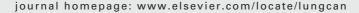


available at www.sciencedirect.com







Effects of different combinations of gefitinib and irinotecan in lung cancer cell lines expressing wild or deletional EGFR

Tatsu Shimoyama^{a,b}, Fumiaki Koizumi^a, Hisao Fukumoto^a, Katsuyuki Kiura^b, Mitsune Tanimoto^b, Nagahiro Saijo^a, Kazuto Nishio^{a,*}

Received 8 November 2005; accepted 1 March 2006

KEYWORDS

Gefitinib; CPT-11; SN-38; EGFR; Combination; Lung cancer Summary EGFR mutations are a major determinant of lung tumor response to gefitinib, an EGFR-specific tyrosine kinase inhibitor. Obtaining a response from lung tumors expressing wildtype EGFR is a major obstacle. The combination of gefitinib and cytotoxic drugs is one strategy against lung cancers expressing wild-type EGFR. The DNA topoisomerase inhibitor irinotecan sulfate (CPT-11) is active against lung cancer. We examined the sensitivity of lung cancers expressing wild- or mutant-type EGFR to the combination of gefitinib and CPT-11. The in vitro effect of gefitinib and SN-38 (the active metabolite of CPT-11) was examined in seven lung cancer cell lines using the dye formation assay with a combination index. When administered concurrently, gefitinib and SN-38 had a synergistic effect in five of the seven cell lines expressing wild-type EGFR, whereas the combination was antagonistic in PC-9 cells and a PC-9 subline resistant to gefitinib and expressing deletional mutant EGFR (PC-9/ZD). When administered sequentially, treatment with SN-38 followed by gefitinib had remarkable synergistic effects in the PC-9 and PC-9/ZD cells. In an in vivo tumor-bearing model, this combination had a scheduledependent synergistic effect in the PC-9 and PC-9/ZD cells. An immunohistochemical analysis of the tumors in mice treated with CPT-11 and gefitinib demonstrated that the number of Ki-67 positive tumor cells induced by CPT-11 treatment was decreased when CPT-11 was administered in combination with gefitinib. In conclusion, the sequential combination of CPT-11 and gefitinib is considered to be active against lung cancer.

 $\ensuremath{\mathbb{C}}$ 2006 Elsevier Ireland Ltd. All rights reserved.

* Corresponding author at: Tel.: +81 3 3542 2511; fax: +81 3 3547 5185.

E-mail address: knishio@gan2.res.ncc.go.jp (K. Nishio).

1. Introduction

Lung cancer is one of the leading causes of cancer-related death, despite the use of conventional chemotherapy regi-

^a Shien-Lab and Medical Oncology, National Cancer Center Hospital, Pharmacology Division, National Cancer Center Research Institute, Tsukiji 5-1-1, Chuo-ku, Tokyo 104-0045, Japan

^b Department of Medicine II, Okayama University Medical School, 2-5-1 Shikata-cho, Okayama 700-8558, Japan

14 T. Shimoyama et al.

mens. The epidermal growth factor receptor (EGFR) is frequently expressed in non-small cell lung cancer (NSCLC) and is correlated with a poor prognosis. Gefitinib ('Iressa') is an orally active, selective EGFR-tyrosine kinase inhibitor that blocks signal transduction pathways. Its clinical efficacy has been show in refractory NCSLC patients, but the survival benefit of this agent remains unclear. EGFR mutations have been identified in NSCLC, and lung cancers carrying the EGFR mutation have been reported to be hyperresponsive to gefitinib [1,2]. Mutant EGFR is a major determinant of lung tumor response to gefitinib, but the hyperresponsiveness of tumors expressing mutant EGFR has been observed in a small population. Now, obtaining a clinical benefit in lung tumors expressing wild-ype EGFR is a major obstacle. The combination of gefitinib and cytotoxic drugs is one strategy against lung cancers expressing wild-type EGFR. The DNA topoisomerase I inhibitor irinotecan (CPT-11) is a key drug in the treatment of patients with lung cancer and has been shown to prolong survival. SN-38 is the active metabolite of CPT-11 in vitro. The objective of this study was to determine the potential therapeutic utility of gefitinib when combined with CPT-11 therapy to lung cancer cell according to the treatment schedule and EGFR status.

Acquired resistance to gefitinib is also of clinical interest. Recently, Kobayashi et al. [3] reported that an EGFR mutation was related to the development of acquired resistance to gefitinib. We have established subclone PC-9/ZD cells that are resistant to gefitinib [4]. Our results suggested that another mechanism of resistance was active in PC-9/ZD cells. The effect of the combination of gefitinib and SN-38 in these PC-9/ZD cells was also examined.

2. Materials and methods

2.1. Drugs and chemicals

Gefitinib (*N*-(3-chloro-4-fluorophenyl)-7-methoxy-6-[3-(morpholin-4-yl)propoxy]quinazolin-4-amine) was provided by AstraZeneca (Cheshire, UK). Gefitinib was dissolved in dimethyl sulfoxide (DMSO) for the in vitro study. CPT-11 and SN-38 were obtained from Yakult Honsha (Tokyo, Japan) and were dissolved in dimethyl sulfoxide (DMSO) for both of the in vitro studies.

2.2. Cells and cultures

Human NSCLC cell lines PC-9, PC-7, and PC-14 derived from untreated patients with pulmonary adenocarcinoma were provided by Professor Y. Hayata, Tokyo Medical College. A small cell lung cancer cell line, H69, was established at the National Cancer Institute (Bethesda, MD, USA). The gefitinib-resistant subline, PC-9/ZD, was established from intrinsic hypersensitive cell PC-9 [5] in our laboratory [4]. A small cell lung cancer cell line, SBC-3, and an adenocarcinoma cell line, A549, were obtained from the Japanese Cancer Research Resources Bank (Tokyo, Japan). All cell lines were maintained in RPMI1640 (Nikken Bio Med. Lab., Kyoto, Japan) supplemented with 10% heat-inactivated fetal calf serum, 100 µg/ml streptomycin, and 100 units/ml

penicillin in an incubator at 37 °C and 100% humidity in 5% CO_2 and air, as described previously [6].

2.3. RT-PCR

Specific primers designed for EGFR CDS were used to detect the EGFR mRNA, as described elsewhere [1]. Sixteen first-strand cDNAs were synthesized from the cells' RNA using an RNA PCR Kit (TaKaRa Biomedicals, Ohtsu, Japan). After the reverse transcription of 1 μg of total RNA with Oligo(dT)-M4 adaptor primer, the whole mixture was used for PCR with two oligonucleotide primers (5'-AATGTGAGCAGAGGCAGGGA-3' and 5'-GGCTTGGTTTGGAGCTTCTC-3). PCR was performed with an initial denaturation at 94 °C for 2 min and 25 cycles of amplification (denaturation at 94 °C for 30 s, annealing at 55 °C for 60 s, and extension at 72 °C for 105 s).

2.4. Western blot analysis

The cultured cells were washed twice with ice-cold phosphate buffered saline (PBS), lysate in EBC buffer (50 mM Tris-HCI, pH 8.0; 120 mM NaCI; 0.5% Nonidet P-40; 100 mM NaF; 200 mM Na orthovanadate; and 10 mg/ml each of leupeptin, aprotinin and phenylmethylsulfonyl fluoride). The lysate was cleared by centrifugation at $20,000 \times g$ for 5 min, and the protein concentration of the supernatant was measured using a BCA protein assay (Pierce, Rockford, IL, USA). For immunoblotting, 20 µg samples of protein were electrophoretically separated on a 7.5% SDS-polyacrylamide gel and transferred to a polyvinylidene difluoride (PVDF) membrane (Millipore, Bedford, MA, USA). The membrane was then probed with rabbit polyclonal antibodies against EGFR, HER2/neu, Her3 and Her4 (Santa Cruz Biotech, Santa Cruz, CA, USA) and phospho-EGFR specific for Tyr 845, Tyr 1045, and Tyr 1068 (numbers 2231, 2235 and 2234; Cell Signaling, Beverly, MA, USA).

2.5. Growth-inhibition assay

We used the tetrazolium dye (3,(4,5-dimethyl-2-thiazolyl)-2,5-diphenyl-2H-tetrazolium bromide, MTT) assay to evaluate the cytotoxicity of various drug concentrations. After incubation for 72 h at 37 °C, 20 µl of MTT solution (5 mg/ml in PBS) was added to each well; the plates were then incubated for a further 4h at 37 °C. After centrifuging the plates at $200 \times g$ for 5 min, the medium was aspirated from each well and 180 µl of dimethylsulfoxide was added to each well to dissolve the formazan. Optical density was measured at 562 and 630 nm using a Delta Soft ELISA analysis program interfaced with a Bio-Tek Microplate Reader (EL-340; Bio-Metallics, Princeton, NJ, USA). Each experiment was performed in six replicate wells for each drug concentration and was independently performed three or four times. The IC₅₀ value was defined as the concentration needed for a 50% reduction in the absorbance, as calculated based on the survival curves. Percent survival was calculated as follows:

Download English Version:

https://daneshyari.com/en/article/2144551

Download Persian Version:

https://daneshyari.com/article/2144551

<u>Daneshyari.com</u>