Primary Resistance to Brigatinib in a Patient with Lung Adenocarcinoma Harboring ALK G1202R Mutation and LIPI-NTRK1 Rearrangement

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Purpose: Anaplastic lymphoma kinase (ALK) inhibitors have transformed the management of non-small-cell lung cancer (NSCLC) patients with ALK gene rearrangement. This paper reports a new resistance mechanism to a second-generation ALK inhibitor, brigatinib.

Case Report: A 43-year-old woman who had no history of smoking was diagnosed with stage IVa (T2bN2M1b) lung adenocarcinoma. After the first-line chemotherapy failed, the patient received crizotinib due to the presence of EML4-ALK fusion by next-generation sequencing (NGS). The patient had disease progression after 8 months on crizotinib, and a second NGS identified the ALK G1202R resistance mutation. Therefore, she was switched to brigatinib. After only 53 days of treatment with brigatinib, the patient developed a new 1.6×1.2 cm lesion in the mediastinal lymph node. A third NGS testing revealed a new form of NTRK rearrangement (LIPI-NTRK1). The patient died 16 months after diagnosis.

Conclusion: This paper provides new insights into the primary resistance to brigatinib in NSCLC patients carrying ALK G1202R mutation. The new fusion form of NTRK rearrangement was detected, which may provide potential treatment options after brigatinib resistance.

Keywords: NTRK1, ALK, primary resistance, brigatinib, NSCLC

Introduction

Anaplastic lymphoma kinase (ALK) rearrangements are present in approximately 5% of non-small-cell lung cancer (NSCLC) patients, defining a specific molecular subgroup.1 The ALK gene copy number gain and amplification (ALK-A) are well studied in different types of cancer, and could affect the rearrange of the ALK gene in NSCLC.2 Crizotinib is effective as the first-in-class inhibitor in ALK-positive NSCLC,3 but most patients acquire resistance against ALK inhibitors through different molecular mechanisms,4 including the co-occurrence of ALK-A and ALK rearrangement.2 Brigatinib is a next-generation oral ALK inhibitor to treat metastatic ALK+ NSCLC patients who have progressed on or are intolerant of crizotinib.5 Efficacy data confirmed thatbrigatinib is an effective therapeutic strategy after crizotinib failure.5,6 Previous reports showed that brigatinib can overcome resistance to first- and second-generation ALK inhibitors caused by secondary mutations such as ALK G1202R.7,8 According to the preliminary data of the Phase 2 ALTA trial, the investigator-assessed median progression-free survival (PFS) was 12.9 months in patients treated with brigatinib.9 However, there are some conflicting views on the resistance mechanisms...
of brigatinib. Sharma et al reported that the G1202R mutation might cause acquired resistance to brigatinib. This may result from the steric clash between the side chain of G1202R and the extended solubilization group of brigatinib.\(^{10}\)

Chromosomal rearrangements involving neurotrophic tyrosine kinase 1 (NTRK1) occur at an extremely low frequency of approximately 0.1% to 1% in NSCLC and they are highly therapeutically actionable drivers of tumor growth.\(^{11,12}\) Entrectinib is an orally available small molecule for the treatment of various solid tumors harboring NTRK1/NTRK2/NTRK3 or ALK gene fusions.\(^{13}\) Despite durable responses to TRK-directed therapy in patients with NTRK-rearranged tumors, it is expected that most patients eventually develop acquired resistance. Moreover, NTRK-rearranged NSCLCs seem to be oncogene-dependent and not combined with ALK or ROSI gene rearrangements.\(^{14,15}\)

We here reported a case of lung adenocarcinoma carrying the G1202R ALK mutation and a new oncogenic NTRK fusion variant who was resistant to brigatinib treatment.

**Case Presentation**

A 43-year-old female never-smoker presented with prolonged paroxysmal cough and was diagnosed with stage IVa (T2bN2M1b) lung adenocarcinoma in Jun 2017. She underwent an endobronchial ultrasound-guided transbronchial needle aspiration (EBUS-TBNA) and immunohistochemistry showed that the nucleoli had obvious heterotypic cells arranged like adenoid and CK (+), TTF-1 (+), NapsinA (+), P63 (+). Enhanced computed tomography (CT) revealed an upper left lung lesion (3.6×2.9 cm) with mediastinal lymph node metastasis (2.1×1.0 cm) and hepatic S4 segment metastasis (1.0×0.9 cm). She received two courses of docetaxel combined with cisplatin chemotherapy as the first-line treatment, and CT revealed no change in the lesion (Figure 1). To explore potential targeted treatment, next-generation sequencing (NGS) analysis was performed on the patient’s peripheral blood using a 21 gene panel. The patient was found to carry the classical EML4-ALK fusion. Therefore, crizotinib was commenced at 250 mg bid on September 5th 2017. A follow-up CT conducted on January 17th, 2018 revealed a 61% regression in her primary lung lesion (1.4×1.2 cm), indicating that the patient had achieved partial response (PR). In May 2018, eight months after the onset of crizotinib treatment, the patient was discovered to have tumor progression (PD) due to brain metastases (1.9×1.6 cm) by head MRI and acquired resistance to crizotinib was suspected.

A second blood-based NGS showed the presence of the p. G1202R ALK mutation was observed. The patient was started on brigatinib (180 mg daily with a seven-day lead-in at 90 mg) on May 18\(^{th}\), 2018. Brigatinib is a next-generation oral ALK inhibitor used in the second-line after progression on crizotinib. However, a CT scan conducted after 53 days of brigatinib treatment revealed a new mediastinal lymph node (1.6×1.2 cm), and the appearance of new pericardial metastases. A third NGS testing was therefore performed, and a new type of NTRK arrangement (LIPI-NTRK1, Figure 2) was identified in addition to the two previous alternations. The patient’s shortness of breath significantly increased due to the hydrothorax on the left side of the chest. After a pleural puncture, the symptoms were slightly relieved. Given her physical conditions, the patient refused to switch to another regimen, including NTRK inhibitors. She was able to benefit from ALK TKI therapy for 8 months and died on September 24\(^{th}\), 2018, with overall survival of 16 months from the time of diagnosis.

**Discussion**

In this case report, a new form of NTRK rearrangement (LIPI-NTRK1) was identified, which included inversion of LIPI exon 1 and NTRK1 exons 8–17. In addition, the classical EML4-ALK fusion and the ALK G1202R primary resistance mutation were also detected. Despite multiple lines of targeted treatment guided by NGS testing, this patient failed to benefit from the treatment of brigatinib due to the emergence of resistance mutations.

ALK-TKIs are widely used in clinical practice, but patients’ responses are heterogeneous due to the emergence of resistance genes. Numerous researchers have explored the mechanisms of primary resistance to ALK-TKIs for ALK+ NSCLC patients.\(^{3,4}\) The G1202R mutation is reported to be one of the common resistance mechanisms to first- and second-generation ALK inhibitors (crizotinib, alectinib, and ceritinib).\(^{16}\) Interestingly, brigatinib is a highly potent and selective ALK inhibitor, and it maintained substantial activity against all 17 secondary ALK mutants tested in cellular assays and exhibited a superior inhibitory profile compared with crizotinib, ceritinib, and alectinib at clinically achievable concentrations. It has been reported to have potent effects on the refractory G1202R mutant in vitro and in vivo.\(^{8,17}\) Brigatinib has been demonstrated to produce a 12.8 month PFS and a 64% intracranial ORR in crizotinib-refractory patients. However, the patient discussed here did not benefit from brigatinib. Based on the patient’s NGS results, it is possible that the G1202R mutation caused primary resistance to brigatinib.
After brigatinib failure, the patient underwent another NGS testing, and the result showed \textit{LIPI-NTRK1} fusion, a new type of NTRK rearrangement, suggesting that this patient developed a potential treatment to NTRK inhibitors. However, the \textit{ALK} G1202R mutation is analogous to resistance mutations that affect the kinase solvent front and will directly interfere with binding with entrectinib and other TKIs with TRK activity. Functional studies have subsequently confirmed that cancer cells harboring these mutations are cross-resistant to all TKIs with anti-TRK activity. As a result, we speculate that both brigatinib and NRTK inhibitors have limited efficacy in this patient due to the G1202R mutation. Meanwhile, given her
poor physical conditions, switching to an NRTK inhibitor or other targeted drugs is not a viable option for this patient.

In summary, we reported a case of ALK-positive NSCLC with primary resistance to brigatinib due to the G1202R mutation. This is also the first report of the LIPI-NTRK1 fusion variant. Although verification of the fusion with fluorescent in situ hybridization or immunohistochemistry was not an option for us due to the lack of tissue, our case provided useful information to better understand the primary resistance to ALK-TKIs. In addition, NGS is valuable to explore the comprehensive genetic characteristics and potential drug targets for patients who experience primary resistance to ALK inhibitors.

Abbreviations
ALK, Anaplastic lymphoma kinase; NSCLC, non-small-cell lung cancer; ALK-A, ALK amplification; PFS, progression-free survival; NTRK1, neurotrophic tyrosine kinase 1; EBUS-TBNA, endobronchial ultrasound-guided transbronchial needle aspiration; NGS, next generation sequencing; PR, partial response; PD, progression disease.

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Author Contributions
Lizhu Lin contributed to the concept and design and interpretation of data. All authors contributed to acquisition, analysis, and interpretation of data. Zhiwei Xiao and Xuewu Huang drafted the article as the co-first authors. All authors contributed equally in drafting and revising the article and 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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Figure 2 A new fusion form of NTRK rearrangement involving inversion of the LIPI gene (exons 1) and the NTRK gene (exons 8-17) was identified.
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