



## OPEN ACCESS

## EDITED BY

Mehmet Altan,  
University of Texas MD Anderson  
Cancer Center, United States

## REVIEWED BY

Ilaria Attili,  
European Institute of Oncology (IEO),  
Italy  
Jinjuan Yao,  
Memorial Sloan Kettering Cancer  
Center, United States

## \*CORRESPONDENCE

Xiaoyan Zhang  
xiaoyan.zhang@genetronhealth.com  
Tonghui Ma  
tonghuima0818@sina.com  
Cuiying Zhang  
cenyao\_2006@126.com

†These authors have contributed  
equally to this work and share first  
authorship

## SPECIALTY SECTION

This article was submitted to  
Precision Medicine,  
a section of the journal  
Frontiers in Medicine

RECEIVED 27 June 2022

ACCEPTED 14 September 2022

PUBLISHED 06 October 2022

## CITATION

Yang H, Li H, Fang Y, Li Z, Zhu J, Liu H,  
Lu C, Zhang X, Ma T and Zhang C  
(2022) A non-functional 5' *ALK* fusion  
validated at the RNA level as a classical  
*EML4-ALK* that responds well to the  
novel *ALK* inhibitor ensartinib: A case  
report.  
*Front. Med.* 9:979032.  
doi: 10.3389/fmed.2022.979032

## COPYRIGHT

© 2022 Yang, Li, Fang, Li, Zhu, Liu, Lu,  
Zhang, Ma and Zhang. This is an  
open-access article distributed under  
the terms of the [Creative Commons  
Attribution License \(CC BY\)](https://creativecommons.org/licenses/by/4.0/). The use,  
distribution or reproduction in other  
forums is permitted, provided the  
original author(s) and the copyright  
owner(s) are credited and that the  
original publication in this journal is  
cited, in accordance with accepted  
academic practice. No use, distribution  
or reproduction is permitted which  
does not comply with these terms.

# A non-functional 5' *ALK* fusion validated at the RNA level as a classical *EML4-ALK* that responds well to the novel *ALK* inhibitor ensartinib: A case report

Hong Yang<sup>1†</sup>, Haojing Li<sup>1†</sup>, Yu Fang<sup>2†</sup>, Zhijun Li<sup>1</sup>, Jianhua Zhu<sup>2</sup>, Huan Liu<sup>3</sup>, Chao Lu<sup>3</sup>, Xiaoyan Zhang<sup>2\*</sup>, Tonghui Ma<sup>2\*</sup> and Cuiying Zhang<sup>1\*</sup>

<sup>1</sup>Department of Oncology, Inner Mongolia People's Hospital, Hohhot, China, <sup>2</sup>Department of Translational Medicine, Genetron Health (Beijing) Technology, Co., Ltd., Beijing, China, <sup>3</sup>Department of Oncology, Inner Mongolia Medical University, Hohhot, China

**Background:** Currently, many targeted drugs are approved for treatment of *ALK* fusion non-small cell lung cancer. However, it has been previously assumed that patients with 5' non-oncogenic kinase (5' NOK) fusion detected by DNA next-generation sequencing (NGS) would not benefit from *ALK* inhibitors because of lack of an intact kinase domain.

**Case description:** A novel 5' NOK fusion form, *ALK-CYP27C1* (A19:C5), was detected by DNA NGS in surgical tissue specimens of a patient with recurrent lung adenocarcinoma. The patient achieved 29 months of progression-free survival with ensartinib treatment. The results of RNA NGS from the same operative tissue identified *EML4-ALK* (E13:A20) fusion variant type I.

**Conclusion:** This is the first case to provide real-world evidence of effective treatment of a patient with the 5' NOK fusion form at the DNA level but functional *EML4-ALK* at the RNA level, illustrating the need for RNA testing in 5' NOK patients.

## KEYWORDS

5' NOK fusion, *ALK*, RNA NGS, ensartinib, NSCLC

## Introduction

Lung cancer is one of the most common cancer types in China and is the leading cause of cancer death (1). With the development of targeted drugs, an increasing number of patients whose tumors harbor driver mutations, especially fusions, can receive targeted therapy. *ALK* fusion is an important driver mutation that accounts for approximately 3–7% of non-small cell lung cancer (NSCLC) cases (2–4), and the most common form is *EML4-ALK* (5, 6). Conventional diagnostic strategies for fusion include immunohistochemistry (IHC) and fluorescence *in situ* hybridization (FISH); however, both are low-throughput (7). Additionally, a previous report has shown that the false-negative rate of FISH and IHC is 24 and 12.6%, respectively, which may lead to patients not receiving *ALK* inhibitor treatment in time (8). A total of 303 patients with *ALK* IHC-positive NSCLC were reevaluated in the ALEX trial study, of which 203 were FISH-positive, 39 were FISH-negative, and 61 were FISH-uninformative, and the corresponding ORR was 90.6, 28.6, and 96%, respectively (9). This suggested that the results were differences between DNA level (FISH) and protein level (IHC) in fusion detection, and that the patients with *ALK* IHC-positive and FISH-negative or *ALK* IHC-positive and FISH-uninformative results could also benefit from *ALK* inhibitors.

NGS has been recommended by the National Comprehensive Cancer Network (NCCN) guidelines for genetic testing. Because of the high throughput of DNA NGS detection techniques, driver mutations are more easily detected with these approaches than by FISH or IHC. Meanwhile, more fusion forms of the *ALK* gene have been identified, including novel partners and breakpoints, some of which may be transcribed to non-functional or other fusion forms at the RNA level (10–12). 5' NOK fusion is a novel form at the DNA level that does not have an intact kinase domain; therefore, it has been suggested that patients with such a fusion cannot benefit from *ALK* inhibitor treatment.

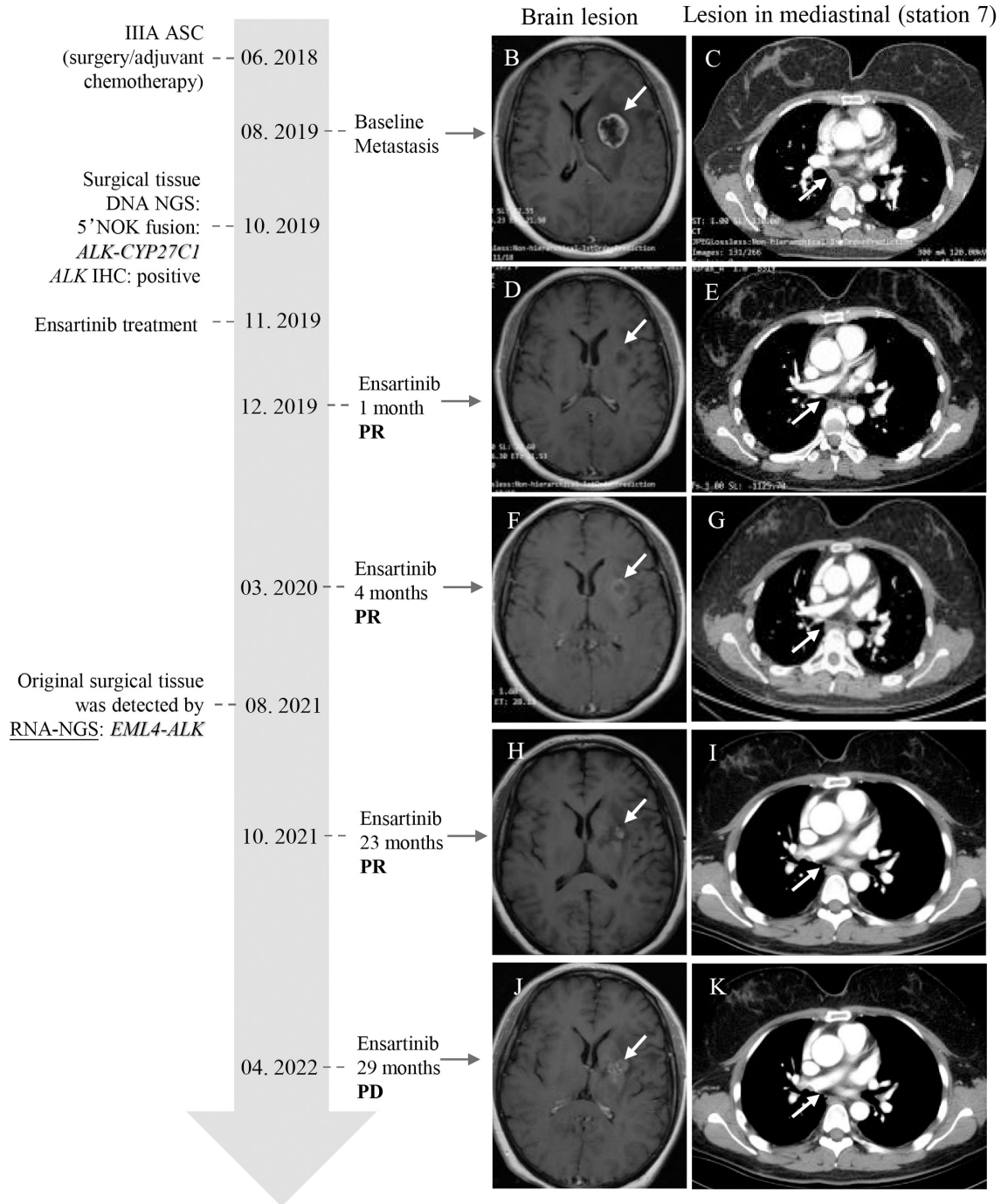
Here, we report a case of lung adenosquamous carcinoma that recurred after surgical resection, harboring a novel 5' NOK fusion form, *ALK-CYP27C1*, which benefited from the *ALK* inhibitor ensartinib. RNA NGS revealed that it was *EML4-ALK* (E13:A20, V1), which suggested that further detection is required at the RNA level when a single 5' NOK is revealed by DNA NGS.

## Case description

The patient was a 49-year-old Mongolian female with no history of smoking. She was admitted to our hospital with a cough of 2 months. After systematic examination and immunohistochemistry, adenocarcinoma of the right lung was considered. She then underwent a right middle and

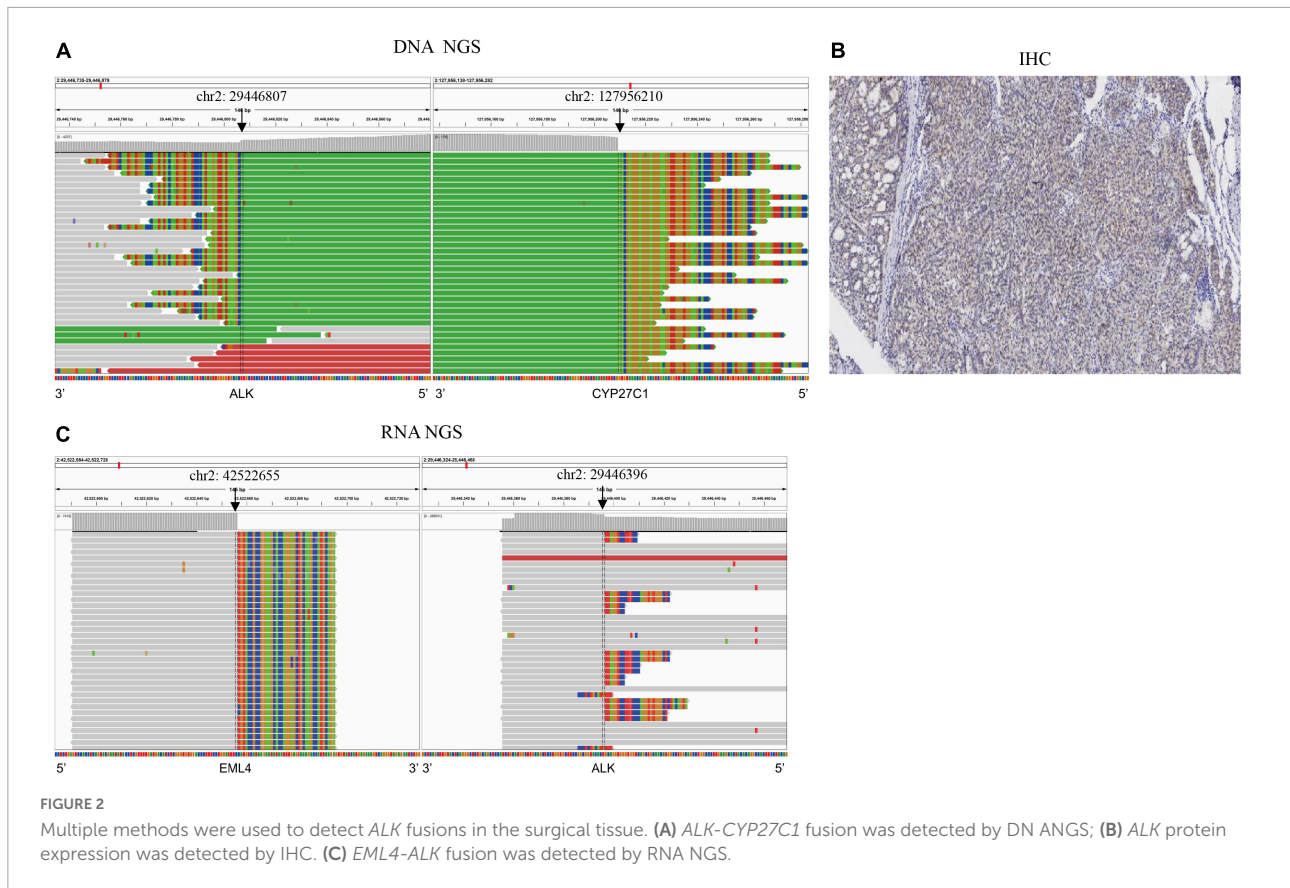
lower lobectomy in June 2018. The postoperative pathology indicated lung adenosquamous carcinoma, and the pathologic stage was IIIA; thus, she accepted pemetrexed plus carboplatin as standard adjuvant chemotherapy (Figure 1A). However, 14 months after the operation, in August 2019, she was referred to the hospital for headache. The brain MRI revealed a nodule in the left basal ganglia with edema, compression of the ventricle, and right midline shift (Figure 1B). The chest CT showed lymphadenopathy in the mediastinum (station 7) (Figure 1C). The patient then received radiotherapy for the brain lesion in September 2019. Mannitol and a small dose of dexamethasone were administered to reduce intracranial pressure, and the headache symptoms were slightly relieved, but the brain lesion did not shrink significantly (data not shown). To determine the next treatment, in October 2019, DNA NGS was performed on her previous surgical tissue specimen using OncoPanscan™ at the Diagnostic Laboratory of Genetron Health (Supplementary Data 1). A 5' NOK fusion *ALK-CYP27C1* (A19:C5) was identified in DNA level, of which a breakpoint occurred in chromosome 2: 29,446,807 (intron 19-*ALK*) and chromosome 2: 127,956,210 (intron 4-*CYP27C1*) (Figure 2A). The *ALK* fusion part was in the 5-terminal region without a kinase domain, which was previously considered to not express the *ALK* protein. The *ALK* fusion was validated by another DNA panel, Oncofocus™ (Genetron Health) (Supplementary Figure 1 and Supplementary Data 1). To confirm whether the *ALK* protein was expressed, IHC was further performed on a Ventana platform, and *ALK* protein expression was positive (Figure 2B). According to the positive results of *ALK* IHC, the patient was enrolled in a donation project about a new *ALK* inhibitor, ensartinib, on 19 November 2019, and received 225 mg orally once daily. One week after the treatment, the dizziness and headache symptoms were relieved, and no adverse events were observed. The brain and mediastinal lesions (station 7) were significantly reduced after 1 month of medication (Figures 1D,E; December 2019). The chest lymph node disappeared after an additional 3 months of ensartinib treatment (Figures 1F,G; March 2020). The brain lesion shrank again after 23 months of ensartinib treatment (Figures 1H,I; October 2021). Because different *ALK* fusions affect the efficacy of *ALK* inhibitors differently, previous articles have reported differences between DNA NGS and RNA NGS in fusion detection, and RNA NGS had become increasingly popular in recent years. We hoped to perform RNA NGS detection to guide subsequent treatment. In addition, we also wanted to explore the cause of efficient treatment of ensartinib against the 5' NOK fusion of *ALK*. Therefore, RNA NGS was conducted on the same surgical tissue specimen using Fusioncapture™ (Genetron Health) in August 2021, which involved all exons of the *ALK* gene for any type of *ALK* fusion detected (Supplementary Data 2). Intriguingly, the result showed *EML4-ALK* (V1, E13:A20) instead of *ALK-CYP27C1* (Figure 2C). When the patient had been using ensartinib for 29 months, the brain lesion began

**A**



**FIGURE 1**

Timeline of the 5' NOK *ALK* fusion clinical diagnosis and treatment and MRI/X-ray computed tomography of lesions of the brain and mediastinal lymph node (station 7) of the patient. (A) Timeline of patient clinical diagnosis and treatment. (B,C) Baseline: brain lesion (24\*21 mm) and lymphadenopathy in the mediastinum (station 7) (short diameter, 10 mm). (D,E) One month after ensartinib treatment: brain lesion reduction (18\*14 mm) and the lesion in the mediastinum (station 7) was reduced (short diameter, 4 mm). (F,G) Four months after ensartinib treatment: brain lesion reduction (16\*15 mm) and the lesion in the mediastinum (station 7) disappeared. (H,I) Twenty-three months after ensartinib treatment: the brain lesion shrank again (15.8\*13.8 mm), and there was no lesion in the mediastinal region (station 7). (J,K) Twenty-nine months after ensartinib treatment: the brain lesion progressed (28\*21 mm), but the mediastinal region (station 7) was still responding. The white arrow indicates the lesion. ASC: adenosquamous carcinoma; PR: partial response; PD: progressive disease.



to progress, while mediastinal lesions (station 7) continued to respond (Figures 1J,K, April 2022). Following 29 months of clinical and radiological response, the patient experienced disease progression in the brain only; following stereotactic radiotherapy, the patient resumed therapy but had clinical progression. The patient did not feel unwell during follow-up.

## Discussion

Here, we present a case of postoperative recurrent lung adenosquamous carcinoma in which a surgical tissue was identified by DNA NGS as a novel form of 5' NOK fusion, *ALK-CYP27C1* (A19:C5) while also being positive for the *ALK* protein by IHC. Treatment with the *ALK* inhibitor ensartinib was administered for nearly 29 months until the brain lesion progressed. Then, the patient received local stereotactic radiotherapy and continued to take ensartinib. The effect was similar to the results of a randomized clinical trial of ensartinib on patients with *ALK*-positive NSCLC that showed a median PFS of 25.8 months (13).

To the best of our knowledge, this is the first report on a 5'NOK form of *ALK* fusion in DNA NGS that benefited from an *ALK* inhibitor (ensartinib). Subsequently, the same surgical tissue specimen was further verified by RNA NGS.

No *ALK-CYP27C1* fusion was detected at the RNA level, but *EML4-ALK* (V1, E13:A20), a common fusion form of *ALK*, was detected. Recent studies have shown that clinical benefits were different among various forms of *EML4-ALK*, and that *EML4-ALK* (E13:A20, V1) can respond well to *ALK* inhibitors, including ensartinib (5, 14, 15). In the present case, the patient was diagnosed as having an *EML4-ALK* (E13:A20, V1) fusion at the RNA level. This may be the reason the patient responded well to ensartinib. Here, we found for the first time a 5' NOK fusion that could have resulted in a functional *EML4-ALK* transcript and responded well to the *ALK* inhibitor ensartinib.

Although various techniques can be used to identify gene fusion, including IHC, FISH, DNA NGS, and RNA NGS, each technique has its own limitations. The ALEX trial showed that because of low sensitivity, the result of DNA level detection by FISH was not enough for fusion detection and required a verification by protein level detection (i.e., IHC) (9), and that IHC was not as specific as FISH (3, 7). Moreover, the detection flux of FISH and IHC is very low. The NGS technology has the advantage of high-throughput, and DNA NGS has been widely conducted in clinical practice. However, previous reports showed that DNA NGS results were different from RNA NGS results in fusion detection, such as many uncommon fusion forms (intergenic-breakpoint fusions, novel partner fusions, and intragenic fusions) identified at the DNA level by DNA NGS,

had been verified as common fusions, or uncommon functional fusions, or no transcription at the RNA level (16, 17). For RNA NGS, high-quality RNA is also difficult to obtain (3). The reasons for different fusion being detected at the DNA and RNA levels in the same sample are unclear. A possible explanation may be chromothripsis, which can result in thousands of chromosomal rearrangements (18). In this case, *ALK-CYP27C1* has been detected by DNA NGS, while *EML4-ALK* was found at the RNA level, which may be due to *EML4-ALK* rearrangement being mediated by chromothripsis. Based on the evidence of previous reports and this case, we suggest that validation of the RNA level is needed for fusions without an intact kinase domain. These results indicate that a combination of multiple methods, especially DNA NGS and RNA NGS, is necessary for fusion detection.

## Conclusion

To our knowledge, this is the first case of a 5' NOK fusion that benefited from an *ALK* inhibitor. This demonstrates the necessity of further analysis from the RNA level of 5' NOK fusion per DNA NGS to identify whether there is another functional structural form. Precise analysis of such a fusion (5' NOK) form may yield clinical benefits for patients. In clinical practice, it would be better to analyze DNA and RNA simultaneously to reduce the turnaround time.

## Data availability statement

The original contributions presented in this study are included in the article/**Supplementary material**, further inquiries can be directed to the corresponding author/s.

## Author contributions

CZ, TM, and XZ contributed to conception and design of the study. HY and YF wrote the first draft of the manuscript. HAL and ZL prepared figures and background research. JZ prepared

figures. HUL and CL prepared background research. All authors contributed to manuscript revision, read, and approved the submitted version.

## Funding

This study was supported by the Scientific Research Foundation for Middle-Aged and Young Scientist of San Sheng TCP (NO.008) and Foundation for Study Encouragement of Beijing Medical (NO.YXJL-2020-0785-0315).

## Conflict of interest

Authors YF, JZ, XZ, and TM were employed by Genetron Health (Beijing) Technology, Co., Ltd.

The remaining authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

## Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.

## Supplementary material

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fmed.2022.979032/full#supplementary-material>

### SUPPLEMENTARY FIGURE 1

*ALK-CYP27C1* fusion was validated by DNA NGS in another DNA panel.

## References

- Cao W, Chen HD, Yu YW, Li N, Chen WQ. Changing profiles of cancer burden worldwide and in China: A secondary analysis of the global cancer statistics 2020. *Chin Med J.* (2021) 134:783–91. doi: 10.1097/CM9.0000000000001474
- Yang CY, Yang JCH, Yang PC. Precision management of advanced non-small cell lung cancer. *Annu Rev Med.* (2020) 71:117–36. doi: 10.1146/annurev-med-051718-013524
- Murakami Y, Mitsudomi T, Yatabe Y. A screening method for the *ALK* fusion gene in NSCLC. *Front Oncol.* (2012) 2:24. doi: 10.3389/fonc.2012.00024
- Liu S, Huang T, Liu M, He W, Zhao Y, Yang L, et al. The genomic characteristics of *ALK* fusion positive tumors in Chinese NSCLC patients. *Front Oncol.* (2020) 10:726. doi: 10.3389/fonc.2020.00726
- Yoshida T, Oya Y, Tanaka K, Shimizu J, Horio Y, Kuroda H, et al. Differential crizotinib response duration among *ALK* fusion variants in *ALK*-positive non-small-cell lung cancer. *J Clin Oncol.* (2016) 34:3383–9. doi: 10.1200/JCO.2015.65.8732
- Zhao R, Zhang J, Han Y, Shao J, Zhu L, Xiang C, et al. Clinicopathological features of *ALK* expression in 9889 cases of non-small-cell lung cancer and

genomic rearrangements identified by capture-based next-generation sequencing: A Chinese retrospective analysis. *Mol Diagn Ther.* (2019) 23:395–405. doi: 10.1007/s40291-019-00389-y

7. Wu YC, Chang IC, Wang CL, Chen TD, Chen YT, Liu HP, et al. Comparison of IHC, FISH and RT-PCR methods for detection of ALK rearrangements in 312 non-small cell lung cancer patients in Taiwan. *PLoS One.* (2013) 8:e70839. doi: 10.1371/journal.pone.0070839

8. Cabillic F, Gros A, Dugay F, Begueret H, Mestroux L, Chiforeanu DC, et al. Parallel FISH and immunohistochemical studies of ALK status in 3244 non-small-cell lung cancers reveal major discordances. *J Thorac Oncol.* (2014) 9:295–306. doi: 10.1097/JTO.000000000000072

9. Mok T, Peters S, Camidge DR, Noé J, Gadgeel S, Ignatius Ou SH, et al. Outcomes according to ALK status determined by central IHC or FISH in patients with ALK-positive NSCLC enrolled in the phase III ALEX study. *J Thorac Oncol.* (2020) S1556-0864:30815–7.

10. Li W, Guo L, Liu Y, Dong L, Yang L, Chen L, et al. Potential unreliability of uncommon ALK, ROS1, and RET genomic breakpoints in predicting the efficacy of targeted therapy in NSCLC. *J Thorac Oncol.* (2021) 16:404–18. doi: 10.1016/j.jtho.2020.10.156

11. Li W, Liu Y, Li W, Chen L, Ying J. Intergenic breakpoints identified by DNA sequencing confound targetable kinase fusion detection in NSCLC. *J Thorac Oncol.* (2020) 15:1223–31. doi: 10.1016/j.jtho.2020.02.023

12. Ou SHI, Zhu VW, Nagasaka M. Catalog of 5' fusion partners in ALK-positive NSCLC Circa 2020. *JTO Clin Res Rep.* (2020) 1:100015. doi: 10.1016/j.jtocrr.2020.100015

13. Horn L, Wang Z, Wu G, Poddubskaya E, Mok T, Reck M, et al. Ensartinib vs crizotinib for patients with anaplastic lymphoma kinase-positive non-small cell lung cancer: A randomized clinical trial. *JAMA Oncol.* (2021) 7:1617–25. doi: 10.1001/jamaoncol.2021.3523

14. Yang Y, Zhou J, Zhou J, Feng J, Zhuang W, Chen J, et al. Efficacy, safety, and biomarker analysis of ensartinib in crizotinib-resistant, ALK-positive non-small-cell lung cancer: A multicentre, phase 2 trial. *Lancet Respir Med.* (2020) 8:45–53. doi: 10.1016/S2213-2600(19)30252-8

15. Zhang SS, Nagasaka M, Zhu VW, Ou SHI. Going beneath the tip of the iceberg. Identifying and understanding EML4-ALK variants and TP53 mutations to optimize treatment of ALK fusion positive (ALK+) NSCLC. *Lung Cancer.* (2021) 158:126–36. doi: 10.1016/j.lungcan.2021.06.012

16. Shi M, Wang W, Zhang J, Li B, Lv D, Wang D, et al. Identification of RET fusions in a Chinese multicancer retrospective analysis by next-generation sequencing. *Cancer Sci.* (2022) 113:308–18. doi: 10.1111/cas.15181

17. Xia P, Zhang L, Li P, Liu E, Li W, Zhang J, et al. Molecular characteristics and clinical outcomes of complex ALK rearrangements identified by next-generation sequencing in non-small cell lung cancers. *J Transl Med.* (2021) 19:308. doi: 10.1186/s12967-021-02982-4

18. Kodama T, Motoi N, Ninomiya H, Sakamoto H, Kitada K, Tsukaguchi T, et al. A novel mechanism of EML4-ALK rearrangement mediated by chromothripsis in a patient-derived cell line. *J Thorac Oncol.* (2014) 9:1638–46. doi: 10.1097/JTO.0000000000000311