A Patient of Advanced NSCLC with a New EGFR Exon 19 Insertion Mutation and its Response to EGFR-TKIs

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ABSTRACT

Epidermal growth factor receptor tyrosine kinase inhibitors (EGFR-TKIs) are the standard therapy for patients with advanced non-small-cell lung cancer (NSCLC) harbouring common EGFR mutations. However, about 10% of EGFR mutations are uncommon mutations and their response to EGFR-TKIs remains unclear. The present case reports a 75-year, female patient with advanced NSCLC, presenting with a new subtype of EGFR exon 19 insertion mutation (IPVAIL insertion), who showed obvious symptom improvement after EGFR-TKIs treatment but a relatively short time of progression-free survival (PFS) and succumbed to tumor 133 days (4.4 months) after diagnosis. In conclusion, patients harbouring new subtype of EGFR exon 19 insertion mutations, IPVAIL insertion may have a poor prognosis. Further experiences are required to characterise these uncommon mutations.

Key Words: Gefitinib, afatinib, exon 19 mutation, non-small cell lung cancer, epidermal growth factor receptor.

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INTRODUCTION

Non-small-cell lung cancer (NSCLC) is the most common type of all lung cancers (about 85%).¹ Activating EGFR mutations are main oncogenic drivers in NSCLC. With the application of the next-generation sequencing (NGS), more and more uncommon EGFR mutations are being detected. However, EGFR exon 19 insertions, first reported by He *et al.* in 2012, are very rare (about 0.5% of all EGFR mutations).² Their sensitivity to EGFR-TKIs in patients with NSCLC is uncertain. Here, we report a case of a patient with advanced NSCLC harbouring IPVAIL insertion, a new subtype never reported before and its response to EGFR-TKIs.

CASE REPORT

In August 2017, a 75-year Chinese female patient presented at our hospital with a one-year history of repeated epigastria discomfort and pain in her lower back. The symptoms of appetite and weight loss aggravated in recent three months. The eastern cooperative oncology group (ECOG) performance status score of the patient was one.³ A serum tumor markers test panel was used for tumor screening and showed the level of cancer

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Received: February 26, 2019; Revised: July 08, 2019; Accepted: July 20, 2019 antigen-199 (CA-199), cancer antigen-153 (CA-153), cancer antigen-125 (CA-125) and carcinoembryonic antigen (CEA) obviously elevated. The levels were 279.7 unit/ml (normal level <37 unit/ml), 122.2 unit/ml (normal level: <30 unit/ml), 75.53 unit/ml (normal level: <35 unit/ml) and 161.97 nanogram/ml (normal level: <5 nanogram/ml), respectively. Gastroscopy suggested gastric ulcer, and colonoscopy was negative. Enhanced computed tomography (CT) of abdomen and pelvic cavity revealed multiple patchy high-density shadows in centrum, pelvis and proximal femur. Positron emission tomography/ computed tomography (PET/CT) (Figure 1) suggested extensive bone destruction with increased radioactive intake and maximum standard uptake value (SUV) of 14.1. It also showed a mass with a slight increase in radioactive intake (SUV of 3.3) partially obstructing the bronchial cavity in the lower lobe of left lung. Although CA-199 and CA-125 were elevated in this patient, enhanced CT of abdomen and PET/CT showed no evidence of pancreatic/ovarian metastasis or primary pancreatic/ovarian cancers. Chest CT enhancement scan (Figure 2A) revealed blocked bronchial cavity in the lower lobe of left lung with pulmonary atelectasis, and a small amount of effusion in the left thoracic cavity. The patient refused to undergo bronchoscopy. We performed CT-guided biopsy of the right ilium and sacrum. Pathological results showed metastatic adenocarcinoma. Combined with immuno-histochemicalo results, CK (AE1/AE3) +++, CK7 +++, CK20 -, CDX2 -, TTF-1 +++, NapsinA +++, Ki-67 30%+, P53 +), the patient was eventually diagnosed with lung adenocarcinoma with multiple bone metastasis. NGS of 168 cancer-related



Figure 1: Positron emission tomography/computed tomography (PET/CT).



Figure 2: (A) Chest contrast enhanced CT before afatinib; **(B)** Thoracic high-resolution CT after afatinib; **(C)** Thoracic high-resolution CT after gefitinib.

 Table I:
 EGFR mutation subtype occurring in exon 18-21 and their frequency

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EGFR mutation subtype	Frequency (%)
Common mutations	
Exon 19 deletion	48
Exon 21 point mutation (L858R)	43
Uncommon mutations	
Exon 18 point mutation (G719X, E709X)	3-4
Exon 19 insertion	0.5
Exon 20 insertion	10 (1-17)
Exon 20 point mutation (S768I)	1
Exon 20 point mutation (T790M)	1 (inherited)
Exon 21 point mutation (L861Q)	2

EGFR: Epidermal growth factor receptor.

genes (Burning Rock Biotech, Guangzhou, People's Republic of China) was performed on DNA derived from the biopsy specimen and circulating tumor DNA (ctDNA) derived from whole blood sample. Both showed the patient harboured a mutation (NM_005228.3, c.2217_2234dup) in the EGFR gene exon 19, resulting in a six amino acids insertion (p.Lys745_ Glu746insIleProValAlalleLys). She took afatinib at a starting dose of 40 mg/day from September 10, 2017 and showed great improvement in her pain and other symptoms.

However, about one month later, thoracic high-resolution scanning CT (Figure 2B) revealed solid mass beside the left parahilar enlarged and the left pleural effusion increased compared to the images taking before afatinib. The patient presented symptoms of nausea, vomiting, asthenia and anorexia. Limited effectiveness of afatinib was observed in our patient. We recommended the patient for the genetic test of EGFR T790M. However, because of the fact that the third-generation TKI (AZD9291) was not been included in the coverage of the Chinese healthcare insurance system in October, 2017, patients ought to pay about RMB 50,000 (about 7,500 dollars) a month for taking the third-generation TKI. This was an unaffordable price for our patient. Thus, the patient refused to take genetic test. Moreover, the patient also refused to take chemotherapy. Taking into account the economic situation of the patient and effectiveness of gefitinib in patients with EGFR exon 19 insertion mutation of previous literature, we decided to try the first generation TKI, which had already been included in the coverage of the Chinese healthcare insurance system and the cost of medicine was only about RMB 1,000 (150 dollars) per month. So gefitinib was started at a standard dose of 250 mg/day from October 16, 2017. Thoracic high-resolution scanning CT on October 31, 2017 (Figure 2C) showed that the left pleural effusion was obviously resolved. From November 25, 2017, the patient began to be unconscious, incontinent of urine and feces. Finally, the patient died on December 20, 2017 and her overall survival (OS) was 133 days (4.4 months).

DISCUSSION

Activating EGFR mutations are main oncogenic drivers in NSCLC. In Asian NSCLC patients, EGFR mutations are detected in about 40-60% of cases.⁴ EGFR mutations can be divided into two categories: common mutations and uncommon mutation. Exon 19 deletions and exon 21 point mutation (L858R) are the two most common mutations, which also called classical mutations. They account for approximately 48% and 43% of all EGFR mutations, respectively. The remaining 10% of EGFR mutations are reported to occur in exon 18-21 and their subtype and frequency are listed in Table I.⁵⁻⁷ However, due to the complexity of uncommon mutations and the lack of large cohort studies, treatment aiming at EGFR uncommon mutations still remains a big challenge.

Exon 19 insertions are one subtype of the rare EGFR mutations. Park *et al.* reported two of three cases with exon 19 insertions responding to gefitinib, but time to progression (TTP) was relatively short (mean, 5.9 months).⁸ lyevleva *et al.* reported three patients with exon 19 insertions receiving gefitinib and TTP were 5 months, 9 months, and 11 months.⁹ Lin *et al.* also reported three patients received EGFR-TKIs (two gefitinib, one erlotinib) and their progression-free survival (PFS) were 5 months, 22 months and 2 months, respectively.¹⁰ He *et al.* reported three of four patients harbouring exon 19 insertions sensitive to erlotinib or afatinib (TTP: 14 months, 19 months and 50 months) and they suggested that patients with exon 19 insertions should be treated with EGFR-TKs.²

In our patient, we found an 18-base pair insertion between the 2217th and 2234th nucleotides in EGFR exon 19, which lead to a six residue insertion in the protein structure between codon 745 and codon 746 (p.Lys745_Glu746insIleProValAlalleLys). According to the published literature, all the reported EGFR exon 19 insertion subtypes are non-frame shift mutations. Except for one patient, who had 15-base pair insertion, to all other had 18-base pair insertion.^{2,10} In addition, existing literature shows insertion types to include KIPVAI, IPVAIK, VPVAIK and TPVAIK2, whereas our patient had IPVAIL insertion, a new type never reported before. Its response to EGFR-TKIs treatment was less than satisfactory. Our patient's overall survival was only 133 days (4.4 months).

Despite the limitations of a case report, the present case demonstrates that patients harbouring this rare subtype of EGFR exon 19 insertion mutation may have a poor prognosis. Further experiences are required to characterise these rare mutations.

PATIENT'S CONSENT:

Written informed consents for publication of this case report and accompanying images were obtained both from the patient during her lifetime and from the patient's daughter after the patient died.

CONFLICT OF INTEREST:

Authors declared no conflict of interest.

AUTHORS' CONTRIBUTION:

NZ: Contributed to the conception of the work, the acquisition of data for the work and the writing of the manuscript.

CD: Contributed to the design of the work, the interpretation of the results and the revising the manuscript.

SW: Contributed to the interpretation of the results and the revising the manuscript.

YY: Contributed to the conception of the work, the interpretation of results for the work and the revising the manuscript.

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