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CASE REPORT

Acquisition of T790M resistance mutation in a patient with advanced adenocarcinoma harboring uncommon *EGFR* mutations: a case report and literature review

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Department of Thoracic Oncology and Respiratory Medicine, Tokyo Metropolitan Cancer and Infectious Diseases Center Komagome Hospital, Tokyo, Japan **Background:** Osimertinib is the standard of care for patients with non-small-cell lung cancer (NSCLC) patients harboring acquired *EGFR* T790M resistance mutation. However, the frequency of T790M resistance mutation acquisition and the efficacy of osimertinib in patients harboring uncommon *EGFR* mutations, which accounts for approximately 10% of *EGFR* mutations, remain unclear.

Case presentation: We present the case of a 72-year-old Japanese woman with recurrent NSCLC harboring uncommon *EGFR* mutations, who was subsequently found to have acquired the T790M resistance mutation and was treated with osimertinib. She died 9 days later because of worsening respiratory failure with disease progression.

Conclusion: The findings of the present case suggest that the T790M resistance mutation can occur even when the patient harbors an uncommon *EGFR* mutation after *EGFR*-tyrosine kinase inhibitor treatment, and the prognosis could be poor despite the presence of an acquired T790M resistance mutation.

Keywords: non-small-cell lung cancer, *EGFR*-TKIs, osimertinib, uncommon *EGFR* mutation, acquired resistance

Introduction

EGFR mutations are some of the major mutations in non-small-cell lung cancer (NSCLC). Exon 19 deletion and exon 21 L858R point mutation, which are referred to as common *EGFR* mutations, are activating *EGFR* mutations associated with *EGFR*-tyrosine kinase inhibitors (TKIs). Approximately 10% of *EGFR* mutations represent uncommon *EGFR* mutations.¹ First-generation *EGFR*-TKIs show insufficient clinical benefit for uncommon *EGFR* mutations;² however, second-generation *EGFR*-TKIs, such as afatinib, show a 60% response rate.³ Because prospective data on survival are absent and uncommon *EGFR* mutations are heterogeneous, there is no established standard of care (SOC) for patients harboring uncommon *EGFR* mutations.

Acquired resistance to *EGFR*-TKIs develops after a median of 10-12 months, and it is most commonly mediated by an *EGFR* exon 20 T790M mutation. The AURA 3 Phase III trial demonstrated the superiority of osimertinib over platinum therapy plus pemetrexed, which had been the SOC for patients with NSCLC harboring acquired resistance to prior *EGFR*-TKIs.⁴ However, the frequency of T790M resistance mutation acquisition after treatment with *EGFR*-TKIs for uncommon *EGFR* mutations remains

Correspondence: Makiko Yomota Department of Thoracic Oncology and Respiratory Medicine, Tokyo Metropolitan Cancer and Infectious Diseases Center Komagome Hospital, 3-18-22 Honkomagome, Bunkyo, Tokyo 113-8677, Japan Tel +81 3 3823 2101 Email myomota@gmail.com



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© 2019 Hakozaki and formota. This work is published and licensed by Dove Medical Press Limited. The full terms of this license are available at https://www.dovepress.com/terms.php hereby accept the Terms. Non-commercial uses of the work are permitted without any further permission from Dove Medical Press Limited, provided the work is properly attributed. For permission for commercial use of this work, please see paragraph 4.2 and 5 of our Terms (http://www.dovepress.com/terms.php). unclear, although 50%–60% of cases of *EGFR* common mutations, including exon 19 deletion and exon 21 L858R mutation, acquire the T790M resistance mutation. Under these circumstances, the optimal treatment approach for patients harboring uncommon *EGFR* mutations remains unclear.

Herein, we present the case of a patient with recurrent NSCLC harboring uncommon *EGFR* mutations, who was subsequently found to have acquired the T790M resistance mutation and was treated with osimertinib.

Case presentation

A 72-year-old Japanese woman presented with abnormal chest opacity at an annual health checkup. She was a nonsmoker and had no specific medical history. Her Eastern Cooperative Oncology Group (ECOG) performance status was zero. Chest computed tomography (CT) revealed a pulmonary nodule measuring 2.5×1.6 cm in the left lower lobe. She underwent left lower lobectomy and systemic lymph node dissection. Based on the analysis of the nodule, she was diagnosed with adenocarcinoma (pT2aN2M0 Stage IIIA) harboring an EGFR exon 18 G719X mutation. She underwent four cycles of adjuvant chemotherapy with cisplatin plus vinorelbine. Recurrence with multiple intrapulmonary metastases and malignant pleural effusion were observed later. Genetic analysis of the pleural effusion at that time showed an EGFR exon 18 G719X mutation, as previously detected. Subsequently, she was treated with gefitinib for 15 months, and then, increased pleural effusion and carcinomatous lymphangiomatosis were noted. Both EGFR exon 18 G719X and exon 20 T790M mutations were detected in her plasma (CobasTM EGFR Mutation Test v2, Hoffman-La Roche Ltd., Basel, Switzerland). Her ECOG performance status declined to 2 because of worsening dyspnea, and she was treated with osimertinib. She died 9 days later from worsening respiratory failure with disease progression.

Discussion and conclusion

Tumor genotyping for an acquired T790M resistance mutation at disease progression has become a standard component of care in patients with NSCLC harboring *EGFR* mutations to guide subsequent treatment. Previous studies have shown that approximately 50–60% of patients treated with first or second-generation *EGFR*-TKIs acquire the T790M resistance mutation at the time of progressive disease (PD). However, in these studies, almost all patients had common *EGFR* mutations, such as exon 19 deletion and exon 21 L858R mutation. Therefore, the prevalence of T790M resistance mutation acquisition in patients harboring uncommon *EGFR* mutations and the efficacy of third-generation *EGFR*-TKIs in this population remain unclear.

In the AURA Phase I/II trial, osimertinib was administered to 222 patients with NSCLC harboring both T790M-mediated and non-T790M-mediated resistance to *EGFR*-TKIs in the expansion cohort. Of these 222 patients, 10 (4.5%) initially had uncommon *EGFR* mutations, and 4 of these 10 patients (40%) acquired the T790M resistance mutation.⁵ In a study of 125 patients who were re-biopsied at disease progression with initial *EGFR*-TKIs (mostly first-generation therapy), only 6 (5%) initially had uncommon mutations, and only 1 of these 6 patients (17%) acquired the T790M resistance mutation.⁶ In another study on 37 patients who were treated with afatinib, 3 (8%) initially had uncommon *EGFR* mutations, and none of these 3 patients (0%) acquired the T790M resistance mutation at the time of PD (Table 1).⁷ These results may indicate that patients with NSCLC harboring uncommon *EGFR* mutations

Table I Patients with uncommon EGFR mutations who weresubsequently treated with osimertinib in previous studies

	EGFR mutation status at the	T790M	Best
	baseline	status	objective
		at PD	response
AURA (N=222)			
Ι	Exon 18 G719X	(+)	PR
2	Exon 18 G719X	(+)	SD
3	Exon 18 G719X	(+)	SD
4	Exon 18 G719X+exon 20 S7681	(+)	PR
5	Exon 18 G719X	(-)	SD
6	Exon 18 G719X	(-)	PD
7	Exon 18 G719X	(-)	PD
8	Exon 18 G719X	(-)	PD
9	Exon 18 G719X+exon 20 S7681	(-)	SD
10	Exon 20 insertion	(-)	PD
Kawamura et al ⁷ (N=I25)			
Ι	Uncommon EGFR mutation ^a	(+)	NA
2	Uncommon EGFR mutation ^a	(-)	NA
3	Uncommon EGFR mutation ^a	(-)	NA
4	Uncommon EGFR mutation ^a	(-)	NA
5	Uncommon EGFR mutation ^a	(-)	NA
6	Uncommon EGFR mutation ^a	(-)	NA
Tanaka et al ⁶ (N=37)			
I	Exon 18 ^b	(-)	NA
2	Exon 20 ^b	(-)	NA
3	Exon 20 ^b	(-)	NA
Notes: "Exon 18 G719X, exon 20 insertion, or exon 21 L861Q. "Details were not			

Notes: ^aExon 18 G719X, exon 20 insertion, or exon 21 L861Q. ^bDetails were not provided.

Abbreviations: PR, partial response; SD, stable disease; PD, progressive disease; NA, not available.

are less likely to acquire the T790M resistance mutation when compared with those who have common *EGFR* mutations (50%–60%). In the AURA 2 Phase II (N=210) and AURA 3 Phase III (N=419) trials, of all participants who had acquired T790M resistance, 8 of 210 (4%) and 11 of 419 (3%) participants initially had uncommon *EGFR* mutations.^{4,8} Considering the prevalence of uncommon *EGFR* mutations among all *EGFR* mutations (approximately 10%), the proportion of patients harboring uncommon *EGFR* mutations in these trials was less. This may reflect the rarity of T790M resistance acquisition among patients harboring uncommon *EGFR* mutations.

In addition, there are limited data on the efficacy of osimertinib among patients with T790M resistance acquisition who previously had uncommon EGFR mutations besides the 4 patients in the AURA trial. Of these 4 patients, 2 showed partial response (PR) and 2 showed stable disease (SD) as the best objective response (Table 1). In contrast, in the present case, osimertinib was found to be inactive with rapid disease progression. We cannot dismiss the poor performance status of the patient at the time of osimertinib initiation; however, her clinical course may have reflected characteristics of the rare EGFR mutation status. The contribution of mechanisms of acquired resistance other than T790M mutation may have been substantial, and cytotoxic chemotherapy could be a choice of treatment if the general condition permits. Prior studies and the present case describe some characteristics of patients with T790M resistance mutation acquisition who previously had uncommon EGFR mutations. However, further cases should be accumulated to obtain practical clinical information that can guide optimal treatment. In addition, it may be reasonable to utilize comprehensive methods, such as next-generation sequencing, particularly in small patient subgroups for which conformational clinical trials may not be feasible.

The findings of the present case indicate that the T790M resistance mutation can occur even when the patient harbors an uncommon *EGFR* mutation after *EGFR*-TKI treatment and that the prognosis could be poor despite the presence of an acquired T790M resistance mutation. Further understanding of the clinical characteristics may help in the optimization of strategies to provide the maximum clinical benefit in patients with NSCLC harboring uncommon *EGFR* mutations.

Ethics approval and informed consent

The case report was waivered by the Ethics Committee of Tokyo Metropolitan Cancer and Infectious diseases Center Komagome Hospital. The clinical information presented in this case report was obtained through Tokyo Metropolitan Cancer and Infectious Diseases Center Komagome Hospital's medical records.

Consent for publication

Written informed consent was obtained from the patient for publication of this case report.

Data availability

All relevant data are within the manuscript.

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Author contributions

Both authors contributed to data analysis, drafting and revising the article, gave final approval of the version to be published, and agree to be accountable for all aspects of the work.

Disclosure

The authors report no conflicts of interest in this work.

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