

OPEN ACCESS

EDITED BY
José Díaz-Chávez,

Instituto Nacional de Cancerología (INCAN),

REVIEWED BY

Michael Pogorzelski, Essen University Hospital, Germany Lishui Niu, Central South University, China

*CORRESPONDENCE

Zhe Yang,

sdslyyyz@sina.com

Xiaomei Li,

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

RECEIVED 28 September 2024 ACCEPTED 11 September 2025 PUBLISHED 08 October 2025

CITATION

Lei L, Zhan X, Li J, Ding Y, Li J, Zhou F, Feng A, Li X and Yang Z (2025) Addition of bevacizumab to gefitinib plus chemotherapy as first-line therapy in EGFR L858R mutate advanced nonsmall cell lung cancer patients. Front. Pharmacol. 16:1503171. doi: 10.3389/fphar.2025.1503171

COPYRIGHT

© 2025 Lei, Zhan, Li, Ding, Li, Zhou, Feng, Li and Yang. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). 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.

Addition of bevacizumab to gefitinib plus chemotherapy as first-line therapy in EGFR L858R mutate advanced non-small cell lung cancer patients

Lingli Lei^{1†}, Xiang Zhan^{1,2†}, Jixian Li¹, Yi Ding¹, Jiaxin Li¹, Fengge Zhou³, Alei Feng³, Xiaomei Li^{3*} and Zhe Yang^{1,3*}

¹Shandong Provincial Hospital, Shandong University, Jinan, Shandong, China, ²Department of Radiotherapy, the Second Hospital of Hebei Medical University, Jinan, Shandong, China, ³Tumor Research and Therapy Center, Shandong Provincial Hospital Affiliated to Shandong First Medical University, Jinan, Shandong, China

Background: For advanced non-small cell lung cancer patients with the epidermal growth factor receptor L858R mutation, the efficacy of the combination of tyrosine kinase inhibitor (TKI) and chemotherapy is suboptimal. Currently, it is unclear whether the combination of bevacizumab, gefitinib, and chemotherapy could improve the survival.

Materials and methods: Data was retrospectively collected at Shandong Provincial Hospital between June 2019 and December 2022. The patients were divided into two groups, ATC group receiving bevacizumab, gefitinib, and chemotherapy, and TC group receiving gefitinib and chemotherapy. After propensity score matching (PSM), progression-free survival (PFS) and overall survival (OS) were calculated along with the objective response rate (ORR) and disease control rate (DCR).

Results: The study enrolled 217 patients, 58 in the ATC group and 149 in the TC group, and was adjusted to 55 and 118 after PSM, respectively. Both the ORR and DCR were higher in the ATC group compared to the TC group (ORR: 76.4% versus 65.3%; DCR: 89.1% versus 80.1%). After 41.30 months of follow-up, the first-line PFS in the ATC group was significantly longer than in the TC group, while OS was not (PFS: 22.26 months versus 19.18 months, P = 0.02; OS: 42.18 months versus 39.42 months, P = 0.91). Univariate and multivariate analyses indicated that ATC treatment and the absence of brain metastases positively predict PFS, with no variables that dependently predict OS.

Conclusion: The combination of bevacizumab, gefitinib, and chemotherapy significantly benefits patients with the L858R mutation in first-line PFS but not OS.

KEYWORDS

non-small cell lung cancer, epidermal growth factor receptor, bevacizumab, gefitinib, prognosis

1 Introduction

Lung cancer is one of the most prevalent malignant tumors and the leading cause of tumor-related deaths either worldwide or specifically within the Chinese population (Sung et al., 2021; Han et al., 2024). EGFR-mutated non-small cell lung cancer plays a significant role in the landscape of lung cancer. The L858R mutation in exon 21 and the exon 19 deletion are the most prevalent types of EGFR mutations, constituting around 80%–90% of cases (Hsu et al., 2018). The extensive use of tyrosine kinase inhibitors (TKI) that target mutant EGFR has revolutionized the standards of treatment and management for patients with this disease. The inevitable challenge of resistance to TKI drugs has necessitated the adoption of combination treatment strategies such chemotherapy to augment the effectiveness of TKIs in patients. A series of clinical trials have confirmed the reliability of chemotherapy in enhancing the efficacy of TKIs (Wu et al., 2013; Hosomi et al., 2020; Planchard et al., 2023). Moreover, the combination of bevacizumab, a VEGFR antibody, with TKIs has also demonstrated efficacy as a treatment approach. Two clinical trials carried out in China and Japan have demonstrated that the initial introduction of Bevacizumab in conjunction with erlotinib extends patients' progression-free survival (PFS) compared to using erlotinib alone treatment while maintaining acceptable safety profiles (Zhou et al., 2021; Kawashima et al., 2022).

While L858R and 19del variants exhibit sensitivity to TKIs, variations in treatment response exist. Researches indicate that individuals carrying the L858R mutation demonstrate a diminished response to identical treatments (Lee et al., 2013; Sheng et al., 2016). The underlying reason may be owing to the impact of structural variances in the EGFR protein resulting from the mutation site on the affinity for binding to TKI drugs (Kumar et al., 2008). Another pertinent factor to consider is the drug-resistant mutation T790M, which has been observed to exhibit a greater propensity to occur concurrently with L858R (Pao et al., 2005).

The purpose of this study is to investigate whether the addition of Bevacizumab to TKI drugs in combination with chemotherapy, a stronger treatment regimen, can bring more therapeutic benefits for patients with L858R mutation. In the era of precision medicine, providing more evidence-based medicine for EGFR-mutated lung cancer.

2 Materials and methods

2.1 Study design and patients

Patients who were diagnosed with metastatic non-small cell lung cancer and initiated first-line therapy at the Oncology Department

Abbreviations: NSCLC, Non-small cell lung cancer; EGFR, Epidermal Growth Factor Receptor; TKI, Tyrosine kinase inhibitor; PSM, propensity score matching; PFS, Progression-free survival; OS: Overall survival; ORR, Objective response rate; DCR: Disease control rate; CR, Complete response; PR, Partial response; SD, Stable disease. HR, Hazard ratio IQR, Interguartile range; CI, Confidence interval.

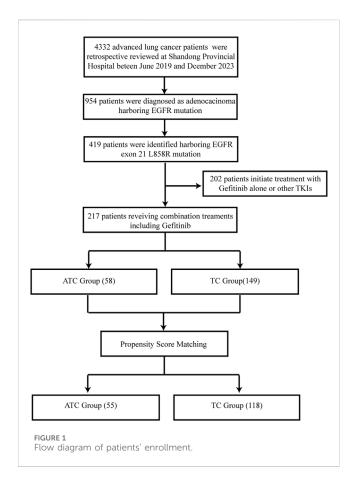
of Shandong Provincial Hospital between June 2019 and December 2021 were retrospectively screened. Clinical characteristics, such as age, gender, smoking status, and clinical stage, are extracted from medical records. The treatment process and efficacy evaluation are retroactively examined. Specifically selected are adult patients with the EGFR exon 21 L858R mutation identified through first- or second-generation sequencing, and an ECOG score of ≥2 in firstline treatment involving the combination of gefitinib and chemotherapy. Patients with concurrent malignant tumors, significant organ dysfunction, mental disorders, substance abuse history, or who died from other causes during follow-up are excluded. Additionally, patients are categorized into two groups based on the use of bevacizumab in the initial treatment regimen: the AGC group receiving anti-angiogenic drugs bevacizumab, gefitinib, and chemotherapy, and the GC group receiving gefitinib and chemotherapy.

2.2 Treatment methods

The chemotherapy protocol, incorporating Pemetrexed in conjunction with platinum agents like Carboplatin or Cisplatin, is delivered via intravenous infusion every 3–4 weeks over 4-6 cycles. The administration of pemetrexed and cisplatin was conducted at a dosage of 500 mg and 75 mg per square meter of body surface area, respectively. Carboplatin was administered based on an area under the concentration-time curve of 5 mg/mL. Bevacizumab and chemotherapy are administered concurrently by intravenous injection at a dosage of 15 mg/kg. After completion of the chemotherapy cycle, Bevacizumab is continued for maintenance treatment until disease progression occurs. Gefitinib is administered orally at a dose of 250 mg once daily after confirmation of the L858R mutation in the patient's genetic testing report.

2.3 Definitions and assessments

The response evaluation during treatment is evaluated through imaging studies, including computed tomography (CT) of the neck, chest, and upper abdomen, and magnetic resonance imaging (MRI) scans of the brain. During the chemotherapy cycles, imaging evaluations are conducted before the start of the second cycle of chemotherapy, and then every 2 cycles thereafter. After the completion of chemotherapy, imaging evaluations are conducted every 3 months. For patients without initial brain metastases, brain imaging is performed every 6 months or when clinical symptoms occur. Evaluations were conducted according to the Response Evaluation Criteria in Solid Tumors (RECIST, version 1.1). PFS was measured as the primary endpoint, while the OS, objective response rate (ORR), and disease control rate (DCR) were calculated as the secondary endpoints. Treatment-related adverse events (AEs) were graded and documented according to the Common Terminology Criteria for Adverse Events version 5.0 (CTCAE v5.0), covering hematological and non-hematological toxicities (e.g., general condition, gastrointestinal reactions) as well as targeted therapy-specific reactions (e.g., rash, paronychia, hemorrhage, and venous thrombosis). AE monitoring was conducted



throughout each treatment cycle. For patients followed up outside the hospital, standardized telephone interviews combined with electronic medical record systems were used for tracking.

2.4 Statistical analysis

The addition of bevacizumab to chemotherapy and gefitinib as a first-line treatment strategy is not a standard clinical practice, marked by a limited sample size within this cohort. Consequently, propensity score matching was employed to mitigate confounding variables. PSM analysis without replacement was conducted using the nearest neighbor method with the setting of a 1:4 matching ratio and a caliper of 0.1, proceeded by the matched package of R (Version 4.3.3). The Clinicopathologic data of patients, including age, ECOG score, smoking status, and metastatic sites were collected and entered into the PSM model. Patients' baseline data were compared using appropriate statistical methods both before and after matching. Categorical variables were compared using methods like Fisher's exact test or χ2 test, while continuous variables such as age were assessed using an independent t-test. Time-to-event variables such as the follow-up period and PFS were estimated using the Kaplan-Meier method, and comparisons of variation between curves were assessed using logrank tests. The Cox proportional-hazards model was carried out to analyze the response and treatment-by-covariate interaction effects of pre-specified patient subgroups as well as for univariate and multivariate analysis to assess the influence of potentially independent prognosis factors on survival functions. Statistical analyses were performed using GraphPad Prism 9. HR with 95% confidence intervals (CI) was presented, and a two-sided P-value <0.05 was considered statistically significant.

3 Results

3.1 Patients' baseline characteristics

Following the commencement of the study, we conducted a retrospective review of 4,332 lung cancer patients' electronic medical records at Shandong Provincial Hospital between June 2019 and December 2022. Among them, 954 were identified as metastatic lung adenocarcinoma patients with sensitizing EGFR mutations, including 217 with the L858R mutation, who underwent first-line treatment involving a combination therapy with gefitinib. The ATC group consisted of 58 patients, while the TC group included 149 patients. The overall patient enrollment process is shown in Figure 1. Upon propensity score matching, the patient numbers in the two groups were adjusted to 55 and 118, respectively. The baseline clinical characteristics of the patients before and after matching are presented in Table 1. Before matching, the median ages of the ATC group and TC group were 57 and 61 years, respectively, with the majority of patients aged 65 or younger. For the whole population, females and patients with no history of smoking formed the primary composition of both groups. The central nervous system and bones were identified as the predominant sites of metastasis. Before matching, the baseline data displayed statistical differences solely in ECOG scores between the two groups. Processing of PSM balanced clinical characteristics more with no significant variations noted among the diverse variables.

3.2 Tumor response

The median follow-up period for the entire population was 41.30 months. In the ATC group, the median follow-up was 41.91 months, while in the TC group, it was 40.89 months. Before PSM, within the first-line treatment, the ATC group witnessed 5 patients achieving complete response (CR), 39 achieving a partial response (PR), and 8 achieving stable disease, whereas in the TC group, these numbers were 7, 94, and 22, respectively. The ORR assessed in the overall cohort amounted to 70.0%, while the DCR reached a notable value of 84.5%. Upon comparing the response rates between the two groups, it was observed that the ATC group exhibited a higher ORR in contrast to the TC group (75.9% versus 67.8%, P = 0.26). The DCR was also higher in the ATC group, with rates of 89.7% compared to 82.6% in the TC group (P = 0.20). Following PSM, the ORR and DCR reached 68.8% and 83.2%, respectively. While the ATC group exhibited higher ORR (76.4% versus 65.3%, P = 0.14) and DCR (89.1% versus 80.1%, P = 0.16) values compared to the TC group, the difference was not statistically significant among the matched pairs (Table 2).

TABLE 1 Characteristics of baseline. IQR: interquartile range.

	Before matching			After matching		
	A + T + C	T + C	<i>P</i> -value	A + T + C	T + C	<i>P</i> -value
	(N = 58)	(N = 149)		(N = 55)	(N = 118)	
Age, median (IQR), years	57.0 [45.0, 78.0]	61.0 [38.0, 76.0]	0.09	57.0 [52.0, 64.0]	59.0 [55.0, 65.0]	0.57
>65 y	13 (22.4%)	34 (22.8%)	0.95	13 (23.6%)	21 (17.8%)	0.37
≤65 y	45 (77.6%)	115 (77.2%)		42 (76.4%)	97 (82.2%)	
Sex			0.88			1.00
Male	20 (34.5%)	48 (32.2%)		19 (34.5%)	41 (34.7%)	
Female	38 (65.5%)	101 (67.8%)		36 (65.5%)	77 (65.3%)	
ECOG score			< 0.01			0.06
0	28 (48.3%)	106 (71.1%)		28 (50.9%)	79 (66.9%)	
1	30 (51.7%)	43 (28.9%)		27 (49.1%)	39 (33.1%)	
Smoking Status			0.45			0.57
Never	16 (27.6%)	32 (21.5%)		16 (29.1%)	28 (23.7%)	
Former	42 (72.4%)	117 (78.5%)		39 (70.9%)	90 (76.3%)	
Brain Metastases			0.07			0.34
No	41 (70.7%)	83 (55.7%)		38 (69.1%)	71 (60.2%)	
Yes	17 (29.3%)	66 (44.3%)		17 (30.9%)	47 (39.8%)	
Bone Metastases			0.82			1.00
No	31 (53.4%)	84 (56.4%)		31 (56.4%)	67 (56.8%)	
Yes	27 (46.6%)	65 (43.6%)		24 (43.6%)	51 (43.2%)	
Pleural Effusion or Metastases			0.61			0.77
No	45 (77.6%)	122 (81.9%)		42 (76.4%)	94 (79.7%)	
Yes	13 (22.4%)	27 (18.1%)		13 (23.6%)	24 (20.3%)	
Intrapulmonary Metastasis			0.39			0.67
No	42 (72.4%)	118 (79.2%)		41 (74.5%)	93 (78.8%)	
Yes	16 (27.6%)	31 (20.8%)		14 (25.5%)	25 (21.2%)	
Adrenal Metastasis			0.48			1.00
No	55 (94.8%)	135 (90.6%)		52 (94.5%)	111 (94.1%)	
Yes	3 (5.2%)	14 (9.4%)		3 (5.5%)	7 (5.9%)	
Liver Metastases			0.61			0.73
No	47 (81.0%)	114 (76.5%)		44 (80.0%)	90 (76.3%)	
Yes	11 (19.0%)	35 (23.5%)		11 (20.0%)	28 (23.7%)	
Metastases of Other Sites	1		0.31			0.80
No	49 (84.5%)	135 (90.6%)		47 (85.5%)	104 (88.1%)	
Yes	9 (15.5%)	14 (9.4%)		8 (14.5%)	14 (11.9%)	

3.3 Progression-free survival

Before PSM, a total of 164 progress events (79.2%) were observed, with 42 patients (72.4%) in the ATC group and

122 patients (81.9%) in the TC group. The median PFS of the entire cohort was 18.77 months, whereas it was 22.18 months in the ATC group, showing a statistical difference compared to 16.16 months in the TC group (P < 0.01) (Supplementary Figure

TABLE 2 Summary of survival analysis and therapeutic effects of the two groups.

	Tar arrangers arra arranap		3			
	Before matching			After matching		
	ATC group	TC group	<i>P</i> -value	ATC group	TC group	<i>P</i> -value
	(N = 58)	(N = 149)		(N = 55)	(N = 118)	
mPFS (month)	22.18	16.16	<0.01	22.26	19.18	0.02
PFS Rate (%)						
6 months	96.6	85.9	0.06	98.2	84.7	0.02
12 months	81.8	61.7	0.05	78.2	64.4	0.09
18 months	61.8	43.0	0.06	58.2	50.0	0.38
mOS (month)	42.18	39.42	0.75	43.99	43.54	0.91
OS Rate (%)						
12 months	98.3	93.3	0.12	98.2	93.2	0.13
24 months	69.0	68.5	0.94	69.1	73.7	0.45
36 months	46.6	45.0	<0.01	43.6	49.2	0.43
ORR (%)	75.9	67.8	0.26	76.4	65.3	0.14
DCR (%)	89.7	82.6	0.20	89.1	80.1	0.16
Best response (case)						
Complete response	5	7	0.45	4	6	0.82
Partial response	39	94	0.58	38	71	0.26
Stable disease	8	22	0.86	7	18	0.66
Partial response	39	94	0.58	38	71	0.

S1A). The PFS rates of the patients in the ATC group at the 6th, 12th, and 18th months were higher than that in the TC group, but no statistical differences were observed (Table 2).

After PSM, the first-line PFS of patients in the ATC group demonstrated a significant improvement over that of the TC group patients (22.26 months versus 19.18 months, P = 0.02) (Figure 2A). Additionally, the PFS rates of the ATC group at the 6th, 12th, and 18th months (98.2%, 78.2%, 58.2%) exceeded those of the TC group (P < 0.05 for the 6th month) (Table 2). We also evaluated the differences in treatment outcomes in subgroups of patients with brain metastases, liver metastases, and intrapulmonary metastases. In patients with intrapulmonary metastases, the PFS was 25.37 months in the ATC group and 19.18 months in the TC group, showing a statistically significant difference (P = 0.036) (Figure 2B). However, no significant differences were observed in the subgroups of patients with central nervous system metastases (P = 0.286) and liver metastases (P = 0.902) (Figures 2C,D). The Cox regression model was used to analyze various statistical factors through univariate and multivariate analysis. The results showed that the ATC treatment regimen and brain metastasis were independent protective factors for first-line PFS in patients with the L858R mutation, providing evidence for the analysis of PFS results using the Kaplan-Meier method (Table 3).

3.4 Overall survival

At the date of cut-off, a total of 106 death events (79.2%) were observed for the whole population, with 28 patients (72.4%) in the ATC group and 78 patients (81.9%) in the TC group. The median OS of the entire cohort was 39.42 months, but no significant

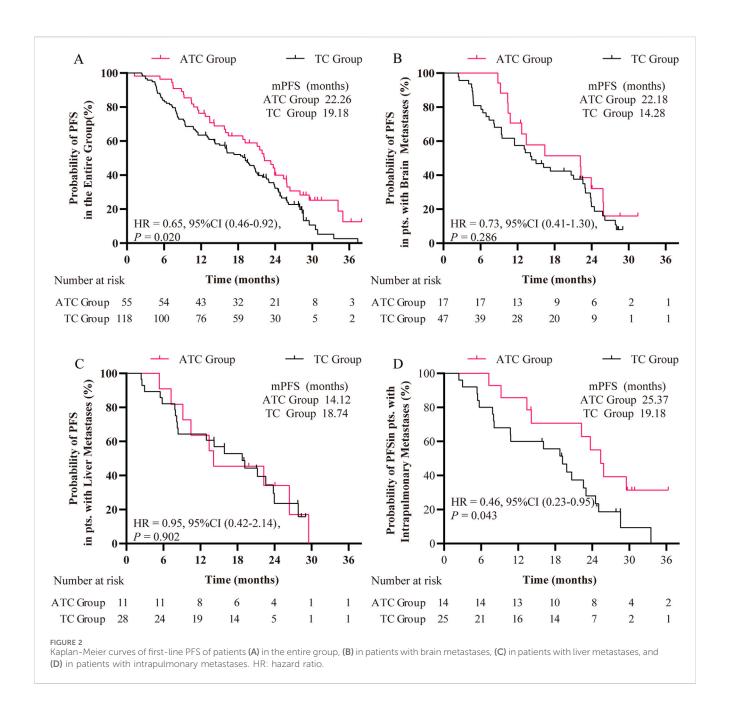
difference was observed between the ATC and TC groups (42.18 months versus 39.42 months, HR = 0.98, P = 0.91) (Supplementary Figure S1B). Following PSM, while the median OS of patients in the ATC group continues to exceed that of the TC group, no statistically significant difference exists between the survival curves (43.99 months versus 43.54 months, HR = 1.08, P = 0.75) (Figure 3A).

Regarding OS rate, prior to PSM, the ATC group exhibited higher OS rates than the TC group at the 12th, 24th, and 36th months, with a significant difference observed at 36th months (P < 0.01) (Table 2). Following PSM, the OS rate of the ATC group surpassed that of the TC group at 12 months, but was lower at 24 and 36 months, with no statistically significant difference (Table 2).

When evaluating the impact of the two treatment regimens on OS in subgroups of patients with brain metastases, liver metastases, and lung metastases after PSM, no statistical differences were observed (Figures 3B–D). After Cox univariate and multivariable regression analyses, no independent factors influencing the OS of patients with the L858R mutation were found after univariate and multivariate analysis (Table 4).

3.5 Saftey

Table 5 summarizes the incidence of AEs during treatment in both groups after PSM. Various grades of myelosuppression were the most common AEs in this study. In the ATC group, the incidence of grade ≥ 3 anemia, leukopenia, neutropenia, and thrombocytopenia was 12.7%, 12.7%, 10.9%, and 9.1%, respectively, which was comparable to the TC



group. However, the incidence of grade 1-2 myelosuppression events was higher in the ATC group. Regarding effects on liver and kidney function, the overall incidence of AEs was similar between the two groups, but more severe liver dysfunction was more common in the ATC group (10.9% vs. 1.7%). The incidence of treatment-related gastrointestinal disorders, such as decreased appetite, nausea, vomiting, and diarrhea, was generally similar between the two groups. Oral mucositis, rash, and paronychia are characteristic AEs associated with gefitinib. In this study, the combination of bevacizumab and gefitinib did not increase the incidence of oral mucositis or paronychia but did increase the incidence of rash (grade 1-2: 69.1%; grade ≥ 3 : 10.9%). Due to the addition of bevacizumab, three patients (5.3%) in the ATC group developed grade ≥ 3 hypertension, and one patient (1.8%) experienced

hemoptysis. No AE-related deaths occurred in this study. Overall, the combination of bevacizumab, gefitinib, and chemotherapy led to an increased incidence of AEs across all grades, but these remained within an acceptable range.

4 Discussion

This retrospective study demonstrates that first-line administration of the anti-angiogenic agent bevacizumab combined with chemotherapy and TKIs in advanced NSCLC patients harboring EGFR-L858R mutations improves ORR and DCR, resulting in significantly prolonged PFS. To our knowledge, this represents the first investigation evaluating the efficacy of a comprehensive regimen integrating anti-angiogenic therapy,

TABLE 3 Univariate and multivariable analysis for first-line PFS.

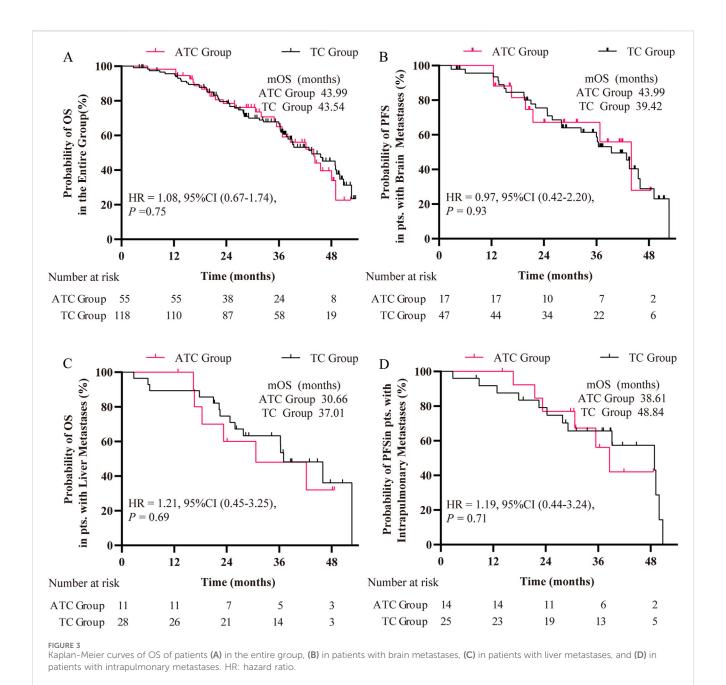
	Univariate analysis		Multivariate analysis			
	HR (95%CI)	<i>P</i> -value	HR (95%CI)	<i>P</i> -value		
Age						
≤65 y	1	0.81	1	0.87		
>65 y	1.06 (0.68-1.63)		1.04 (0.65-1.66)			
Sex						
Male	1	0.96	1	0.99		
Female	0.99 (0.69–1.42)		1.00 (0.60-1.69)			
Smoking st	atus					
Never	1	0.97	1	0.67		
Former	1.01 (0.69-1.48)		0.88 (0.51-1.55)			
ECOG-PS						
0	1	0.82	1	0.40		
1	0.96 (0.67–1.37)		1.17 (0.81-1.68)			
Treatment						
ATC Group	1	0.02	1	0.02		
TC Group	1.59 (1.09–2.33)		1.60 (1.09-2.35)			
Brain metas	stases					
Yes	1	0.03	1	0.03		
No	0.68 (0.48-0.97)		1.51 (1.03-2.21)			
Pleural met	astases or effusio	n				
Yes	1	0.86				
No	0.97 (0.65-1.44)					
Bone metas	stasis					
Yes	1	0.24				
No	1.23 (0.87–1.73)					
Itrapulmona	ary metastasis					
Yes	1	0.54				
No	1.14 (0.76–1.71)					
Adrenal metastasis						
Yes	1	0.08	1	0.08		
No	0.55 (0.28–1.09)		1.94 (0.93-4.08)			
Liver metastasis						
Yes	1	0.34				
No	0.82 (0.54-1.24)					
Metastasis of other sites						
Yes	1	0.86				
No	0.96 (0.57–1.59)					

chemotherapy, and TKIs as initial treatment for this patient population.

L858R and 19Del constitute the predominant EGFR mutation subtypes sensitive to TKIs (Hsu et al., 2018). While both mutations exhibit differential clinical responses to TKI therapy, it is well established that patients harboring the L858R mutation experience inferior clinical outcomes (Lee et al., 2015; Sheng et al., 2016; Chan et al., 2022). Mechanistic insights into this differential response include structural and functional divergences between the two mutants. The variant protein structures exhibit distinct conformational changes, resulting in differential TKIbinding affinity. Drug-binding kinetics analyses confirm higher binding affinity of TKIs to the 19del mutant versus L858R (Carey et al., 2006). Additionally, EGFR amplification levels and phosphorylation sites exhibit inter-subtype heterogeneity, leading to differential downstream pathway activation (Okabe et al., 2007). Notably, L858R demonstrates significantly higher co-occurrence rates with compound EGFR mutations compared to 19del, potentially contributing to primary TKI resistance mechanisms (Zhao et al., 2023).

The development of tyrosine kinase inhibitors (TKIs) has transformed the management of non-small cell lung cancer (NSCLC) with epidermal growth factor receptor (EGFR) mutations. TKIs also show promise as nuclear medicine imaging tracers for NSCLC diagnosis (Fawwaz et al., 2024). Nevertheless, curative outcomes remain elusive due to population heterogeneity along with inevitable drug resistance challenges. In cases where side effects can be tolerated, combination therapy is preferred and is more effective in slowing down the progression of the disease. The commonly used TKI drugs have been confirmed to provide significantly higher benefits when used in combination with chemotherapy compared to single-agent therapy (Sequist et al., 2013; Wu et al., 2013; Hosomi et al., 2020; Planchard et al., 2023). Three randomized trials in Asian advanced NSCLC populations demonstrated that first-line platinum-based chemotherapy concurrent with gefitinib achieved PFS ranging from 15.8 to 20.9 months. Notably, the NEJ005 and NEJ009 trials reported median OS of 41.9 and 50.9 months, respectively (Sugawara et al., 2015; Cheng et al., 2016; Hosomi et al., 2020). In our cohort, patients receiving gefitinib plus chemotherapy exhibited a median PFS of 19.18 months and a median OS of 43.54 months, aligning with published findings.

Preclinical studies demonstrate significant functional crosstalk and synergistic effects between the EGFR and vascular endothelial growth factor (VEGF) pathways in NSCLC, evidenced both *in vitro* and *in vivo* (Le et al., 2021). EGFR and VEGF signaling converge on downstream pathways including RAS/RAF/MAPK and PI3K/AKT/mTOR (Osude et al., 2022). Transcriptional profiling reveals coordinated upregulation of VEGF and VEGFR-2 in EGFR-mutant tumors, suggesting bidirectional pathway crosstalk at transcriptional and translational levels (Hung et al., 2016). Pharmacologically, VEGFR-targeted agents enhance intratumoral penetration of small-molecule TKIs. Gefitinib notably suppresses VEGFR expression through both HIF-1-dependent and HIF1-independent mechanisms, amplifying antitumor efficacy (Wildiers et al., 2003; Pore et al., 2006). In EGFR-mutant NSCLC



cell lines, TKIs like erlotinib suppress VEGF transcription by destabilizing HIF-1 (Ciardiello et al., 2001). And erlotinib combined with bevacizumab reduces tumor microvessel density, inhibits tumor growth, and enhances antitumor efficacy versus monotherapy in EGFR-mutant xenografts (Naumov et al., 2009; Masuda et al., 2017).

Moreover, integrating bevacizumab with TKI therapy yields additional clinical benefits through synergistic suppression of tumor proliferation and metastasis. A Japanese trial demonstrated superior PFS with bevacizumab plus erlotinib versus erlotinib monotherapy (16.9 vs. 13.3 months; P=0.016), without a significant OS difference (50.7 vs. 46.2 months; P=0.97) (Saito et al., 2019; Kawashima et al., 2022). Similarly, a Chinese trial confirmed that simultaneous VEGFR and EGFR inhibition improves first-line PFS (17.9 vs. 11.2 months; P<0.001), with

particular benefit in L858R-mutant patients (19.5 vs. 9.7 months; P=0.001) (Zhou et al., 2021). Our data revealed that the bevacizumab/gefitinib/chemotherapy regimen significantly improved first-line PFS compared to gefitinib/chemotherapy alone (22.26 vs. 19.18 months; P=0.02), though without significant OS extension (43.35 vs. 43.54 months; P=0.91).

Most lung cancer patients present with distant metastases, a primary contributor to the disease's high mortality rate. Brain metastases, occurring in approximately 30% of EGFR-mutant patients, are independently associated with poorer prognosis (Baik et al., 2015; Dormieux et al., 2020). Our analysis further identifies brain metastasis as an independent negative prognostic factor in L858R-mutant patients. Gefitinib, a first-generation EGFR-TKI, demonstrates limited blood-brain barrier (BBB) penetration, restricting its efficacy against intracranial lesions (Ballard et al.,

TABLE 4 Univariate and multivariable analysis for OS.

	Univariate analysis		Multivariate analysis			
	HR (95%CI)	<i>P</i> -value	HR (95%CI)	<i>P</i> -value		
Age						
≤65 y	1	0.75	1	0.73		
>65 y	1.1 (0.63-1.92)		1.106 (0.629–1.944)			
Sex						
Male	1	0.42	1	0.88		
Female	1.21 (0.76-1.92)		0.951 (0.49-1.843)			
Smoking st	tatus					
Never	1	0.48	1	0.47		
Former	1.21 (0.71-2.04)		1.293 (0.647-2.582)			
ECOG-PS						
0	1	0.56				
1	1.14 (0.73-1.79)					
Treatment						
ATC Group	1	0.75	1	0.15		
TC Group	0.92 (0.57-1.49)		0.707 (0.442–1.131)			
Brain meta	stases	'				
Yes	1	0.08	1	0.04		
No	1.48 (0.95-2.31)		0.61 (0.383-0.97)			
Pleural me	tastases or effusi	on				
Yes	1	0.37				
No	0.78 (0.45-1.35)					
Bone meta	stases					
Yes	1	0.37				
No	0.82 (0.52-1.27)					
Intrapulmo	nary metastasis					
Yes	1	0.55				
No	1.17 (0.7–1.96)					
Adrenal metastasis						
Yes	1	0.03				
No	2.24 (1.07-4.7)					
Liver metastasis						
Yes	1	0.2				
No	1.38 (0.84-2.28)					

(Continued in next column)

TABLE 4 (Continued) Univariate and multivariable analysis for OS.

	Univariate	analysis	Multivariate analysis			
	HR (95%CI)	P-value	HR (95%CI)	P-value		
Metastasis of other sites						
Yes	1	0.97				
No	0.99 (0.51-1.92)					

2016; Wrona et al., 2018). A multicenter phase 3 trial revealed that chemotherapy combined with gefitinib significantly prolongs intracranial PFS, systemic PFS, and OS in patients with brain metastases (Hou et al., 2023). In this study, median PFS was 22.18 months in the ATC group versus 14.17 months in the TC arm (p = NS), suggesting equivalent therapeutic robustness. The absence of additional benefit from bevacizumab indicates nonsynergistic activity with this TKI-chemotherapy backbone. Notably, our data demonstrate that incorporating bevacizumab in patients with pulmonary metastases significantly extended PFS when added to gefitinib plus chemotherapy (25.37 vs. 19.18 months; P = 0.043), indicating enhanced responsiveness relative to other metastatic sites - consistent with prior reports (Lin et al., 2016).

Regarding treatment safety assessment, although the ATC regimen increased the risk of specific AEs due to the inclusion of bevacizumab, the majority of treatment-related toxicities remained controllable through dynamic monitoring and graded management strategies. Previous Phase III clinical data in East Asian NSCLC populations reported that gefitinib combined with platinum-based doublet chemotherapy induced myelosuppression (including leukopenia, anemia, and thrombocytopenia) with overall incidence rates ranging from 20% to 60%, and Grade 3-4 hematological toxicities occurring in approximately 10%-30% (Yoshimura et al., 2015; Han et al., 2017; Hosomi et al., 2020). In this study, the observed incidence of hematological toxicities—anemia (52.7%), leukopenia (56.4%), neutropenia and thrombocytopenia (41.8%)—was generally (49.1%),consistent with or slightly lower than those reported in the aforementioned prospective studies. Notably, skin toxicity, a characteristic AE of TKIs, showed a high cumulative incidence in this cohort (ATC group 65.5% vs TC group 57.6%). The increased incidence may be related to bevacizumab-induced vascular endothelial dysfunction and IgE-mediated delayed hypersensitivity. Additionally, bevacizumab-specific vascular toxicities were observed in the ATC group, including Grade 3 hypertension (3 cases, 5.3%) and Grade 3 bleeding (1 case, 1.8%), while no such severe AEs were recorded in the TC group. Compared with safety data from registered clinical trials such as NEJ026, JO25567, and ARTEMIS-CTONG1509, the incidence of Grade 3 hypertension (5.3%) and bleeding events (1.8%) in this study was relatively low, and overall toxicity remained within an acceptable range (Seto et al., 2014; Hosomi et al., 2020; Zhou et al., 2021). It should be noted that, due to the retrospective nature of real-world data collection, the prophylactic

TABLE 5 Treatment-emergent adverse events.

	A + T + C (N = 55)		T + C (N = 118)	
	All grades	Grade ≥3	All grades	Grade ≥3
Anemia	29 (52.7%)	7 (12.7%)	56 (47.5%)	13 (11.0%)
Leukopenia	31 (56.4%)	7 (12.7%)	61 (51.7%)	15 (12.7%)
Neutropenia	27 (49.1%)	6 (10.9%)	51 (43.2%)	12 (10.2%)
Thrombocytopenia	23 (41.8%)	5 (9.1%)	43 (36.4%)	7 (5.9%)
Hepatic aminotransferase increased	30 (54.5%)	6 (10.9%)	58 (49.2%)	2 (1.7%)
Renal Dysfunction	17 (30.9%)	3 (5.3%)	36 (30.5%)	4 (3.4%)
Decreased appetite	25 (45.5%)	2 (3.7%)	42 (35.6%)	3 (2.5%)
Vomiting	15 (27.3%)	2 (3.7%)	28 (23.7%)	1 (0.8%)
Nausea	21 (38.2%)	4 (7.3%)	48 (40.7%)	7 (5.9%)
Diarrhea	21 (38.2%)	3 (5.3%)	50 (42.4%)	5 (4.2%)
Stomatitis	20 (36.4%)	1 (1.8%)	36 (30.5%)	2 (1.7%)
Rash	36 (65.5%)	6 (10.9%)	68 (57.6%)	6 (5.1%)
Paronychia	16 (29.1%)	1 (1.8%)	28 (23.7%)	1 (0.8%)
Asthenia	33 (60.0%)	6 (10.9%)	53 (44.9%)	9 (7.6%)
Hypertension	27 (49.1%)	3 (5.3%)	28 (23.7%)	-
Hemorrhage	15 (27.3%)	1 (1.8%)	18 (15.3%)	-
Superficial vein thrombosis	9 (16.4%)	-	10 (8.5%)	-

use of supportive medications (e.g., G-CSF, recombinant human thrombopoietin) may have mitigated hematological toxicity severity. Meanwhile, recall bias regarding non-severe AEs might have led to underreporting of low-grade toxicities. These factors collectively represent potential reasons for the lower incidence of AEs observed in this study compared to previous clinical trials.

This study has several limitations. First, its single-center retrospective design precludes external validation and introduces inherent confounding biases despite covariate adjustment via propensity score matching (PSM). Second, the sample size remains limited due to infrequent clinical adoption of the bevacizumab/chemotherapy/gefitinib combination per current guidelines. Third, though economic factors critically influence treatment decisions, fluctuating drug pricing and dynamic reimbursement policy adjustments by China's National Healthcare Security Administration precluded formal cost-effectiveness analysis.

In conclusion, this study demonstrates that adding bevacizumab to gefitinib and chemotherapy was associated with improved PFS in patients with advanced EGFR L858R-mutant NSCLC, though without conferring a corresponding OS benefit. The exploratory nature of these results warrants prospective validation before clinical implementation. As novel agents enter the therapeutic landscape, rigorous investigation of optimized combination strategies will advance precision oncology paradigms. Scientifically robust validation through prospectively designed, large-scale randomized controlled trials remains imperative for substantiating these combinatorial approaches.

Data availability statement

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

Ethics statement

The studies involving humans were approved by Ethical Review Committee of Shandong Provincial Hospital. The studies were conducted in accordance with the local legislation and institutional requirements. Written informed consent for participation was not required from the participants or the participants' legal guardians/next of kin in accordance with the national legislation and institutional requirements.

Author contributions

LL: Investigation, Validation and Writing – review and editing. XZ: Investigation, Visualization and Writing – review and editing. JxL: Formal Analysis, Writing – original draft. YD: Investigation, Writing – review and editing. JaL: Investigation, Writing – review and editing. FZ: Data curation, Supervision, Writing – review and editing. AF: Methodology, Supervision, Writing – review and editing. XL: Data curation, Formal analysis, Writing – review and editing. ZY: Writing – review and editing.

Funding

The author(s) declare that financial support was received for the research and/or publication of this article. This work was supported by the Cancer Research Funding of CSCO-Qilu (YQL 2019-0149).

Acknowledgments

We thank all the patients and their family members for their permission to participate in this study.

Conflict of interest

The 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.

Generative AI statement

The author(s) declare that no Generative AI was used in the creation of this manuscript.

References

Baik, C. S., Chamberlain, M. C., and Chow, L. Q. (2015). Targeted therapy for brain metastases in EGFR-mutated and ALK-rearranged non-small-cell lung cancer. *J. Thorac. Oncol.* 10, 1268–1278. doi:10.1097/JTO.0000000000000015

Ballard, P., Yates, J. W. T., Yang, Z., Kim, D.-W., Yang, J. C.-H., Cantarini, M., et al. (2016). Preclinical comparison of osimertinib with other EGFR-TKIs in EGFR-mutant NSCLC brain metastases models, and early evidence of clinical brain metastases activity. *Clin. Cancer Res.* 22, 5130–5140. doi:10.1158/1078-0432.CCR-16-0399

Carey, K. D., Garton, A. J., Romero, M. S., Kahler, J., Thomson, S., Ross, S., et al. (2006). Kinetic analysis of epidermal growth factor receptor somatic mutant proteins shows increased sensitivity to the epidermal growth factor receptor tyrosine kinase inhibitor, erlotinib. *Cancer Res.* 66, 8163–8171. doi:10.1158/0008-5472.CAN-06-0453

Chan, S.-K., Choi, H. C.-W., and Lee, V. H.-F. (2022). Overall survival benefits of first-line treatments for Asian patients with advanced EGFR-mutated NSCLC harboring L858R mutation: a systematic review and network meta-analysis. *JTO Clin. Res. Rep.* 3, 100322. doi:10.1016/j.jtocrr.2022.100322

Cheng, Y., Murakami, H., Yang, P.-C., He, J., Nakagawa, K., Kang, J. H., et al. (2016). Randomized phase II trial of gefitinib with and without pemetrexed as first-line therapy in patients with advanced nonsquamous non-small-cell lung cancer with activating epidermal growth factor receptor mutations. *J. Clin. Oncol.* 34, 3258–3266. doi:10.1200/ ICO.2016.66.9218

Ciardiello, F., Caputo, R., Bianco, R., Damiano, V., Fontanini, G., Cuccato, S., et al. (2001). Inhibition of growth factor production and angiogenesis in human cancer cells by ZD1839 (Iressa), a selective epidermal growth factor receptor tyrosine kinase inhibitor. *Clin. Cancer Res.* 7, 1459–1465.

Dormieux, A., Mezquita, L., Cournede, P. H., Remon, J., Tazdait, M., Lacroix, L., et al. (2020). Association of metastatic pattern and molecular status in stage IV non-small cell lung cancer adenocarcinoma. *Eur. Radiol.* 30, 5021–5028. doi:10.1007/s00330-020-06784-v

Fawwaz, M., Mishiro, K., Purwono, B., Nishii, R., and Ogawa, K. (2024). Synthesis and evaluation of a rociletinib analog as prospective imaging double mutation L858R/T790M in non-small cell lung cancer. *J. Pharm. Pharmacogn. Res.* 12, 231–242. doi:10. 56499/jppres23.1743_12.2.231

Han, B., Jin, B., Chu, T., Niu, Y., Dong, Y., Xu, J., et al. (2017). Combination of chemotherapy and gefitinib as first-line treatment for patients with advanced lung adenocarcinoma and sensitive EGFR mutations: a randomized controlled trial. *Int. J. Cancer* 141, 1249–1256. doi:10.1002/ijc.30806

Han, B., Zheng, R., Zeng, H., Wang, S., Sun, K., Chen, R., et al. (2024). Cancer incidence and mortality in China, 2022. *J. Natl. Cancer Cent.* 4, 47–53. doi:10.1016/j.jncc.2024.01.006

Hosomi, Y., Morita, S., Sugawara, S., Kato, T., Fukuhara, T., Gemma, A., et al. (2020). Gefitinib alone *versus* gefitinib plus chemotherapy for non-small-cell lung cancer with

Any alternative text (alt text) provided alongside figures in this article has been generated by Frontiers with the support of artificial intelligence and reasonable efforts have been made to ensure accuracy, including review by the authors wherever possible. If you identify any issues, please contact us.

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/fphar.2025.1503171/full#supplementary-material

mutated epidermal growth factor receptor: NEJ009 study. J. Clin. Oncol. 38, 115–123. doi:10.1200/JCO.19.01488

Hou, X., Li, M., Wu, G., Feng, W., Su, J., Jiang, H., et al. (2023). Gefitinib plus chemotherapy vs gefitinib alone in untreated EGFR-mutant non-small cell lung cancer in patients with brain metastases: the GAP BRAIN open-label, randomized, multicenter, phase 3 study. *JAMA Netw. Open* 6, e2255050. doi:10.1001/jamanetworkopen.2022.55050

Hsu, W.-H., Yang, J. C.-H., Mok, T. S., and Loong, H. H. (2018). Overview of current systemic management of EGFR-Mutant NSCLC. *Ann. Oncol.* 29, i3–i9. doi:10.1093/annonc/mdx702

Hung, M.-S., Chen, I.-C., Lin, P.-Y., Lung, J.-H., Li, Y.-C., Lin, Y.-C., et al. (2016). Epidermal growth factor receptor mutation enhances expression of vascular endothelial growth factor in lung cancer. *Oncol. Lett.* 12, 4598–4604. doi:10. 3892/ol.2016.5287

Kawashima, Y., Fukuhara, T., Saito, H., Furuya, N., Watanabe, K., Sugawara, S., et al. (2022). Bevacizumab plus erlotinib versus erlotinib alone in Japanese patients with advanced, metastatic, EGFR-Mutant non-small-cell lung cancer (NEJ026): overall survival analysis of an open-label, randomised, multicentre, phase 3 trial. Lancet Respir. Med. 10, 72–82. doi:10.1016/S2213-2600(21)00166-1

Kumar, A., Petri, E. T., Halmos, B., and Boggon, T. J. (2008). Structure and clinical relevance of the epidermal growth factor receptor in human cancer. *J. Clin. Oncol.* 26, 1742–1751. doi:10.1200/JCO.2007.12.1178

Le, X., Nilsson, M., Goldman, J., Reck, M., Nakagawa, K., Kato, T., et al. (2021). Dual EGFR-VEGF pathway inhibition: a promising strategy for patients with EGFR-mutant NSCLC. *J. Thorac. Oncol.* 16, 205–215. doi:10.1016/j.jtho.2020.10.006

Lee, V. H. F., Tin, V. P. C., Choy, T., Lam, K., Choi, C., Chung, L., et al. (2013). Association of exon 19 and 21 EGFR mutation patterns with treatment outcome after first-line tyrosine kinase inhibitor in metastatic non-small-cell lung cancer. *J. Thorac. Oncol.* 8, 1148–1155. doi:10.1097/JTO.0b013e31829f684a

Lee, C. K., Wu, Y.-L., Ding, P. N., Lord, S. J., Inoue, A., Zhou, C., et al. (2015). Impact of specific epidermal growth factor receptor (EGFR) mutations and clinical characteristics on outcomes after treatment with EGFR tyrosine kinase inhibitors *versus* chemotherapy in EGFR-mutant lung cancer: a meta-analysis. *J. Clin. Oncol.* 33, 1958–1965. doi:10.1200/JCO.2014.58.1736

Lin, J. J., Cardarella, S., Lydon, C. A., Dahlberg, S. E., Jackman, D. M., Jänne, P. A., et al. (2016). Five-year survival in EGFR-mutant metastatic lung adenocarcinoma treated with EGFR-TKIs. *J. Thorac. Oncol.* 11, 556–565. doi:10.1016/j.jtho.2015.12.103

Masuda, C., Yanagisawa, M., Yorozu, K., Kurasawa, M., Furugaki, K., Ishikura, N., et al. (2017). Bevacizumab counteracts VEGF-Dependent resistance to erlotinib in an EGFR-Mutated NSCLC xenograft model. *Int. J. Oncol.* 51, 425–434. doi:10.3892/ijo. 2017.4036

Naumov, G. N., Nilsson, M. B., Cascone, T., Briggs, A., Straume, O., Akslen, L. A., et al. (2009). Combined vascular endothelial growth factor receptor and epidermal growth factor receptor (EGFR) blockade inhibits tumor growth in xenograft models of EGFR inhibitor resistance. *Clin. Cancer Res.* 15, 3484–3494. doi:10.1158/1078-0432. CCR-08-2904

Okabe, T., Okamoto, I., Tamura, K., Terashima, M., Yoshida, T., Satoh, T., et al. (2007). Differential constitutive activation of the epidermal growth factor receptor in non-small cell lung cancer cells bearing EGFR gene mutation and amplification. *Cancer Res.* 67, 2046–2053. doi:10.1158/0008-5472.CAN-06-3339

Osude, C., Lin, L., Patel, M., Eckburg, A., Berei, J., Kuckovic, A., et al. (2022). Mediating EGFR-TKI resistance by VEGF/VEGFR autocrine pathway in non-small cell lung cancer. *Cells* 11, 1694. doi:10.3390/cells11101694

Pao, W., Miller, V. A., Politi, K. A., Riely, G. J., Somwar, R., Zakowski, M. F., et al. (2005). Acquired resistance of lung adenocarcinomas to geftinib or erlotinib is associated with a second mutation in the EGFR kinase domain. *PLoS Med.* 2, e73. doi:10.1371/journal.pmed.0020073

Planchard, D., Jänne, P. A., Cheng, Y., Yang, J. C.-H., Yanagitani, N., Kim, S.-W., et al. (2023). Osimertinib with or without chemotherapy in EGFR-mutated advanced NSCLC. N. Engl. J. Med. 389, 1935–1948. doi:10.1056/NEJMoa2306434

Pore, N., Jiang, Z., Gupta, A., Cerniglia, G., Kao, G. D., and Maity, A. (2006). EGFR tyrosine kinase inhibitors decrease VEGF expression by both hypoxia-inducible factor (HIF)-1-independent and HIF-1-dependent mechanisms. *Cancer Res.* 66, 3197–3204. doi:10.1158/0008-5472.CAN-05-3090

Saito, H., Fukuhara, T., Furuya, N., Watanabe, K., Sugawara, S., Iwasawa, S., et al. (2019). Erlotinib plus bevacizumab versus erlotinib alone in patients with EGFR-Positive advanced non-squamous non-small-cell lung cancer (NEJ026): interim analysis of an open-label, randomised, multicentre, phase 3 trial. *Lancet Oncol.* 20, 625–635. doi:10.1016/S1470-2045(19)30035-X

Sequist, L. V., Yang, J. C.-H., Yamamoto, N., O'Byrne, K., Hirsh, V., Mok, T., et al. (2013). Phase III study of afatinib or cisplatin plus pemetrexed in patients with metastatic lung adenocarcinoma with EGFR mutations. *J. Clin. Oncol.* 31, 3327–3334. doi:10.1200/JCO.2012.44.2806

Seto, T., Kato, T., Nishio, M., Goto, K., Atagi, S., Hosomi, Y., et al. (2014). Erlotinib alone or with bevacizumab as first-line therapy in patients with advanced non-squamous non-small-cell lung cancer harbouring EGFR mutations (JO25567): an open-label, randomised, multicentre, phase 2 study. *Lancet Oncol.* 15, 1236–1244. doi:10.1016/S1470-2045(14)70381-X

Sheng, M., Wang, F., Zhao, Y., Li, S., Wang, X., Shou, T., et al. (2016). Comparison of clinical outcomes of patients with non-small-cell lung cancer harbouring epidermal growth factor receptor exon 19 or exon 21 mutations after tyrosine kinase inhibitors treatment: a meta-analysis. *Eur. J. Clin. Pharmacol.* 72, 1–11. doi:10.1007/s00228-015-1966-0

Sugawara, S., Oizumi, S., Minato, K., Harada, T., Inoue, A., Fujita, Y., et al. (2015). Randomized phase II study of concurrent versus sequential alternating gefitinib and chemotherapy in previously untreated non-small cell lung cancer with sensitive EGFR mutations: NEJ005/TCOG0902. *Ann. Oncol.* 26, 888–894. doi:10.1093/annonc/mdv063

Sung, H., Ferlay, J., Siegel, R. L., Laversanne, M., Soerjomataram, I., Jemal, A., et al. (2021). Global cancer statistics 2020: GLOBOCAN estimates of incidence and mortality worldwide for 36 cancers in 185 countries. *CA Cancer J. Clin.* 71, 209–249. doi:10.3322/caac.21660

Wildiers, H., Guetens, G., De Boeck, G., Verbeken, E., Landuyt, B., Landuyt, W., et al. (2003). Effect of antivascular endothelial growth factor treatment on the intratumoral uptake of CPT-11. *Br. J. Cancer* 88, 1979–1986. doi:10.1038/sj.bjc.6601005

Wrona, A., Dziadziuszko, R., and Jassem, J. (2018). Management of brain metastases in non-small cell lung cancer in the era of tyrosine kinase inhibitors. *Cancer Treat. Rev.* 71, 59–67. doi:10.1016/j.ctrv.2018.10.011

Wu, Y.-L., Lee, J. S., Thongprasert, S., Yu, C.-J., Zhang, L., Ladrera, G., et al. (2013). Intercalated combination of chemotherapy and erlotinib for patients with advanced stage non-small-cell lung cancer (FASTACT-2): a randomised, double-blind trial. *Lancet Oncol.* 14, 777–786. doi:10.1016/S1470-2045(13)70254-7

Yoshimura, N., Kudoh, S., Mitsuoka, S., Yoshimoto, N., Oka, T., Nakai, T., et al. (2015). Phase II study of a combination regimen of gefitinib and pemetrexed as first-line treatment in patients with advanced non-small cell lung cancer harboring a sensitive EGFR mutation. *Lung Cancer* 90, 65–70. doi:10.1016/j.lungcan.2015. 06 002

Zhao, W., Song, A., Xu, Y., Wu, Q., Liu, C., Yin, J. C., et al. (2023). Rare mutation-dominant compound EGFR-Positive NSCLC is associated with enriched kinase domain-resided variants of uncertain significance and poor clinical outcomes. *BMC Med.* 21, 73. doi:10.1186/s12916-023-02768-z

Zhou, Q., Xu, C.-R., Cheng, Y., Liu, Y.-P., Chen, G.-Y., Cui, J.-W., et al. (2021). Bevacizumab plus erlotinib in Chinese patients with untreated, EGFR-mutated, advanced NSCLC (ARTEMIS-CTONG1509): a multicenter phase 3 study. *Cancer Cell* 39, 1279–1291.e3. doi:10.1016/j.ccell.2021.07.005