks of age with an average body weight of 18–20 g were provided by Nanjing Junke Biotechnology Co. Ltd. H358 and H358R cell suspensions (100 μL; 5×107/mL) were separately injected subcutaneously into the left shoulders of mice randomly divided into two groups. When tumors reached about 50–100 mm³, the 2 groups of tumor-bearing mice were again randomly divided into 2 groups (H358 group: control/gefitinib treatment; H358R group: control/gefitinib treatment). The “control” group had no treatment, and the “gefitinib” group received gefitinib (150 mg/kg) daily orally by gavage. Tumor volume and body weight were measured twice a week, and the formula [(length) × (width)²/2] was used to estimate the tumor growth. Twenty-one days after administration, the mice were sacrificed, and tumor tissues were isolated. Animal experiments were carried out in...
accordance with the principles and procedures approved by the Committee on the Ethics of Animal Experiments of Anhui University of Science and Technology. All animals were nursed in the SPF-level animal room of the Central Laboratory of Medical School Anhui University of Science and Technology (NO.: AUST2019-10062).

Statistical analysis
All experiments were performed in triplicate. Group differences were analyzed with Student’s t-test using GraphPad Prism 5 (GraphPad Prism, RRID: SCR_002798) (GraphPad Software, Inc., San Diego, CA, USA). Significance was defined as P<0.05.

Results

Drug resistance of H358<sup>R</sup> cell line to cisplatin
H358 cells were induced by continuous treatment with 2 μmol/L cisplatin for 2 months, then the cells (named H358<sup>R</sup>) were permitted to proliferate and were sub-cultured in the presence of cisplatin. H358<sup>R</sup> cell morphology changed significantly compared to that of its parental cells (Figure 1A). To monitor the resistance, we measured the cells’ IC<sub>50</sub> to cisplatin, which were 12.2±0.8 μM for H358 and 27.9±1.8 μM for H358<sup>R</sup> cells (Figure 1B).

EGFR phosphorylation enhanced in H358<sup>R</sup> cells and suppressed by gefitinib
We first detected EGFR phosphorylation in H358 and H358<sup>R</sup> cells by immunofluorescence assay. As illustrated in Figure 1C, green fluorescence intensity in H358<sup>R</sup> cells was significantly higher than that in its parental cells, suggesting increased EGFR phosphorylation in H358<sup>R</sup> cells. We further measured the phosphorylation level of EGFR and the downstream node molecules ERK/AKT in these cells with western blot assay. The results documented that phosphorylation of EGFR/ERK/AKT in H358<sup>R</sup> cells was significantly increased, and gefitinib inhibited their phosphorylation in a dose-dependent manner (Figure 1D).

Gefitinib increased inhibition of proliferation by G0/G1 arrest in H358<sup>R</sup> cells
To test the sensitivity of H358<sup>R</sup> cells with increased EGFR phosphorylation to gefitinib, we tested their proliferation activity and that of the parental cells after gefitinib treatment separately by MTT (Figure 2A) and clone formation assay (Figure 2B). The results showed that the inhibitory effect of gefitinib on H358<sup>R</sup> cells was further enhanced. We also analyzed whether the effect of gefitinib on cell proliferation was associated with cell-cycle dysregulation in H358<sup>R</sup> cells based on DNA content by flow cytometry analysis. With H358 cells treated with gefitinib, there was a moderate increase in the percentage of cells in G0/G1 phase compared with the percentage in controls. The percentage of H358<sup>R</sup> cells in G0/G1 phase was increased more compared with the values of its parental cells H358 (Figure 2C).

Gefitinib promoted apoptosis in H358<sup>R</sup> cells
Annexin V-FITC/PI staining and flow cytometry analysis were performed to evaluate the apoptotic effect of gefitinib in H358 and H358<sup>R</sup> cells. As shown in Figure 3A, annexin V-FITC/PI staining revealed that the green fluorescence intensity of H358<sup>R</sup> cells treated with gefitinib was significantly more than that of H358 cells. In Figure 3B, flow cytometry analysis also displayed that gefitinib had a significantly higher apoptosis ratio in H358<sup>R</sup> cells.

Gefitinib increased the anti-tumor effect in H358<sup>R</sup> xenograft in vivo
H358 and H358<sup>R</sup> tumor-bearing nude mouse model were prepared to evaluate the anti-tumor effect of gefitinib. As shown in Figure 4A, when mice were treated with the same dose of gefitinib, the growth rate of the H358<sup>R</sup> xenograft tumor (n=3) was significantly slower than that of H358 tumors (n=3). Gefitinib had moderate anti-tumor activity in H358 xenograft because the wild-type EGFR NSCLC cell line is not sensitive to gefitinib. However, the tumor growth of H358<sup>R</sup> xenograft was significantly suppressed, and there was a steady trend of reduced tumor volume, especially during the first 15 days of gefitinib treatment. The growth inhibition of H358 and H358<sup>R</sup> xenograft tumor at the 21st day after treatment was 28.0±1.4% and 52.7±3.1%, respectively (Figure 4B,C), and the difference between the 2 groups was statistically significant. The weights of the tumor-bearing mice were stable, and weights of the treated mice were slightly lower than those of the tumor-bearing control mice. Thus, these in vivo findings are evidence that the anti-tumor effect of gefitinib was increased in cisplatin-
Figure 1 Resistance of cisplatin-resistant cell line H358R and phosphorylation of EGFR in H358R cells. (A) Morphological changes of H358R cells compared with those of their parental cells (200×). (B) Cell viability after cisplatin treatment with MTT assay and IC_{50} were calculated as described by Chou and colleagues (11). (C) Immunofluorescence green fluorescence intensity of EGFR in H358R and their parental cells (200×). (D) Western blot analysis of the phosphorylation of EGFR and downstream node molecules ERK/AKT and anti-phosphorylation effect of gefitinib (GFB) in H358R cells. *, P<0.05.
resistant H358^R xenograft with no overt toxic side effects.

**Discussion**

NSCLC accounts for most lung cancers and remains the leading cause of cancer-related deaths worldwide. The molecular driver EGFR mutation plays an important role in NSCLC, and it is a target for personalized therapy (12-15). EGFR-TKIs are efficacious in treatment of patients with NSCLC that harbor a mutated, activated form of EGFR (16). However, effectiveness of the inhibitors is lost in wild-type EGFR NSCLC patients, which limits their clinical application (17). Despite much effort to develop new treatment methods, platinum-based chemotherapy remains the first-line treatment in most wild-type EGFR NSCLC.

Drug resistance with resultant tumor progression after platinum-based chemotherapy is common (18), and many patients have difficulty tolerating other chemotherapy drugs. This difficult situation requires an effective drug with less toxic side effects. Gefitinib, an aniline quinazoline derivative, is the first generation of EGFR-TKI that competes for the Mg-ATP binding site on the EGFR-TK catalytic region. It inhibits EGFR phosphorylation and
activation of mitogen-activated protein kinase; blocks tumor growth, metastasis, and angiogenesis; and promotes tumor-cell apoptosis (19,20). Gefitinib’s beneficial effects have been demonstrated as first-line treatment in mutant EGFR NSCLC, but in wild-type EGFR NSCLC patients, the indication of gefitinib is debated (21).

The aim of our study was to evaluate EGFR modifications in cisplatin-resistant wild-type EGFR NSCLC cells and the sensitivity of gefitinib treatment before and after cisplatin resistance. Our work provided evidence of activation of EGFR induced by treatment of cisplatin in the wild-type EGFR NSCLC cell line H358.
This activation results in gefitinib sensitization, enhanced inhibition of EGFR and downstream activation of Ras/Raf/MAPK and PI3K/AKT/mTOR pathways (Figure 5). These events explain the enhanced anti-proliferative and pro-apoptotic effects of gefitinib in our in vitro experiments.

Sensitization to gefitinib induced by cisplatin pretreatment may involve the activation of the proto-oncogene tyrosine-protein kinase Src in the EGFR pathway. Cisplatin-elicited DNA damage activates nuclear as well as cytoplasmatic signaling pathways that deliver compensatory survival signals, of which Src-EGFR-ERK is an important pathway for EGFR high-expression cancers (18,22). Abnormal activation of Src mediates cisplatin resistance in several cell types (23-25). Ligand-independent, Src-mediated activation of EGFR induced by cisplatin treatment also has been reported (26). Src kinase participates in signaling pathways that control diverse biological activities, including MAPK and PI3K/AKT/mTOR pathways (27-29). In our study, the EGFR downstream node molecule ERK of the MAPK pathway as

Figure 4 Enhanced anti-tumor effect of gefitinib (GFB) in H358R xenograft in vivo. (A) Tumor growth curves of gefitinib-treated H358 and H358R xenograft (*, P<0.05). (B,C) Tumor volume of H358 and H358R xenograft treated with gefitinib (GFB), respectively.

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well as AKT are activated in cisplatin-resistant H358 cells, a finding that is like that of Kong et al. (30). The activities of ERK/AKT were significantly reduced with application of gefitinib, suggesting that gefitinib inhibits EGFR pathway activation in the presence of cisplatin resistance.

We found that gefitinib is effective also in xenograft models in vivo. As seen in tumor growth curves, the first 2 weeks of gefitinib treatment significantly and stably inhibited tumor growth in the cisplatin-resistant mouse model. With longer treatment, however, the tumors developed, which may be due to the gradual desensitization of gefitinib, but the overall growth rate was still significantly lower than that of control animals. These findings suggest a new perspective of gefitinib and other EGFR-TKIs: sequential maintenance therapy in cisplatin-resistant wild-type EGFR NSCLC. This possibility will require testing with more cell types and more EGFR inhibitors.

The results of our study concur with those of other research that suggest that EGFR-TKI maintenance could be more efficient in wild-type EGFR NSCLC controlled by initial platinum-based therapy (31-33). Our experiments also show that cisplatin resistance is associated with abnormal activation of EGFR. Therefore, we suspect that combination with EGFR targeting strategies may enhance anti-tumor response in cisplatin-resistant wild-type EGFR NSCLC, which we will study in future experiments.

Conclusions

Abnormally activated EGFR in the cisplatin-resistant cell line H358R increased the sensitivity of these cells to gefitinib. This finding suggests that patients with cisplatin-resistant wild-type EGFR NSCLC could benefit from maintenance treatment with gefitinib.

Figure 5 Signal pathway of cisplatin-resistant cells sensitized to gefitinib.
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Footnote

Reporting Checklist: The authors present the study in accordance with the ARRIVE reporting checklist. Available at http://dx.doi.org/10.21037/tcr-20-1441

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Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. Animal experiments were carried out in accordance with the principles and procedures approved by the Committee on the Ethics of Animal Experiments of Anhui University of Science and Technology. All animals were nursed in the SPF-level animal room of the Central Laboratory of Medical School Anhui University of Science and Technology (NO.: AUST2019-10062).

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