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Clinical development of EGFR-tyrosine kinase inhibitors in Japan

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Abstract Although the initial impact of the epidermal growth factor receptor-tyrosine kinase inhibitor (EGFR-TKI) gefitinib may have been less than spectacular in the field of non-small cell lung cancer (NSCLC), this EGFR-TKI does offer a therapy that, at least in the short term, markedly reduces tumors without bone marrow suppression including neutropenia and without causing severe nausea and vomiting even in NSCLC patients with the worst prognosis. This raises the possibility of putting the disease under control if only temporarily. Now we must be aware that overcoming gene mutation in lung cancer is the next significant milestone for new therapeutics. This report discusses clinical trials of EGFR-TKIs focusing on Japanese contributions to current knowledge, *EGFR* mutation, and future directions. A Japanese phase I clinical trial saw the first super-responders to gefitinib. Two randomized phase II trials identified Japanese, females, and those with adenocarcinoma of the lung as specific populations sensitive to gefitinib. Unexpectedly, in the context of first-line chemotherapy four phase III trials gave completely negative results for additional clinical benefit by EGFR-TKIs combined with standard chemotherapy. However, subset analysis suggested efficacy of this treatment strategy in nonsmokers and patients harboring activated-type *EGFR* mutations. In the settings of second-line and later therapy, two independent randomized placebo-controlled trials, BR.21 with erlotinib and ISEL with gefitinib, revealed better duration of overall survival, time to progression, and response rate in the EGFR-TKI versus control groups, although the result was nonsignificant in the latter study. Data suggesting that adenocarcinoma, Asian race, female, and nonsmoker are associated with better response to EGFR-TKI may be closely related with phenotype of *EGFR* mutations, making this parameter a "response predictive marker." On the other hand, some reports have stated that gene amplification of *EGFR* by FISH analysis shows better correlation with clinical benefit of EGFR-TKIs than that assessed by other means in large-scale phase III trials (BR21 and ISEL). Further validation of response predictive markers is needed. Recent studies of EGFR-TKIs in NSCLC provide novel biological insights and have given birth to the concept of patient selection for this disease. Further investigation of the biological significance of *EGFR* mutation and its validation as response predictive marker will lead to better treatments to come for NSCLC.

Keywords EGFR-TKI · Gefitinib · Erlotinib · *EGFR* mutation

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Introduction

Epidermal growth factor receptor-tyrosine kinase inhibitors (EGFR-TKIs) have been clinically available for the treatment of nonsmall cell lung cancer

(NSCLC) for the past 4 years. In the course of clinical development of EGFR-TKIs, in comparison with conventional anticancer agents many unexpected findings were observed such as relating to tumor shrinkage, specific responder subsets, adenocarcinomatous disease, and gene mutation. Hence although knowledge concerning EGFR-TKIs and *EGFR* gene mutation is advancing in the laboratory setting, clinically it is unclear how we should use EGFR-TKIs in NSCLC and which patients might benefit most from these agents. In this review, clinical trials of EGFR-TKIs are recounted and a key factor for drug sensitivity, *EGFR* mutation, is discussed.

Clinical trials of EGFR-TKIs

Four phase I trials of EGFR-TKI including one Japanese study were performed in a total of 254 patients [4, 8]. These trials defined diarrhea and liver function test abnormality as dose-limiting factors. Five of 23 patients demonstrated partial responses (PRs) without dose–response tendency (Table 1). Toxicity profiles were quite different to those commonly observed with conventional anticancer agents. Ten percent of patients failed treatment at doses >600 mg/day and these early studies could not identify an optimal dosing schedule. Based on the results of phase I, the phase II IDEAL1 study was conducted in 210 previously treated advanced NSCLC patients in Japan, Australia, and Europe [1]. In this large-scale international study, a similar objective tumor response rate (20%) to those of previous studies was observed. There was no difference of clinical response between patients receiving 250 mg/day and those on 500 mg/day, whereas toxicity was more severe in the higher-dose group. Subset analysis revealed startling clinico-pathological subpopulations with especially high drug sensitivity to EGFR-TKI namely Japanese patients, females, nonsmokers, and those with adenocarcinoma (Table 2). In particular, Japanese females exhibited an overall response rate >50% in this analysis. For the first time, unlike conventional anticancer agents these results suggested that EGFR-TKIs are efficacious in specific subpopulations. While that phase II trial was ongoing, two large phase III trials in untreated NSCLC were begun in the USA and Europe [2, 3]. The rationale of these two clinical trials, INTACT1 and INTACT2, was based on preclinical studies that suggested synergistic effects of taxane plus gefitinib against cancer cells in vitro and in vivo. Hence, gefitinib or placebo was added onto standard chemotherapy regimens cisplatin/gemcitabine (INTACT1) and carboplatin/paclitaxel (INTACT2) [2, 3]. Both trials showed that there was no evidence for prolonged survival time with add-on gefitinib for either standard chemotherapy schedule. The same negative result was observed in another phase III trial using the same design with erlotinib as well as gefitinib [5]. However, in this trial subset analysis suggested enhanced efficacy of EGFR-TKI therapy among nonsmokers and those harboring activated-type *EGFR* mutations. Two subsequent studies of second-line and later treatment, BR.21 and ISEL, gave conflicting results for overall survival time, time to progression, and response rate: the former suggested additional benefit of add-on EGFR-TKI and the latter gave negative results [10, 11].

Table 1 Antitumor activity of gefitinib in Japanese phase I study

	Total	PR (%)
All cases	31	5 (16)
NSCLC	23	5 (22)
Histology		
Adenocarcinoma	19	5 (26)
Squamous cell carcinoma	4	0 (0)
Gender		
Male	15	1 (7)
Female	8	4 (50)

PR partial response

To clarify the clinical benefit of EGFR-TKIs in *EGFR* mutation-positive NSCLC, prospective phase II (WJTOG0403) and phase III (WJTOG3405) studies are now underway (Fig. 1). The results of these investigations aim to give us data that will enable us better to understand *EGFR* mutational status and whether mutant EGFR phenotype confers clinical benefit in patients.

EGFR mutation and drug sensitivity

To use gefitinib effectively in clinical settings we must first identify patient populations who respond well to this agent. As mentioned above, data from IDEAL1 revealed that gefitinib is highly effective in Japanese, females, adenocarcinomatous histology, good performance status (PS), and nonsmokers (Table 2). Since the target molecule of EGFR-TKIs is EGFR, some correlation between expression patterns of EGFR protein and clinical outcome was widely speculated. However, IDEAL1 and 2 found no correlation between these parameters clinically, questioning the concept of molecular-targeting drugs. However, the answer to this question was provided by the striking findings regarding *EGFR* gene mutations [7, 9]. These *EGFR* mutations, located on the ATP binding site (exon 19–21) of

Table 2 Overall survival by patient characteristics: IDEAL1

Characteristic	Evaluable (*n*)	MST, days (95%CI)	*P*-value[a]	ORR, % (*n*)
All patients	209	241 (205–276)		18.7 (39/208)
Dose				
250 mg/day	103	232 (161–318)	0.716	18.4 (19/103)
500 mg/day	106	243 (203–309)		19.0 (20/105)
Age				
<65 years	145	238 (198–284)	0.5598	19.4 (28/144)
>65 years	64	241 (188–371)		17.2 (11/64)
Gender				
Female	61	397 (261–439)	0.0025	34.4 (21/61)
Male	148	212 (161–243)		12.2 (18/147)
WHO PS				
0–1	182	268 (234–318)	<0.0001	21.0 (38/181)
2	27	83 (57–121)		3.7 (1/27)
Histology				
Adenocarcinoma	131	300 (236–371)	<0.0001	26.0 (34/131)
Other	78	198 (129–232)		6.5 (5/77)
Smoking history				
Yes	104	186 (127–241)	<0.0001	12.5 (13/104)
No	53	414 (357–534)		37.7 (20/53)

MST mean survival time, *ORR* overall response rate

[a] Log-rank test

EGFR tyrosine kinase domain, are missense or deletion mutations causing substitution or partial deficiency of amino acid. Based on the results of basic studies, structural changes of the ATP binding site were found to increase binding affinity for ATP and gefitinib. In other words, under physiological conditions *EGFR* mutations are activating mutations that constitutively increase tyrosine kinase activity, and it is speculated that signals via EGFR are thereby abnormally enhanced and have greater impact on malignant transformation such as cancer cell proliferation. Fortunately, since these mutations are thought to have more highly augmented binding affinity for gefitinib than for ATP, they may display overwhelmingly high sensitivity induced by EGFR-TKIs. What is surprising is the correlation between frequency of *EGFR* mutations and clinical antitumor effects. We compared mutation rates and projected response rates obtained from IDEAL 1 and 2 and from 154 subjects in the clinical study in which our institute participated, and found that the *EGFR* mutation was highly correlated with clinical response (Table 3). In addition, it was reported at the American Society of Clinical Oncology (ASCO) meeting 2005 that *EGFR* gene mutation is closely related to

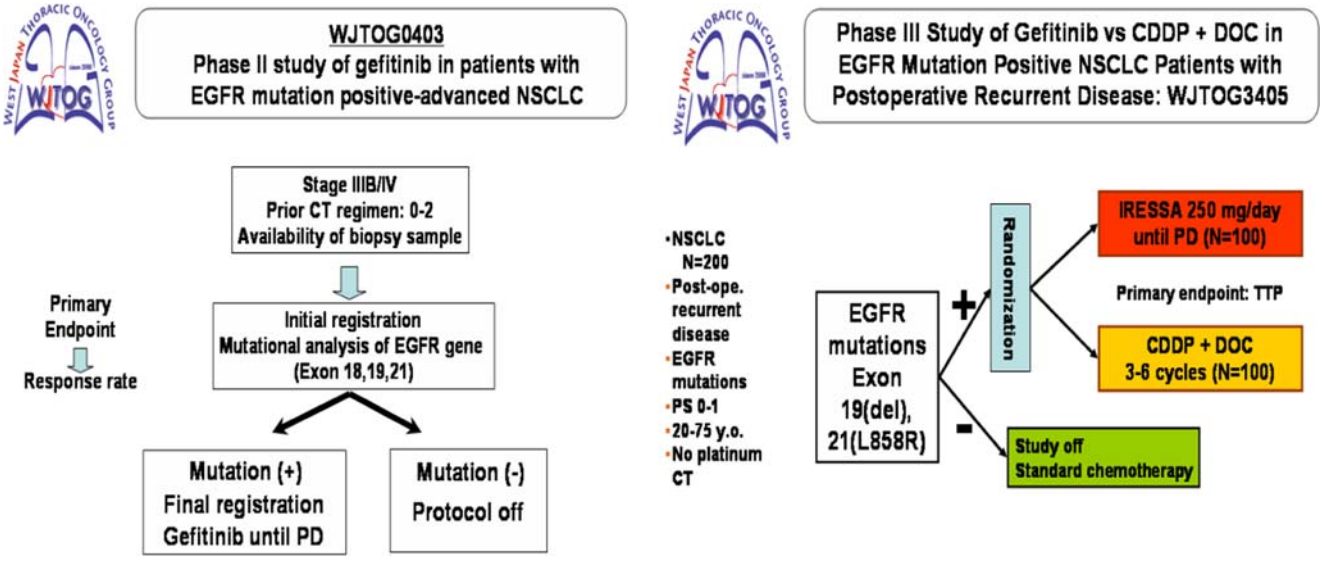

Fig. 1 Trial design of two ongoing prospective phase II (WJTOG0403) and phase III (WJTOG3405) studies investigating clinical benefit of EGFR-TKIs in *EGFR* mutation-positive NSCLC

Table 3 Estimated response rate (RR) for gefitinib and *EGFR* mutation in patients with NSCLC

Patient population	Estimated RR (%)	*EGFR* mutation (%)	
		Guillermo	Mitsudomi
Euro-American	10	2	–
Japanese	28	26	40
Japanese-adenocarcinoma	35	32	49
Female Japanese-adenocarcinoma	50	57	62

gefitinib sensitivity [6]. It is thought that the reason for the high response rate associated with Japanese race, female, adenocarcinoma, good PS, and nonsmokers is high frequency of *EGFR* mutations in these populations.

Future challenges

To establish clinical usage of EGFR-TKIs there are many issues to be addressed such as: (1) precisely identifying the site of *EGFR* mutations associated with drug sensitivity; (2) conducting a prospective clinical study of *EGFR* mutation and drug sensitivity; (3) establishing techniques to detect *EGFR* mutation precisely; (4) investigating efficacy of EGFR-TKI therapy in patients without *EGFR* mutations; (5) identifying patients responsive to EGFR-TKIs among those without *EGFR* mutations and clarifying the mechanism of action of EGFR-TKIs; and (6) clarifying mechanisms of EGFR-TKI resistance and developing drugs to overcome this resistance.

Combined use with conventional anticancer agents

Currently, gefitinib is the only EGFR-TKI available in Japan. How should we use gefitinib in combination with other anticancer agents? Large-scale clinical studies in Caucasian NSCLC patients indubitably have shown that concomitant use of conventional anticancer agents and gefitinib has no clinical usefulness in that patient population. Considering the association between gefitinib sensitivity and *EGFR* gene mutations, however, it seems too early to make a similar conclusion in Japanese patients in whom *EGFR* gene mutations might be more frequent. Therefore, it is important clinically to test gefitinib in Japanese patients concomitantly taking conventional anticancer drugs. In addition, in the context of combination therapeutic regimens not only simultaneous administration with conventional anticancer agents but sequential and maintenance therapies should be evaluated. To this end, the WJTOG phase III clinical trial is currently ongoing. Patients enrolled in this trial are divided into two groups: those taking three courses of two chemotherapeutic agents including one platinum-based drug followed by three courses of gefitinib, and the group on six courses of two drugs including one platinum drug alone. This trial, expected to terminate in April 2005, is aimed to show conclusively whether serial/sequential gefitinib therapy is useful in Japanese patients with NSCLC.

Conclusions

The advent of EGFR-TKIs convinced us that biological study of these agents in NSCLC could improve prognosis of these patients. Although the improvement elicited by gefitinib may be small so far, this agent does at least provide a new form of therapy that over the short term leads to markedly reduced tumor size without bone marrow suppression including neutropenia and no severe nausea and vomiting even in those patients with the worst prognosis. This raises the possibility of placing this rapidly fatal disease under some control. Doctors must be aware that making inroads towards understanding the implications of gene mutation in lung cancer will be a milestone for new therapeutics.

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