

# Alectinib efficacy in advanced lung adenocarcinoma with coexistence of a novel ALK-MTUS2 and STRN3-ALK double fusion: A case report and literature review

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**Abstract.** In total, >90 unique anaplastic lymphoma kinase (ALK) rearrangement fusion partners have been identified, each with a unique sensitivity to ALK tyrosine kinase inhibitors, rendering targeted therapy challenging. In the present study, the efficacy of alectinib in a patient with advanced lung adenocarcinoma harbouring a novel ALK-microtubule-associated tumour suppressor candidate 2 (MTUS2) fusion and a rare Striatin 3 (STRN3)-ALK fusion was assessed. A 20-year-old non-smoker was hospitalised with a persistent cough. Subsequent positron emission tomography/computed tomography revealed a tumour in the right lower lobe, with mediastinal, hilar, lymph node and bone metastases. Cranial magnetic resonance imaging also revealed a cerebral metastasis. Bronchoscopic biopsy revealed an adenocarcinoma in the right lower lobe nodule, resulting in a clinical stage IVB diagnosis (cT2bN3bM1c). Targeted next-generation sequencing of tissue and blood samples revealed ALK-MTUS2 and STRN3-ALK fusions. The patient was treated with alectinib as the first-line therapy and a durable partial response was achieved after 3 months, with disappearance of the brain metastases. To the best of our knowledge, the present study represents the first discovery of simultaneous ALK-MTUS2 and STRN3-ALK fusions. The clinical outcome offers

potential therapeutic options for patients with rare ALK rearrangements and underscores the necessity for further research on the functions of these fusions.

## Introduction

Anaplastic lymphoma kinase (ALK) rearrangement occurs in an estimated 4-5% of patients diagnosed with non-small cell lung cancer (NSCLC), a characteristic associated with a younger age, non-smoking status and advanced disease stage at diagnosis (1-3). These patients also face a heightened risk of brain metastasis, affecting 50-60% of cases (4). Among them, 16.2% present with multiple ALK fusions, a prevalence that is rising due to the adoption of next-generation sequencing (NGS) in detecting these fusions (5). While the echinoderm microtubule-associated protein-like 4 (EML4)-ALK fusion is the most common rearrangement, non-canonical fusions and particularly dual ALK fusion variants are less frequently observed (6). A multicentre national registry by the French National Cancer Institute demonstrated that patients harbouring actionable genetic alterations, predominantly epidermal growth factor receptor (EGFR) mutations and ALK rearrangements, exhibited a 4.7-month extension in median overall survival time when treated with targeted therapeutics, underscoring the clinical utility of molecular profiling (1). Concurrently, real-world evidence highlights notable heterogeneity in first-line treatment duration of response (8.3-13.9 months) among ALK-rearranged cases, a variability strongly associated with the specific ALK fusion partner gene (2). Therefore, investigating the treatment response and its durability in patients with dual ALK fusion variants holds distinct clinical implications.

Alectinib, an effective ALK tyrosine kinase inhibitor (TKI), demonstrates promising results in treating advanced ALK-positive NSCLC and is particularly effective against brain metastases (4). Despite its recognized utility, research on the efficacy of alectinib in patients with NSCLC and concurrent brain metastasis harbouring rare dual fusion mutations remains scarce. To the best of our knowledge, the present study is the first to report a novel case of a patient with rare dual

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Striatin 3 (STRN3)-ALK and ALK-microtubule-associated tumour suppressor candidate 2 (MTUS2) fusions, who demonstrated a positive response to standard-dose alectinib therapy.

### Case report

At initial presentation (September 2023), a 20-year-old man who had never smoked and had no significant medical history was initially admitted to the Affiliated Hospital of Nanjing University of Chinese Medicine (Nanjing, China) due to a persistent cough for 8 months. Chest computed tomography (CT) showed a soft tissue mass in the lower lobe of the right lung near the hilum and in the dorsal segment of the lower lobe, with enlarged lymph nodes in the mediastinum (Fig. 1A and F). Pulmonary function indicated moderate obstructive hypoventilation, and the bronchodilator test was negative. Post-bronchodilator spirometry showed: Forced expiratory volume in the first second (FEV1), 2.95 l; FEV1%, 63.9%; forced vital capacity (FVC), 3.85 l; FVC%, 70.1%; and FEV1/FVC, 76.53%. Cranial magnetic resonance imaging (MRI) revealed left cerebellar hemisphere and right frontal lobe enhancement nodules, which were considered to be metastatic tumours (Fig. S1A and E). Positron emission tomography/CT indicated hypermetabolic nodules in the posterior basal segment of the lower lobe of the right lung, suggesting lung cancer with multiple lymph node and bone metastases (Fig. S2). Adenocarcinoma was confirmed by bronchoscopic biopsy of the nodules in the basal segment of the lower lobe of the right lung.

Subsequently, two rare STRN3-ALK and ALK-MTUS2 fusions as well as a heterozygous germline frameshift mutation in the ataxia-telangiectasia mutated (ATM) gene (p.M3011Dfs\*6, in exon 63; Fig. S3) were identified through targeted NGS (Geneseeq Prime™; Nanjing Geneseeq Technologies, Inc.) technology (Table 1) that covered exons, fusion-related introns and the microsatellite loci of 437 cancer-related genes with a target region of 1.53 Mb (Data S1). The STRN3-ALK fusion site involved exon 3 of STRN3 and exon 20 of ALK (Fig. 2). The sequence analysis of the ALK-MTUS2 cDNA showed that the 5' untranslated region (5'UTR) of MTUS2 was fused to exon 19 of ALK (Fig. S4). To the best of our knowledge, this rare variant has not been previously reported. Haematoxylin-eosin staining showed histological differentiation typical of lung adenocarcinoma with alveolar and solid structures (Fig. 3A). Immunohistochemical analysis showed diffuse strong positivity for ALK (D5F3) (Fig. 3B) as well as diffuse positive expression of cytokeratin 7, thyroid transcription factor 1 and Napsin A in tumour cells; p53 showed ~80% positive expression, Ki67 exhibited ~15% positive expression, programmed death-ligand 1 (22C3) had a tumour proportion score of ~20% and p40 protein was negatively expressed (Fig. S5).

The final diagnosis was clinical stage IVB (cT2bN3bM1c) lung adenocarcinoma based on the eighth edition of the Tumour-Node-Metastasis classification (7). CT imaging revealed unresectable tumour invasion of major vessels and the trachea (Fig. 1A and F), and the patient presented with multiple distant metastases (bone metastasis and brain metastasis), assessed as clinical stage IV lung cancer, which was not amenable to surgical resection. Therefore, a decision was made

to proceed with systemic therapy. The patient was treated with alectinib (600 mg orally twice a day) and zoledronic acid injection (5 mg intravenously once every 4 weeks) to treat the bone metastasis, after discussion with the Department of Respiratory Medicine and multidisciplinary team consultation. Following initiation of alectinib therapy (September 2023), the patient exhibited a rapid clinical response and the cough improved considerably after just 1 week of receiving alectinib. At present, the patient has maintained treatment for 15 months. Follow-up cranial MRI at 3 months post-treatment initiation (December 2023) demonstrated complete resolution of the brain metastases (Fig. S1B and F), with no recurrence observed during subsequent follow-up (Fig. S1C, D, G and H). CT imaging revealed significant lung tumour shrinkage 3 months after initial treatment (December 2023), and serial chest CT scans throughout the 15-month treatment (December 2024) and follow-up period showed sustained partial response without disease progression, and no adverse events were observed (Fig. 1).

### Discussion

ALK, a receptor tyrosine kinase from the insulin receptor family, is prevalent in various cancer types, such as NSCLC, breast cancer and colorectal cancer (8). Typically, ALK resides on cell membranes and engages with growth factors. In NSCLC, the EML4-ALK fusion is the most frequent rearrangement (8). Recent advancements in genetic testing have identified additional fusion partners of the ALK gene (3,9). The present case study reports initial evidence of the potential efficacy of alectinib in treating two uncommon ALK fusions, ALK-MTUS2 and STRN3-ALK, in a patient diagnosed with lung adenocarcinoma.

The MTUS2 gene is located on chromosome 13q12.3 and has 21 exons. MTUS2 modulates the dynamics at the distal growing tip, or plus-end, of microtubules (10). In an NGS-based study of breast cancer genomics, MTUS2 emerged as a potential driver of the disease (11). Additionally, MTUS2 may be a potential therapeutic target for Alzheimer's disease (12). To date, and to the best of our knowledge, the ALK-MTUS2 fusion in NSCLC has been unreported. Notably, the patient described in the present study harboured an ALK-MTUS2 fusion, where the 5'UTR of MTUS2 was fused to ALK exon 19. Although the patient showed a clinical response to alectinib therapy, this fusion results in the loss of the tyrosine kinase domain of the ALK gene (ALK exon 20-29), which may impair its kinase activity and potentially reduces its clinical significance as a driver mutation (6).

ATM gene mutation is a rare and harmful germline mutation (13). A study reported that 4.7% of patients with lung cancer carry pathogenic or suspected pathogenic germline variants, with the most common being in ATM, checkpoint kinase 2 and breast cancer susceptibility gene 2 (13). In a study of patients with NSCLC, 562 pathogenic mutations were identified in the ATM gene, 62.8% of which were missense mutations (14). In the patient reported in the present study, a heterozygous germline frameshift mutation was detected in exon 63 of the ATM gene (p.M3011Dfs\*6), causing a deletion of bases 9031 to 9034 and a frameshift starting at amino acid 3011. This mutation led to a premature termination codon and a truncated

Table I. Targeted next-generation sequencing detected genetic alterations in both the blood and tumour samples from the patient.

Gene	c.HGVS	p.HGVS	Functional region	Allele frequency (%)	
				Plasma	Tissue
ATM	c.9031_9034del	p.M3011Dfs*6	EX63	48.63	44.16
STRN3-ALK		Fusion	EX3:EX20	0.01	38.52
ALK-MTUS2		Fusion	EX19:5'UTR	6.58	23.85

HGVS, human genome variation society; c.HGVS, description of coding DNA variants by HGVS; p.HGVS, description of protein variants by the HGVS; ALK, anaplastic lymphoma kinase; MTUS2, microtubule-associated tumour suppressor candidate 2; STRN3, Striatin 3; 5'UTR, 5'untranslated region.

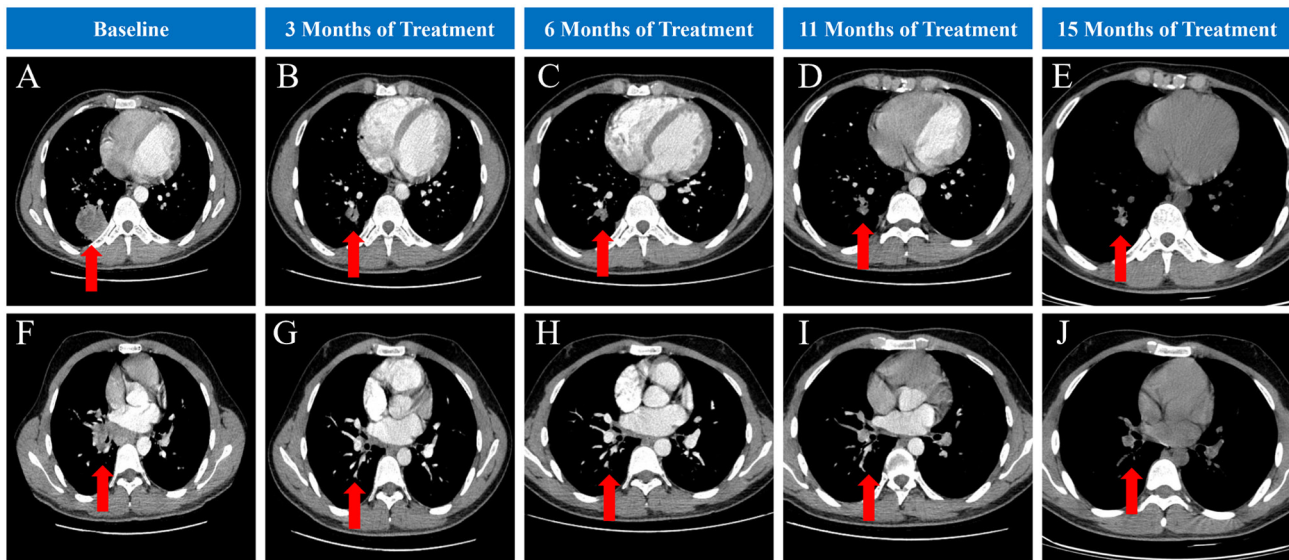


Figure 1. Dynamic monitoring via CT of the patient's response to alectinib. (A) CT shows a large mass in the right lower lobe of the lung at baseline. CT shows partial response after (B) 3, (C) 6, (D) 11 and (E) 15 months of alectinib treatment. (F) CT shows an enlarged right hilar lymph node. CT shows a significant reduction in tumour size after (G) 3, (H) 6, (I) 11 and (J) 15 months of alectinib treatment. CT, computed tomography.

protein that may disrupt critical functional domains. Previous literature has reported the pathogenicity of the STRN-ALK fusion in thyroid cancer, indicating that targeting STRN-ALK can inhibit tumour cell proliferation (15). Meanwhile, ATM, a gene commonly mutated in lung cancer (16), may synergistically promote tumour cell proliferation by regulating the cell cycle in conjunction with the STRN-ALK fusion. This may explain the mechanism behind the advanced lung cancer observed in the young patient reported in the present study.

The STRN3 gene product, a protein with a coiled-coil domain, mediates the constitutive activation of ALK through a dimerization-driven mechanism (17). The literature on STRN3-ALK fusions was reviewed to explore previous therapeutic interventions, with 12 published articles on patients with NSCLC harbouring this fusion identified. The findings are summarised in Table II, which outlines the characteristics and therapeutic approaches of the disease (6,17-27). Of the 12 patients, 10 were men and 8 were non-smokers, with ages ranging from 29 to 70 years old. Additionally, 11 patients identified were diagnosed with adenocarcinoma, and the histological type was not described for 1 patient (21). Only 2 patients had

metastatic lung cancer (18,24), while the remaining patients were diagnosed with primary tumours. The majority of patients underwent NGS, mostly using tissue for testing, and the structural mutations were all characterized as STRN3-ALK (fusion breakpoint S3, A20). Various other mutations coexisted with STRN3-ALK, including ALK-DnaJ homolog subfamily C member 27 fusion, phosphoinositide-dependent kinase 1-ALK fusion, tumour protein p53 (TP53), EGFR and breast cancer susceptibility gene 1. Upon identifying the STRN3-ALK mutation, alectinib was administered in 2 cases (18,20), while crizotinib (17) and gefitinib (26,27) were each administered in 1 and 2 cases, respectively. Alectinib was administered as a first-line treatment to 4 out of 12 patients, with efficacy sustained for 6-19 months (19,22,23,25). In total, 2 patients achieved short-term partial response with alectinib but were switched to crizotinib due to disease progression, showing sustained partial response (18,23). One study suggests hepatocyte growth factor receptor amplification may confer alectinib resistance (18), while another implicates TP53 and phosphatidylinositol 4,5-bisphosphate 3-kinase catalytic subunit  $\alpha$  isoform mutations (23). In summary, alectinib demonstrates therapeutic efficacy as

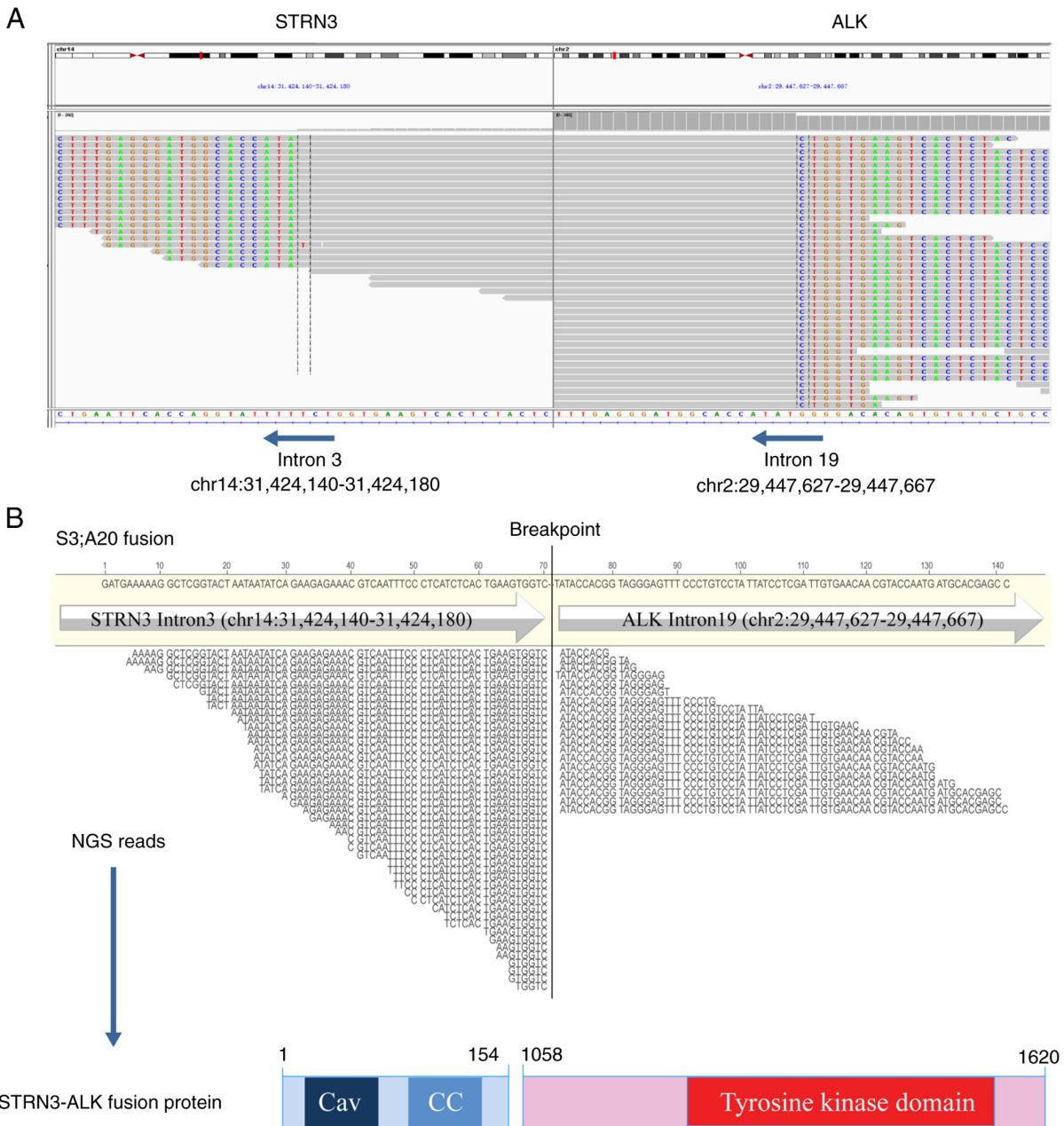


Figure 2. A fusion variant of STRN3 intron 3 with ALK intron 19 was identified by NGS analysis and was considered to cause a rare STRN3-ALK fusion transcript in which exon 3 of STRN3 was fused to exon 20 of ALK (S3:A20). (A) Paired-end sequencing data indicated the somatic intrachromosomal STRN-ALK fusion, as demonstrated by the Integrative Genomics Viewer. (B) Diagram depicting the STRN3-ALK fusion (S3:A20). CC, coiled coil domain; Cav, caveolin-binding domain; Chr, chromosome; NGS, next-generation sequencing; ALK, anaplastic lymphoma kinase; STRN3, Striatin 3.

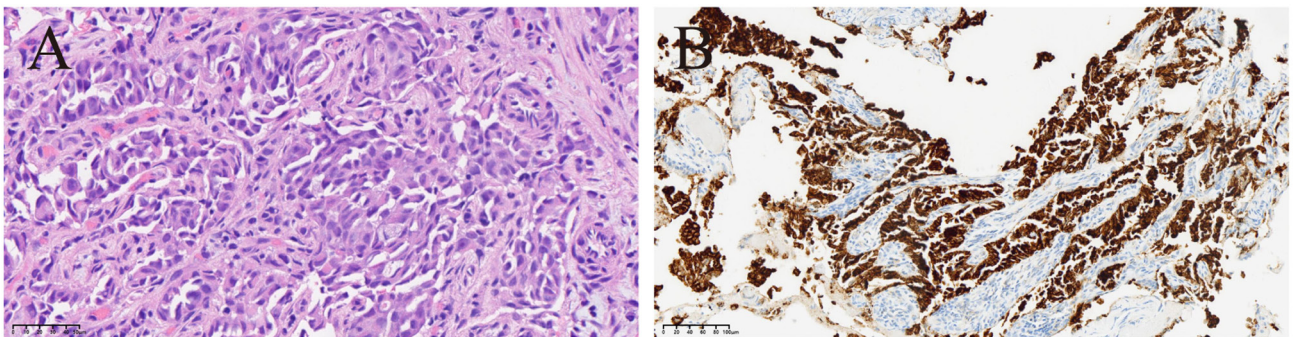


Figure 3. Tumour tissue histological typing. (A) The histological typing identified adenocarcinoma of the lung with acinar and solid patterns (haematoxylin-eosin; magnification, x400). (B) Immunohistochemical analysis showed diffuse strong positivity for anaplastic lymphoma kinase (D5F3) (magnification, x200).

Table II. Case reports of patients with NSCLC<sup>a</sup> harbouring the STRN3-ALK (S3:A20) fusion.

First author, year	Sex	Age (years)	Smoking status	Stage	Co-mutations	PD-L1 expression level	TKIs administered before identifying STAN3-ALK	Treatments, lines	Response	Duration of response (months)	(Refs.)
Yang <i>et al</i> , 2017	M	59	Never	Postoperative metastases	MYC amplification and TP53 R181C	Not reported	No	Crizotinib, first	CR	6, ongoing	(24)
Nakanishi <i>et al</i> , 2017	M	51	Never	IV	-	Not reported	Alectinib	Docetaxel, first	PD	-	(20)
Ren <i>et al</i> , 2019	F	52	Never	IV	CDKN2A p9del	Not reported	Crizotinib	Ceritinib, first	PR	26, ongoing	(17)
Zhou <i>et al</i> , 2019	M	43	Smoker	IV	EGFR ex19del and EGFR T790M	Not reported	Gefitinib	Gefitinib + crizotinib, first	PR	6	(27)
Su <i>et al</i> , 2020	M	64	Never	IV	GRM8 E508K and SETD2	Negative	No	Alectinib, first	PR	19, ongoing	(22)
Nagasaka <i>et al</i> , 2020	M	66	Never	IV	E1553K TP53 L43fs and MYC amplification	98% (22C3)	No	Alectinib, first	CR	6, ongoing	(19)
Li <i>et al</i> , 2021	M	42	Never	Postoperative metastases	MET amplification, TP53 C991T, TP53 C742T and BRCA1 A5347C	Not reported	Alectinib	Crizotinib, first	PR	11, ongoing	(18)
Zeng <i>et al</i> , 2021	F	29	Never	IV	PDK1-ALK (P7: A20) and TP53 ex9 splicing mutation	Not reported	No	Alectinib, first	PR	7, ongoing	(25)

Table II. Continued.

First author, year	Sex	Age (years)	Smoking status	Stage	Co-mutations	PD-L1 expression level	TKIs administered before identifying STAN3-ALK	Treatments, lines	Response	Duration of response (months)	(Refs.)
Sun <i>et al</i> , 2021	M	65	Smoker	IV	PIK3CA G106V, KRAS G12C and TP53 R267H	Not reported	No	Alectinib, first; crizotinib, second; pemetrexed + carboplatin, third	PD for alectinib; PR for crizotinib	5, for crizotinib	(23)
Zeng <i>et al</i> , 2021	M	38	Never	IV	EGFR L858R, EGFR T790M, RB1 R445, TP53 T284Afs62 and EML4-ALK fusion	Not reported	Gefitinib	Osimertinib + crizotinib, first	SD	5, ongoing	(26)
Zhang <i>et al</i> , 2022	M	67	Smoker	IV	ALK-DNAJC27 (A19: D4), ALK E974Q, EPHA5 L63S, PRKDC K3196N, PFWD2 S90C and RPL5 V88Ffs*2	Negative	No	Crizotinib, first; ensartinib, second	PR for crizotinib; PR for ensartinib	18, for crizotinib; 13, ongoing, for ensartinib	(6)
Song <i>et al</i> , 2023 <sup>a</sup>	M	70	Smoker	IV	TP53 R158L	Not reported	No	Ensartinib, first	PR for ensartinib	6, ongoing, for ensartinib	(21)

<sup>a</sup>All cases were diagnosed as adenocarcinoma except for the study by Song *et al* (21), in which the pathology was not reported. M, male; F, female; CR, complete response; PR, partial response; SD, stable disease; PD, progressive disease; STRN3, Striatin 3; ALK, anaplastic lymphoma kinase; PD-L1, programmed death-ligand 1; MYC, MYC proto-oncogene protein; TP53, tumour protein p53; CDKN2A, cyclin-dependent kinase inhibitor 2A; EGFR, epidermal growth factor receptor; GRM8, glutamate metabotropic receptor 8; SETD2, SET domain containing 2, hSET2, p231HBP; MET, hepatocyte growth factor receptor; BRCA1, breast cancer type 1 susceptibility protein; PDK1, pyruvate dehydrogenase kinase 1; PIK3CA, phosphatidylinositol-4,5-bisphosphate 3-kinase catalytic subunit  $\alpha$ ; KRAS, GTPase Kras; RB1, retinoblastoma-associated protein; DNAJC27, DnaJ homolog subfamily C member 27; EPHA5, ephrin type-A receptor 5; PRKDC, DNA-dependent protein kinase catalytic subunit; RPL5, large ribosomal subunit protein uL18.

first-line treatment for advanced STRN3-ALK fusion-positive adenocarcinoma. However, reported follow-up remains limited, and resistance mechanisms require further investigation.

Alectinib, a second-generation ALK-TKI, shows superior efficacy and reduced toxicity compared with crizotinib in the treatment of ALK-positive NSCLC, as evidenced by the ALEX (28) and ALESIA (29) studies. Additionally, a real-world study in a Chinese population confirmed its significant systemic and central nervous system benefits (30). As aforementioned, alectinib has shown promising clinical efficacy in treating patients with NSCLC that harbour the STRN-ALK fusion gene (19,25). Compared with previously reported cases of lung adenocarcinoma with a STRN3-ALK fusion, the patient described in the present is unique as being the youngest (20 years old), with no smoking history, no identifiable exposure to conventional risk factors and harbouring both a novel ALK-MTUS2 fusion and a rare STRN3-ALK fusion. The patient also presented with brain, bone and multiple lymph node metastases. After the 15 months of treatment (December 2024), the patient has responded well to alectinib treatment, with complete resolution of brain metastases and no recurrence observed. Considering the age and good baseline health of the patient, a full therapeutic dose of 600 mg alectinib twice daily was prescribed. Follow-up assessments at 3, 6 and 15 months revealed satisfactory therapeutic outcomes, with no adverse events observed.

The present study has certain limitations. First, it is a single case report, and the efficacy of alectinib in ALK dual fusions needs to be further validated through large-scale clinical trials. Second, the present study only involves the identification of the STRN3-ALK and ALK-MTUS2 dual fusion genes, without exploring the potential functional changes brought by these gene fusions, which requires further research and validation in the future. Third, the current follow-up time for the reported patient is relatively limited, and the specific efficacy of alectinib still requires longer-term follow-up observation.

In conclusion, the present study reports a lung adenocarcinoma case with an ALK-MTUS2 fusion and concurrent STRN3-ALK fusion, highlighting its innovation. Follow-up results confirmed the efficacy of the ALK TKI, alectinib, in NSCLC with this rare fusion type. The findings expand the spectrum of ALK rearrangements and explore the therapeutic strategies for lung adenocarcinoma.

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### Availability of data and materials

The high-throughput sequencing data generated in the present study may be found in the NCBI SRA database under accession number SRP584787 or at the following URL: <https://www.ncbi.nlm.nih.gov/sra/?term=SRP584787>. All other data generated in the present study may be requested from the corresponding author.

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### Authors' contributions

XQ contributed to conceptualization, writing the original draft and reviewing and editing the manuscript. YW and XW contributed to conceptualization, reviewing and editing the manuscript and obtaining MRI and PET/CT images. SX reviewed and edited the manuscript, obtained CT scan images and advised on patient treatment. YZ reviewed and edited the manuscript and carried out the high-throughput sequencing experiments. HH and XZ contributed to conceptualization, supervision, reviewing and editing the manuscript and advised on patient treatment. XQ, YW, XW, SX, YZ, HH and XZ confirm the authenticity of all the raw data. All authors read and approved the final version of the manuscript.

### Ethics approval and consent to participate

Written informed consent was obtained from the patient, including consent to participate.

### Patient consent for publication

Written informed consent was obtained from the patient, including consent for publication of the findings.

### Competing interests

The authors declare that they have no competing interests.

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