

ORIGINAL RESEARCH

Real-World Data Of Osimertinib In Patients With Pretreated Non-Small Cell Lung Cancer: A Retrospective Study

This article was published in the following Dove Press journal: Cancer Management and Research

Yuxin Mu Puyuan Xing Xuezhi Hao Yan Wang Junling Li

National Cancer Center, National Clinical Research Center for Cancer, Cancer Hospital, Chinese Academy of Medical Sciences and Peking Union Medical College, Beijing, People's Republic of China **Purpose:** Osimertinib is an oral, irreversible epidermal growth factor receptor (EGFR) tyrosine kinase inhibitor (TKI) targeted for both EGFR sensitizing mutations and T790M resistance mutation in patients with non-small-cell lung cancer (NSCLC). We assessed efficacy and safety of osimertinib in patients with pretreated NSCLC in a real-world setting. **Patients and methods:** Ninety-four patients with advanced NSCLC who received osimertinib after progression of prior EGFR-TKIs or chemotherapy treatments were retrospectively collected.

Results: In patients evaluable for response analysis (n = 91), overall objective response rate (ORR) was 47.3%, and disease control rate (DCR) was 90.1%. Median duration of response (DoR) in responding patients was 12.5 months (95% confidence interval [CI], 10.7 to 14.3). Median progression-free survival (PFS) was 8.5 months (95% CI, 7.4 to 9.6) in 2nd line group, 9.1 months (95% CI, 6.6 to 11.6) in ≥3rd line group, and 8.6 months (95% CI, 7.2 to 10.0) in overall population. For subgroup analysis, DCR and median PFS were 91.9% and 8.6 months (95% CI, 7.2 to 10.0) in patients with detectable T790M mutation, respectively, while 80.0% and 3.2 months (95% CI, 0.5 to 5.9) for those without. Median PFS was significantly longer for T790M-positive patients co-occurring with exon19del than with L858R (17.9 months vs 7.3 months; P<0.001). Among 45 patients with metastases to the central nervous system (CNS), median systemic PFS was 8.8 months (95% CI, 6.9 to 10.7), while intracranial time to progression (iTTP) was not reached. Safety profile was acceptable, no adverse events (AEs) related deaths was observed.

Conclusion: Osimertinib was highly active in patients with pretreated advanced NSCLC who harbored EGFR T790M mutation, with manageable side-effects.

Keywords: osimertinib, non-small-cell lung cancer, efficacy, safety

Introduction

Non-small-cell lung cancer (NSCLC) accounts for approximately 80% of all lung cancer cases. Approximately 10–15% of Caucasian patients and 30–40% of East Asian patients diagnosed with advanced NSCLC harbor activating epidermal growth factor receptor (EGFR) mutations. EGFR tyrosine kinase inhibitors (TKIs) gefitinib, erlotinib and afatinib are recommended as standard first-line treatment for such patients based on several large-scale prospective clinical trials. EGFR T790M mutation has been identified as the most commonly mechanism of acquired resistance to first-line EGFR-TKIs which were found in approximately 60% of patients. Osimertinib is a third-generation oral, potent, and irreversible EGFR-TKI, which inhibits both EGFR activating mutation

Correspondence: Junling Li
National Cancer Center, National Clinical
Research Center for Cancer, Cancer
Hospital, Chinese Academy of Medical
Sciences and Peking Union Medical
College, Number 17 Panjiayuan Nan Li,
Chao Yang District, Beijing 100021,
People's Republic of China
Tel +86 13801178891
Email lijunling@cicams.ac.cn

and T790M mutation.7 Phase I/II AURA trial reported osimertinib reached an objective response rate (ORR) of 61% and median progression-free survival (PFS) of 9.6 months among patients with T790M mutation.8 The phase III AURA3 study demonstrated the superiority of osimertinib over platinumpemetrexed chemotherapy in patients with T790M positive advanced NSCLC after progression of first-line EGFR-TKI therapy. Data of osimertinib in treatment naive patients of AURA study¹⁰ and FLAURA trial¹¹ also showed highly active of osimertinib in NSCLC patients with activating EGFR mutations. However, there was lack of real-world evidence to illustrate the effectiveness and safety of osimertinib which can reflect the current medical practice. We conducted this retrospective study to assess the real-world clinical impact of osimertinib in patients with advanced NSCLC in our Cancer Center.

Materials And Methods

Data Source And Study Population

The clinical data of patients with advanced NSCLC who received osimertinib after progression of prior EGFR-TKIs or chemotherapy treatments were retrospectively collected from our Cancer Center from Mar 1, 2017 to Jul 1, 2018. Eligible patients were required to be histologically or cytologically confirmed, locally advanced or metastatic NSCLC (stage IIIB and IV), detected with EGFR mutation at least once during disease courses and received osimertinib for at least 3 weeks. The patients who received osimertinib for less than 3 weeks were excluded as they received osimertinib for a short time without tumor response evaluation before molecular testing, and then switched to other regimens once they got the result of molecular testing, according to their medical data. A total of 94 patients met selection criteria.

Assessments

Primary endpoints were disease control rate (DCR) and PFS, secondary objectives included ORR and safety. ORR, DCR and PFS were assessed using Response Evaluation Criteria in Solid Tumor (RECIST) criteria (version 1.1). DCR was calculated as the percentage of patients with response of complete response (CR), partial response (PR), or stable disease (SD) lasting ≥6 weeks before any disease-progression event, while ORR pointed to CR or PR. Radiographic scan was performed to assess the tumor response every 8 to 12 weeks, including CT for chest/abdomen and MRI for brain lesions according to medical

records. All medical data were reviewed by a board-certified oncologist from our cancer center. PFS was defined as the time interval from the start of osimertinib treatment to progressive disease (PD) or death from any causes, whichever occurs first. Intracranial time to progression (iTTP) pointed to the time interval from the start of osimertinib to intracranial PD, regardless of extra-cranial response. Adverse events (AEs) were graded according to National Cancer Institute Common Terminology Criteria for Adverse Events (CTCAE, version 4.0).

Statistics Analysis

The data cutoff was Jul 30, 2018. Statistical analyses were performed with SPSS 23.0 statistical software, P-values were derived from two-sided tests and alpha=0.05 was used as significant level for all statistical testing. ORR and DCR were compared using Chi-square tests and Fisher's exact tests. PFS was analyzed using Kaplan–Meier method, survival curves of different subgroups were compared using log rank test.

Results

Patients And Characteristics

Ninety-four patients met selection criteria. Most patients but two received gene test prior to osimertinib, among whom 77 were T790M positive, 15 were T790M negative. A total of 59.6% (56/94) of the patients were women, 79.8% (75/94) were non-smokers, 56.4% (53/94) with Eastern Cooperative Oncology Group performance status (ECOG PS) 0, and 97.9% (92/94) had adenocarcinoma on histologic analysis. Most patients had received at least one prior EGFR-TKI (96.8%, 91/94), and 38.3% (36/94) had received prior cytotoxic chemotherapy. In the osimertinib ≥3rd treatment line group (n=38), in addition to EGFR-TKIs, 29 patients had received previous chemotherapy, most were pemetrexed-platinum-containing regimens (75.9%, 22/29), and 7/29 (24.1%) had received other anticancer regimens. Patient demographics and baseline characteristics were listed in Table 1.

Clinical Outcomes

At data cutoff, median duration of follow-up was 8.5 months, and 56 (59.6%) patients were still receiving osimertinib treatment.

Tumor Response

A total of 91 patients were evaluable for response analysis, 54 as 2nd line therapy and 37 as ≥3rd line. Of 91 patients,

Table 1 Baseline Patient Demographic And Clinical Characteristics (n=94)

2nd Line (n=56) ≥3rd Line (n=94) Total (n=94) Age, years Median Range 60 58 59 Range 41–86 33–86 33–86 Sex Male Female 24 (42.9) 14 (36.8) 38 (40.4) Female 32 (57.1) 24 (63.2) 56 (59.6) ECOG PS O 32 (57.1) 21 (55.3) 53 (56.4) 34 (36.2) 1 21 (37.5) 13 (34.2) 34 (36.2) 34 (36.2) 2 3 (5.4) 4 (10.5) 7 (7.4) 7 (7.4) Smoking status Nonsmoker 44 (78.6) 31 (81.6) 75 (79.8) Former/current smoker 12 (21.4) 7 (18.4) 19 (20.2) Histology Adenocarcinoma Squamous cell carcinoma 56 (100.0) 36 (94.7) 92 (97.9) Adenosquamous carcinoma Adenosquamous carcinoma 56 (100.0) 1 (2.6) 1 (1.1) Specimen for gene test Tissue Plasma 41 (73.2) 26 (68.4) 67 (71.3) CSF 2 (3.6) 1 (2.6) 3 (3.2) Pleural effusion 2 (3.6) 0 (0.0) 2 (2.1) None† 0 2 (3.6) 0 (0.0)<	Characteristics	No. Of Patients (%)				
Median Range 60 58 59 Range 41–86 33–86 33–86 Sex Male 24 (42.9) 14 (36.8) 38 (40.4) Female 32 (57.1) 24 (63.2) 56 (59.6) ECOG PS 0 32 (57.1) 21 (55.3) 53 (56.4) I 21 (37.5) 13 (34.2) 34 (36.2) 34 (36.2) 2 3 (5.4) 4 (10.5) 7 (7.4) Smoking status Nonsmoker 44 (78.6) 31 (81.6) 75 (79.8) Former/current 12 (21.4) 7 (18.4) 19 (20.2) Smoker 12 (21.4) 7 (18.4) 19 (20.2) Histology Adenocarcinoma 56 (100.0) 36 (94.7) 92 (97.9) Squamous cell 0 (0.0) 1 (2.6) 1 (1.1) carcinoma 56 (100.0) 36 (94.7) 92 (97.9) Specimen for gene test Tissue 11 (19.6) 9 (23.7) 20 (21.3) Plasma 41 (73.2) 26 (68.4) 67 (71.3) 22 (2.1)						
Range 41–86 33–86 33–86 Sex Male 24 (42.9) 14 (36.8) 38 (40.4) Female 32 (57.1) 24 (63.2) 56 (59.6) ECOG PS 0 32 (57.1) 21 (55.3) 53 (56.4) 1 21 (37.5) 13 (34.2) 34 (36.2) 34 (36.2) 2 3 (5.4) 4 (10.5) 7 (7.4) 7 (7.4) Smoking status Nonsmoker 44 (78.6) 31 (81.6) 75 (79.8) 75 (79.8) Former/current 12 (21.4) 7 (18.4) 19 (20.2) 7 (18.4) 19 (20.2) Histology Adenocarcinoma 56 (100.0) 36 (94.7) 92 (97.9) 92 (97.9) Squamous cell 0 (0.0) 1 (2.6) 1 (1.1) 1 (Age, years					
Sex Male Female 24 (42.9) Female 24 (63.2) 26 (59.6) ECOG PS 0 32 (57.1) 21 (55.3) 53 (56.4) 1 21 (37.5) 13 (34.2) 34 (36.2) 2 (37.5) 13 (34.2) 34 (36.2) 3 (5.4) 4 (10.5) 7 (7.4) Smoking status Nonsmoker 44 (78.6) Former/current 12 (21.4) 7 (18.4) 19 (20.2) Malenocarcinoma Adenocarcinoma Adenosquamous cell carcinoma Adenosquamous carcinoma Specimen for gene test Tissue 11 (19.6) Plasma 41 (73.2) 26 (68.4) 67 (71.3) CSF 2 (3.6) Pleural effusion None† 0 (0.0) 2 (5.3) 2 (2.1) Genotypes 1790M-positive exon19del-positive 19 14 28 (82.1) 15 (16.0) 20 (2.1) EBSR-positive exon19del/L858R- negative Unknown† 10 (0.0) 2 (5.3) 2 (2.1) Treatment history Geftinib Erlotinib 16 (28.6) 9 (23.7) 2 (25.6) 15 (39.5) 29 (30.9) Afatnib 0 (0.0) 5 (13.2) 5 (5.3)	Median	60	58	59		
Male Female 24 (42.9) 14 (36.8) 38 (40.4) Female 32 (57.1) 24 (63.2) 56 (59.6) ECOG PS 0 32 (57.1) 21 (55.3) 53 (56.4) 0 32 (57.1) 21 (55.3) 34 (36.2) 34 (36.2) 2 3 (5.4) 4 (10.5) 7 (7.4) Smoking status Nonsmoker 44 (78.6) 31 (81.6) 75 (79.8) Former/current smoker 12 (21.4) 7 (18.4) 19 (20.2) Histology Adenocarcinoma 56 (100.0) 36 (94.7) 92 (97.9) Squamous cell carcinoma 0 (0.0) 1 (2.6) 1 (1.1) Specimen for gene test Tissue 11 (19.6) 9 (23.7) 20 (21.3) Plasma 41 (73.2) 26 (68.4) 67 (71.3) CSF 2 (3.6) 1 (2.6) 3 (3.2) Pleural effusion 2 (3.6) 0 (0.0) 2 (2.1) Monety 46 (82.1) 31 (81.6) 77(81.9) Exonlydel/L858R-	Range	41–86	33–86	33–86		
Female 32 (57.1) 24 (63.2) 56 (59.6) ECOG PS 32 (57.1) 21 (55.3) 53 (56.4) 1 21 (37.5) 13 (34.2) 34 (36.2) 2 3 (5.4) 4 (10.5) 7 (7.4) Smoking status Nonsmoker 44 (78.6) 31 (81.6) 75 (79.8) Former/current smoker 12 (21.4) 7 (18.4) 19 (20.2) Histology Adenocarcinoma 56 (100.0) 36 (94.7) 92 (97.9) Adenocarcinoma 56 (100.0) 36 (94.7) 92 (97.9) Specimen for gene test 11 (19.6) 9 (23.7) 92 (97.9) Specimen for gene test 11 (19.6) 9 (23.7) 20 (21.3) Plasma 41 (73.2) 26 (68.4) 67 (71.3) CSF 2 (3.6) 1 (2.6) 3 (3.2) Pleural effusion 2 (3.6) 1 (2.6) 3 (3.2) Pleural effusion 2 (3.6) 7 (81.9) T790M-positive 46 (82.1) 31 (81.6) 77(81.9) Ex58R-positive 24	Sex					
ECOG PS 0 32 (57.1) 21 (55.3) 53 (56.4) 1 21 (37.5) 13 (34.2) 34 (36.2) 2 3 (5.4) 4 (10.5) 7 (7.4) Smoking status Nonsmoker 44 (78.6) 31 (81.6) 75 (79.8) Former/current 12 (21.4) 7 (18.4) 19 (20.2) Histology Adenocarcinoma 56 (100.0) 36 (94.7) 92 (97.9) Squamous cell 0 (0.0) 1 (2.6) 1 (1.1) carcinoma Adenosquamous 2 (10.0) 1 (2.6) 1 (1.1) Specimen for gene test 7 (11.1) 1 (1.1) Specimen for gene test 7 (1.1) 2 (21.4) 2 (21.4) 2 (21.4) Plasma 41 (73.2) 26 (68.4) 67 (71.3) CSF 2 (3.6) 1 (2.6) 3 (3.2) Pleural effusion 2 (3.6) 0 (0.0) 2 (2.1) None† 0 2 (53.3) 2 (2.1) Genotypes T790M-positive 46 (82.1) 31 (81.6) 77(81.9) exon 19del-positive 19 14 33 CSF 19del-positive 24 15 39 exon 19del-positive 24 15 39 exon 19del-positive 24 15 39 exon 19del-positive 27 5 (2.1) ENORTH 33 6 (8.6) 2 (3.2) 15 (16.0) exon 19del-positive 2 0 2 ENORTH 33 6 (8.6) 1 (2.6) 1 (2.6) 1 (2.6) Treatment history 6 (3.2) 2 (5.3) 2 (2.1) Treatment history 6 (3.2) 2 (5.3) 2 (2.1) Treatment history 6 (3.2) 3 (3.2) 3 (3.2) Frotinib 16 (28.6) 9 (23.7) 25 (26.6) Lotinib 14 (25.0) 15 (39.5) 29 (30.9) Afatinib 0 (0.0) 5 (13.2) 5 (5.3)	Male	24 (42.9)	14 (36.8)	38 (40.4)		
0 32 (57.1) 21 (55.3) 53 (56.4) 1 21 (37.5) 13 (34.2) 34 (36.2) 2 3 (5.4) 4 (10.5) 7 (7.4) Smoking status Nonsmoker 44 (78.6) 31 (81.6) 75 (79.8) Former/current 12 (21.4) 7 (18.4) 19 (20.2) Histology Adenocarcinoma 56 (100.0) 36 (94.7) 92 (97.9) Squamous cell 0 (0.0) 1 (2.6) 1 (1.1) Carcinoma Specimen for gene test Tissue 11 (19.6) 9 (23.7) 20 (21.3) Plasma 41 (73.2) 26 (68.4) 67 (71.3) CSF 2 (3.6) 0 (0.0) 2 (2.1) Pleural effusion 2 (3.6) 0 (0.0) 2 (2.1) Monety 0 2 (5.3) 2 (2.1) Genotypes T790M-positive 46 (82.1) 31 (81.6) 77(81.9) Ex58R-positive 2 1 1 exon 1	Female	32 (57.1)	24 (63.2)	56 (59.6)		
1	ECOG PS					
1	0	32 (57.1)	21 (55.3)	53 (56.4)		
2 3 (5.4) 4 (10.5) 7 (7.4) Smoking status Nonsmoker 44 (78.6) 31 (81.6) 75 (79.8) Former/current smoker 12 (21.4) 7 (18.4) 19 (20.2) Histology Adenocarcinoma 56 (100.0) 36 (94.7) 92 (97.9) Squamous cell carcinoma 0 (0.0) 1 (2.6) 1 (1.1) Adenosquamous carcinoma Specimen for gene test Tissue 11 (19.6) 9 (23.7) 20 (21.3) Plasma 41 (73.2) 26 (68.4) 67 (71.3) CSF 2 (3.6) 1 (2.6) 3 (3.2) Pleural effusion None† 2 (3.6) 0 (0.0) 2 (2.1) Genotypes T790M-positive 46 (82.1) 31 (81.6) 77(81.9) exon 19del-positive 24 15 39 exon 19del-positive 24 15 39 exon 19del-positive 2 0 2 L858R-positive 5 2 7 exon 19del-positive 2 0 2 L858R-positive	ı	l ` ´	` '	. , ,		
Nonsmoker 44 (78.6) 31 (81.6) 75 (79.8) Former/current smoker 12 (21.4) 7 (18.4) 19 (20.2)	2	1 ' '	, ,	` '		
Nonsmoker 44 (78.6) 31 (81.6) 75 (79.8) Former/current smoker 12 (21.4) 7 (18.4) 19 (20.2)	Smoking status					
Former/current smoker Histology Adenocarcinoma Squamous cell Carcinoma Adenosquamous Carcinoma Specimen for gene test Tissue Plasma Plasma Histology Adenocarcinoma Specimen for gene test Tissue Plasma Plasma Alt (73.2) Pleural effusion None† Genotypes T790M-positive exon19del-positive L858R-positive cavon19del-positive L858R-positive L858R-posit	-	44 (78.6)	31 (81.6)	75 (79.8)		
Histology 56 (100.0) 36 (94.7) 92 (97.9) Squamous cell carcinoma 0 (0.0) 1 (2.6) 1 (1.1) Adenosquamous carcinoma 0 (0.0) 1 (2.6) 1 (1.1) Specimen for gene test 11 (19.6) 9 (23.7) 20 (21.3) Plasma 41 (73.2) 26 (68.4) 67 (71.3) CSF 2 (3.6) 1 (2.6) 3 (3.2) Pleural effusion 2 (3.6) 0 (0.0) 2 (2.1) None† 0 2 (5.3) 2 (2.1) Genotypes 7790M-positive 46 (82.1) 31 (81.6) 77(81.9) exon19del-positive 24 15 39 exon19del/L858R-negative 10 (17.9) 5 (13.2) 15 (16.0) exon19del-positive 2 0 2 L858R-positive 5 2 7 exon19del/L858R-negative 3 3 6 Unknown† 0 (0.0) 2 (5.3) 2 (2.1) Treatment history 28 (50.0) 20 (52.6) 48 (51.1) E		l ` ´	` '	` ′		
Adenocarcinoma Squamous cell carcinoma Adenosquamous Carcinoma Adenosquamous Carcinoma Adenosquamous Carcinoma Specimen for gene test Tissue Plasma 41 (73.2) CSF 2 (3.6) Pleural effusion None† 0 0 0 0 0 0 0 0 0 0 0 0 0 0 1 0 0 0 1 0 0 0 1 0 0 0 0 1 0		12 (2)	7 (10.1)	17 (20.2)		
Adenocarcinoma Squamous cell carcinoma Adenosquamous Carcinoma Adenosquamous Carcinoma Adenosquamous Carcinoma Specimen for gene test Tissue Plasma 41 (73.2) CSF 2 (3.6) Pleural effusion None† 0 0 0 0 0 0 0 0 0 0 0 0 0 0 1 0 0 0 1 0 0 0 1 0 0 0 0 1 0	Histology					
carcinoma Adenosquamous carcinoma 0 (0.0) I (2.6) I (1.1) Specimen for gene test Tissue II (19.6) 9 (23.7) 20 (21.3) Plasma 41 (73.2) 26 (68.4) 67 (71.3) CSF 2 (3.6) I (2.6) 3 (3.2) Pleural effusion 2 (3.6) 0 (0.0) 2 (2.1) None† 0 2 (5.3) 2 (2.1) Genotypes T790M-positive 46 (82.1) 31 (81.6) 77(81.9) exon19del-positive 19 14 33 L858R-positive 24 15 39 exon19del/L858R- 3 2 5 negative 10 (17.9) 5 (13.2) 15 (16.0) exon19del/L858R- 3 3 6 negative 0 2 (5.3) 2 (2.1) Treatment history 28 (50.0) 20 (52.6) 48 (51.1) Erlotinib 16 (28.6) 9 (23.7) 25 (26.6) Icotinib 14 (25.0) 15 (39.5) 29 (30.9) Afatinib 0 (0.0) 5 (13.2	- ,	56 (100.0)	36 (94.7)	92 (97.9)		
Adenosquamous carcinoma Specimen for gene test Tissue	Squamous cell	0 (0.0)	1 (2.6)	1 (1.1)		
Carcinoma III (19.6) 9 (23.7) 20 (21.3) Plasma 41 (73.2) 26 (68.4) 67 (71.3) CSF 2 (3.6) 1 (2.6) 3 (3.2) Pleural effusion 2 (3.6) 0 (0.0) 2 (2.1) None† 0 2 (5.3) 2 (2.1) Genotypes 77(81.9) 14 33 1858R-positive 19 14 33 1858R-positive 24 15 39 exon19del/L858R- 3 2 5 negative 10 (17.9) 5 (13.2) 15 (16.0) exon19del/L858R- 3 3 6 exon19del/L858R- 3 3 6 negative Unknown† 0 (0.0) 2 (5.3) 2 (2.1) Treatment history 28 (50.0) 20 (52.6) 48 (51.1) Erlotinib 16 (28.6) 9 (23.7) 25 (26.6) Icotinib 14 (25.0) 15 (39.5) 29 (30.9)	carcinoma					
Specimen for gene test II (19.6) 9 (23.7) 20 (21.3) Plasma 41 (73.2) 26 (68.4) 67 (71.3) CSF 2 (3.6) I (2.6) 3 (3.2) Pleural effusion 2 (3.6) 0 (0.0) 2 (2.1) None† 0 2 (5.3) 2 (2.1) Genotypes 77(81.9) 14 33 Exon19del-positive 19 14 33 Exon19del/L858R- 3 2 5 negative 10 (17.9) 5 (13.2) 15 (16.0) exon19del-positive 2 0 2 Ex58R-positive 5 2 7 exon19del/L858R- 3 3 6 negative Unknown† 0 (0.0) 2 (5.3) 2 (2.1) Treatment history 28 (50.0) 20 (52.6) 48 (51.1) Erlotinib 16 (28.6) 9 (23.7) 25 (26.6) Icotinib 14 (25.0) 15 (39.5) 29 (30.9) Afatinib 0 (0.0) 5 (13.2) 5 (5	Adenosquamous	0 (0.0)	I (2.6)	1 (1.1)		
Tissue	carcinoma					
Plasma 41 (73.2) 26 (68.4) 67 (71.3) CSF 2 (3.6) 1 (2.6) 3 (3.2) Pleural effusion 2 (3.6) 0 (0.0) 2 (2.1) None† 0 2 (5.3) 2 (2.1) Genotypes T790M-positive 46 (82.1) 31 (81.6) 77(81.9) exon19del-positive 19 14 33 L858R-positive 24 15 39 exon19del/L858R- 3 2 5 negative 10 (17.9) 5 (13.2) 15 (16.0) exon19del/L858R- 3 3 6 negative 0 2 7 Unknown† 0 (0.0) 2 (5.3) 2 (2.1) Treatment history 28 (50.0) 20 (52.6) 48 (51.1) Erlotinib 16 (28.6) 9 (23.7) 25 (26.6) Icotinib 14 (25.0) 15 (39.5) 29 (30.9) Afatinib 0 (0.0) 5 (13.2) 5 (5.3)	Specimen for gene test					
CSF 2 (3.6) I (2.6) 3 (3.2) Pleural effusion 2 (3.6) 0 (0.0) 2 (2.1) None† 0 2 (5.3) 2 (2.1) Genotypes T790M-positive 46 (82.1) 31 (81.6) 77(81.9) exon19del-positive 19 14 33 L858R-positive 24 15 39 exon19del/L858R- 3 2 5 negative 10 (17.9) 5 (13.2) 15 (16.0) exon19del-positive 2 0 2 L858R-positive 5 2 7 exon19del/L858R- 3 3 6 negative Unknown† 0 (0.0) 2 (5.3) 2 (2.1) Treatment history 28 (50.0) 20 (52.6) 48 (51.1) Erlotinib 16 (28.6) 9 (23.7) 25 (26.6) Icotinib 14 (25.0) 15 (39.5) 29 (30.9) Afatinib 0 (0.0) 5 (13.2) 5 (53.3)	Tissue	11 (19.6)	9 (23.7)	20 (21.3)		
Pleural effusion 2 (3.6) 0 (0.0) 2 (2.1) None† 0 2 (5.3) 2 (2.1) Genotypes 77(81.9) 2 (2.1) T790M-positive 46 (82.1) 31 (81.6) 77(81.9) exon19del-positive 19 14 33 L858R-positive 24 15 39 exon19del/L858R- 3 2 5 negative 10 (17.9) 5 (13.2) 15 (16.0) exon19del-positive 2 0 2 L858R-positive 5 2 7 exon19del/L858R- 3 3 6 negative Unknown† 0 (0.0) 2 (5.3) 2 (2.1) Treatment history 28 (50.0) 20 (52.6) 48 (51.1) Erlotinib 16 (28.6) 9 (23.7) 25 (26.6) Icotinib 14 (25.0) 15 (39.5) 29 (30.9) Afatinib 0 (0.0) 5 (13.2) 5 (5.3)	Plasma	41 (73.2)	26 (68.4)	67 (71.3)		
None† 0 2 (5.3) 2 (2.1) Genotypes T790M-positive 46 (82.1) 31 (81.6) 77(81.9) exon19del-positive 19 14 33 L858R-positive 24 15 39 exon19del/L858R- 3 2 5 negative 10 (17.9) 5 (13.2) 15 (16.0) exon19del-positive 2 0 2 L858R-positive 5 2 7 exon19del/L858R- 3 3 6 negative Unknown† 0 (0.0) 2 (5.3) 2 (2.1) Treatment history Seffitinib 28 (50.0) 20 (52.6) 48 (51.1) Erlotinib 16 (28.6) 9 (23.7) 25 (26.6) Icotinib 14 (25.0) 15 (39.5) 29 (30.9) Afatinib 0 (0.0) 5 (13.2) 5 (5.3)	CSF	2 (3.6)	I (2.6)	3 (3.2)		
Genotypes T790M-positive exon19del-positive 19 14 33 L858R-positive exon19del/L858R- negative T790M-negative 10 (17.9) 5 (13.2) 15 (16.0) exon19del/L858R- 10 (17.9) 15 (16.0) 2 2 15 (16.0) 2 2 15 (16.0) 2 2 15 (16.0) 2 2 15 (16.0) 2 2 15 (16.0) 2 2 15 (16.0) 2 2 15 (16.0) 2 2 15 (16.0) 2 2 15 (16.0) 2 2 15 (16.0) 2 2 15 (16.0) 2 2 15 (16.0) 2 2 15 (16.0) 2 2 2 2 3 3 6 6 15 (20.0) 2 (20.1) Treatment history Gefitinib 28 (50.0) 20 (52.6) 48 (51.1) Erlotinib 16 (28.6) 9 (23.7) 25 (26.6) 1cotinib 14 (25.0) 15 (39.5) 29 (30.9) Afatinib 0 (0.0) 5 (13.2) 5 (5.3)	Pleural effusion	2 (3.6)	0 (0.0)	2 (2.1)		
T790M-positive 46 (82.1) 31 (81.6) 77(81.9) exon19del-positive 19 14 33 L858R-positive 24 15 39 exon19del/L858R- 3 2 5 negative 10 (17.9) 5 (13.2) 15 (16.0) exon19del-positive 2 0 2 L858R-positive 5 2 7 exon19del/L858R- 3 3 6 negative Unknown† 0 (0.0) 2 (5.3) 2 (2.1) Treatment history Sefitinib 28 (50.0) 20 (52.6) 48 (51.1) Erlotinib 16 (28.6) 9 (23.7) 25 (26.6) Icotinib 14 (25.0) 15 (39.5) 29 (30.9) Afatinib 0 (0.0) 5 (13.2) 5 (5.3)	None†	0	2 (5.3)	2 (2.1)		
exon19del-positive 19 14 33 L858R-positive 24 15 39 exon19del/L858R- 3 2 5 negative 10 (17.9) 5 (13.2) 15 (16.0) exon19del-positive 2 0 2 L858R-positive 5 2 7 exon19del/L858R- 3 3 6 negative Unknown† 0 (0.0) 2 (5.3) 2 (2.1) Treatment history Gefitinib 28 (50.0) 20 (52.6) 48 (51.1) Erlotinib 16 (28.6) 9 (23.7) 25 (26.6) Icotinib 14 (25.0) 15 (39.5) 29 (30.9) Afatinib 0 (0.0) 5 (13.2) 5 (5.3)	Genotypes					
L858R-positive exon19del/L858R- negative T790M-negative 10 (17.9) 5 (13.2) 15 (16.0) exon19del-positive 2 0 2 L858R-positive 5 2 7 exon19del/L858R- negative Unknown† 0 (0.0) 2 (5.3) 2 (2.1) Treatment history Gefitinib 28 (50.0) 20 (52.6) 48 (51.1) Erlotinib 16 (28.6) 9 (23.7) 25 (26.6) Icotinib 14 (25.0) 15 (39.5) 29 (30.9) Afatinib 0 (0.0) 5 (13.2) 5 (5.3)	T790M-positive	46 (82.1)	31 (81.6)	77(81.9)		
exon19del/L858R- negative 3 2 5 T790M-negative 10 (17.9) 5 (13.2) 15 (16.0) exon19del-positive 2 0 2 L858R-positive 5 2 7 exon19del/L858R- negative 3 3 6 Unknown† 0 (0.0) 2 (5.3) 2 (2.1) Treatment history Gefitinib 28 (50.0) 20 (52.6) 48 (51.1) Erlotinib 16 (28.6) 9 (23.7) 25 (26.6) Icotinib 14 (25.0) 15 (39.5) 29 (30.9) Afatinib 0 (0.0) 5 (13.2) 5 (5.3)	exon I 9del-positive	19	14	33		
negative 10 (17.9) 5 (13.2) 15 (16.0) exon19del-positive 2 0 2 L858R-positive 5 2 7 exon19del/L858R- 3 3 6 negative Unknown† 0 (0.0) 2 (5.3) 2 (2.1) Treatment history Gefitinib 28 (50.0) 20 (52.6) 48 (51.1) Erlotinib 16 (28.6) 9 (23.7) 25 (26.6) Icotinib 14 (25.0) 15 (39.5) 29 (30.9) Afatinib 0 (0.0) 5 (13.2) 5 (5.3)	L858R-positive	24	15	39		
T790M-negative 10 (17.9) 5 (13.2) 15 (16.0) exon I 9del-positive 2 0 2 L858R-positive 5 2 7 exon I 9del/L858R-negative 3 6 Unknown† 0 (0.0) 2 (5.3) 2 (2.1) Treatment history 28 (50.0) 20 (52.6) 48 (51.1) Erlotinib 16 (28.6) 9 (23.7) 25 (26.6) Icotinib 14 (25.0) 15 (39.5) 29 (30.9) Afatinib 0 (0.0) 5 (13.2) 5 (5.3)		3	2	5		
exon19del-positive 2 0 2 L858R-positive 5 2 7 exon19del/L858R-negative 3 3 6 Unknown† 0 (0.0) 2 (5.3) 2 (2.1) Treatment history Sefitinib 28 (50.0) 20 (52.6) 48 (51.1) Erlotinib 16 (28.6) 9 (23.7) 25 (26.6) Icotinib 14 (25.0) 15 (39.5) 29 (30.9) Afatinib 0 (0.0) 5 (13.2) 5 (5.3)	-					
L858R-positive		l ` ´		` '		
exon 19del/L858R-negative 3 3 6 Unknown† 0 (0.0) 2 (5.3) 2 (2.1) Treatment history Gefitinib 28 (50.0) 20 (52.6) 48 (51.1) Erlotinib 16 (28.6) 9 (23.7) 25 (26.6) Icotinib 14 (25.0) 15 (39.5) 29 (30.9) Afatinib 0 (0.0) 5 (13.2) 5 (5.3)	•					
negative Unknown† 0 (0.0) 2 (5.3) 2 (2.1) Treatment history Gefitinib 28 (50.0) 20 (52.6) 48 (51.1) Erlotinib 16 (28.6) 9 (23.7) 25 (26.6) Icotinib 14 (25.0) 15 (39.5) 29 (30.9) Afatinib 0 (0.0) 5 (13.2) 5 (5.3)	·					
Unknown† 0 (0.0) 2 (5.3) 2 (2.1) Treatment history Sefitinib 28 (50.0) 20 (52.6) 48 (51.1) Erlotinib 16 (28.6) 9 (23.7) 25 (26.6) Icotinib 14 (25.0) 15 (39.5) 29 (30.9) Afatinib 0 (0.0) 5 (13.2) 5 (5.3)		3	3	6		
Gefitinib 28 (50.0) 20 (52.6) 48 (51.1) Erlotinib 16 (28.6) 9 (23.7) 25 (26.6) Icotinib 14 (25.0) 15 (39.5) 29 (30.9) Afatinib 0 (0.0) 5 (13.2) 5 (5.3)	-	0 (0.0)	2 (5.3)	2 (2.1)		
Gefitinib 28 (50.0) 20 (52.6) 48 (51.1) Erlotinib 16 (28.6) 9 (23.7) 25 (26.6) Icotinib 14 (25.0) 15 (39.5) 29 (30.9) Afatinib 0 (0.0) 5 (13.2) 5 (5.3)	Treatment history					
Erlotinib 16 (28.6) 9 (23.7) 25 (26.6) Icotinib 14 (25.0) 15 (39.5) 29 (30.9) Afatinib 0 (0.0) 5 (13.2) 5 (5.3)		28 (50.0)	20 (52.6)	48 (51.1)		
Icotinib 14 (25.0) 15 (39.5) 29 (30.9) Afatinib 0 (0.0) 5 (13.2) 5 (5.3)		1 ' '	, ,	, ,		
Afatinib 0 (0.0) 5 (13.2) 5 (5.3)		1 ' '	` ′			
				` ′		
			, ,	` ′		
Chemotherapy 7 (12.5) 29 (76.3) 36 (38.3)		l ` ´	, ,	, ,		

(Continued)

Table I (Continued).

Characteristics	No. Of Patients (%)				
	2nd Line (n=56)	≥3rd Line (n=38)	Total (n=94)		
PD sites prior to osimertinib Intracranial only Others	10 (17.9) 46 (82.1)	10 (26.3) 28 (73.7)	20 (21.3) 74 (78.7)		

Note: †Two patients received osimertinib without gene tests.

Abbreviations: ECOG PS, Eastern Cooperative Oncology Group performance status; PD, progression disease; CSF, cerebrospinal fluid.

43 (47.3%) had PR, 39 (42.9%) had SD, 9 (9.9%) had PD. Disease control was achieved in 82 of 91 patients (90.1%), and ORR was 47.3%. DCR was similar between the 2nd line and ≥3rd line treatment groups, and between patients detecting EGFR T790M from plasma ctDNA samples and those with positive tissue-based outcomes. ORR was higher in patients detecting EGFR T790M from tumor samples than those with plasma ctDNA samples, but was not statistically significant (Table 2). Tumor responses were significantly different in terms of various genotypes. Of 74 patients with detectable T790M mutation who could be evaluated for response, ORR and DCR were 51.4% and 91.9%, respectively. In contrast, among 15 patients without detectable T790M mutation, ORR and DCR were 26.7% and 80.0%, respectively. As for T790M co-occurring common EGFR sensitizing mutation status, ORR was 66.7% in the T790M(+)/exon19del(+) group, which was significantly higher than 36.1% in the T790M(+)/L858R (+) group (P=0.011). DCR was 100.0% in the T790M (+)/exon19del(+) group, compared with 86.1% in the T790M(+)/L858R(+) group (P=0.055). Of 43 patients with an objective response, most had initial response at the time of first follow-up scan, with a median time to response of 1.2 months (range: 0.7-6.7).

PFS

At data cutoff, 53 of 94 (56.4%) patients had progressed or died. Median PFS was 8.6 months (95% CI, 7.2 to 10.0), 8.5 months (95% CI, 7.4 to 9.6), and 9.1 months (95% CI, 6.6 to 11.6) in the overall population, 2nd line group and ≥3rd line group, respectively. Subset analysis of PFS by T790M status showed a significant longer PFS in patients who was T790M positive compared with T790M negative population (median, 8.6 months [95% CI, 7.2 to 10.0] vs 3.2 months [95% CI, 0.5 to 5.9]; hazard ratio [HR], 0.51 [95%

Mu et al Dovepress

Table 2 Clinical Activity Summary Of Osimertinib

Types Of Patients	Survival		Tumor Response			
	PFS Months, 95% CI	P value	ORR %, 95% CI	P value	DCR %, 95% CI	P value
Overall population	8.6 (7.2, 10.0)		47.3 (36.8, 57.7)		90.1, (83.9, 96.4)	
Treatment line						
2nd (n=56) ≥3rd (n=38)	8.5 (7.4, 9.6) 9.1 (6.6, 11.6)		44.4, (30.8, 58.1)		90.7 (82.8, 98.7) 89.2 (78.7, 99.7)	
,	7.1 (6.6, 11.6)		51.4, (34.5, 68.2)		87.2 (78.7, 77.7)	
T790M status Positive (n=77) Negative (n=15)	8.6 (7.2, 10.0) 3.2 (0.5, 5.9)	0.041	51.4 (39.7, 63.0) 26.7 (1.3, 52.0)	0.081	91.9 (85.5, 98.3) 80.0 (57.1, 102.9)	0.356
Genotype		<0.001		0.011		0.055
T790M/exon19del (n= 33) T790M/L858R (n=39)	17.9 (5.4, 30.4) 7.3 (4.8, 9.8)	<0.001	66.7 (49.7, 83.6) 36.1 (19.6, 52.6)	0.011	100.0 (100.0, 100.0) 86.1 (74.2, 98.0)	0.033
T790M (+) sample		0.542		0.128		0.919
Tumor (n=19)	9.1 (4.1, 14.1)		68.4 (45.4, 91.4)		94.7 (83.7, 105.8)	
Plasma ctDNA (n=55)	8.5 (7.0, 10.0)		48.1 (34.0, 62.1)		90.4 (82.1, 98.7)	
CNS metastases		0.851				
Yes (n=45)	8.8 (6.9, 10.7)		_*		_*	
No (n=49)	7.8 (5.9, 9.7)					
Local therapy to CNS		0.566				
Yes (n=20)	8.5 (2.9, 14.2)		_*		_*	
No (n=25)	9.1 (6.5, 11.8)					
Smoking status		0.114		0.273		1.000
Never (n=75)	9.0 (8.3, 9.7)		50.0 (38.3, 61.7)		90.5 (83.7, 97.4)	
Ever (n=19)	6.5 (5.0, 8.0)		35.3 (10.0, 60.6),		88.2 (71.2, 105.3)	
Age		0.888		0.859		1.000
≤65 (n=61)	7.8 (5.9, 9.7)		46.6 (33.3, 59.8)		89.7 (81.6, 97.7)	
>65 (n=33)	8.8 (8.2, 9.4)		48.5 (30.5, 66.5)		90.9 (80.6, 101.3)	
Last treatment before osimertinib		0.238		0.148		1.000
EGFR-TKI (n=82)	8.5 (7.1, 9.9)		44.3 (33.1, 55.5)		89.9 (83.1, 96.7)	
Chemotherapy (n=12)	9.1 (1.5, 16.7)		66.7 (35.4, 98.0)		91.7 (73.3, 110.0)	

Note: *Tumor response to CNS was not collected.

Abbreviations: PFS, progression-free survival; CNS, central nervous system; ORR, objective response rate; DCR, disease control rate; EGFR-TKI, epidermal growth factor receptor tyrosine kinase inhibitor.

CI, 0.26 to 0.98]; P=0.041). Among patients with common EGFR sensitizing mutation exon19del or L858R, most received early-generation EGFR-TKIs prior to osimertinib. Of initial early-generation EGFR-TKIs treatment, subgroup analysis demonstrated a significant longer PFS in patients who harbored exon19del compared with L858R (median, 14.2 months [95% CI, 8.7 to 19.7] vs. 12.4 months [95% CI, 9.6 to 15.3]; HR, 0.53 (95% CI, 0.33 to 0.83); P=0.006). Similarly, of third-generation EGFR-TKI osimertinib, a significantly superior PFS was observed in patients harbored T790M mutation co-occurring with exon19del versus L858R (median, 17.9 months [95% CI, 5.4 to 30.4] vs. 7.3

months [95% CI, 4.8 to 9.8]; HR, 0.25 (95% CI, 0.11 to 0.54); P<0.001). Median PFS generally was statistically consistent across other subgroups analyzed: age at baseline (≤65 years vs. over 65 years), last treatment before osimertinib (EGFR-TKI vs. chemotherapy), and sample for detecting T790M status (tumor vs. plasma ctDNA). A nonsignificant trend toward longer PFS was observed in patients who were non-smokers compared with smokers (Table 2). Results of multivariate analysis for PFS showed that T790M/exon19del-positive contributed to significantly longer PFS (Table 3). The Kaplan-Meier curves of PFS was showed in Figure 1. Of 53 patients who had progressed of

Table 3 Cox Regression For PFS

Characteristics	Univariate	Univariate HR (95% CI) p-value		Multivariate		
	HR (95% CI)			p-value		
Male	1.68 (0.97, 2.92)	0.062	1.45 (0.68, 3.08)	0.326		
Age group (≤65)	0.95 (0.53, 1.72)	0.888	0.96 (0.49, 1.88)	0.925		
Smoker	1.65 (0.87, 3.11)	0.119	1.21 (0.53, 2.74)	0.645		
ECOG PS=0	0.67 (0.38, 1.16)	0.160	0.85 (0.45, 1.60)	0.619		
T790M-positive	0.51 (0.26, 0.98)	0.045	1.11 (0.28, 4.43)	0.878		
T790M/Exon I 9del-positive	0.26 (0.12, 0.57)	0.001	0.25 (0.11, 0.56)	0.001		
CNS metastases	1.05 (0.61, 1.81)	0.851	1.15 (0.61, 2.18)	0.655		

Abbreviations: PFS, progression-free survival; HR, hazard ratio; Cl, confidential interval; ECOG PS, Eastern Cooperative Oncology Group performance status; CNS, central nervous system.

osimertinib at data cut-off, 30 (56.6%) continued osimertinib beyond PD (20 of osimertinib monotherapy, 5 combined with local therapy, 3 combined with bevacizumab, 1 combined with icotinib and 1 combined with chemotherapy). Twenty-three (43.4%) patients discontinued osimertinib, among whom 13 switched to chemotherapy, 6 received early-generation EGFR-TKIs and 4 received best supportive care.

OS

Data on overall survival (OS) was immature. At data cutoff, 21 patients (22.3%) had died, 1 with T790M

(+)/exon19del(+), 14 with T790M(+)/L858R(+), and 6 with no detectable T790M mutation. One-year survival rate was 66.3%, a higher percentage of patients who were alive at data cutoff was observed in T790M (+)/exon19del(+) group (32/33, 97.0%) than in T790M(+)/L858R(+) group (25/39, 64.1%).

Osimertinib Activity In Patients With CNS Metastases

Forty-five patients had central nervous system (CNS) metastases at baseline, among whom 20 (44.4%) experienced local therapy to the brain (including surgery,

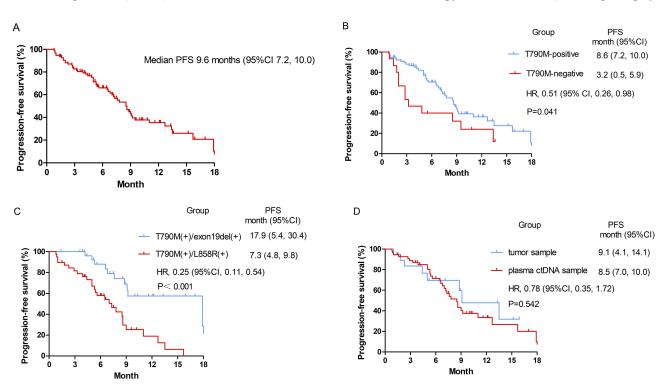


Figure I Progression-free Survival (PFS) in the overall population (**A**), in patients with or without detectable T790M mutation (**B**), in patients of T790M co-occurring with exon19 deletion or L858R mutation (**C**), in patients with T790M detected of tumor samples or plasma ctDNA samples (**D**). Tick marks indicate censored observations. **Abbreviations:** CI, confidence interval; HR, hazard ratio.

radiotherapy and intracranial injection) prior to osimertinib (≤6 months: n=14; >6 months, n=6). PFS was not inferior in patients with CNS metastases than in those without (median, 8.8 months [95% CI, 6.9 to 10.7] vs. 7.8 months [95% CI, 5.9 to 9.7]; HR, 1.05 (95% CI, 0.61 to 1.81); P=0.851). Of patients with locally treated CNS metastases subgroup, median PFS was 8.5 months (95% CI, 2.9 to 14.2), compared with 9.1 months (95% CI, 6.5 to 11.8) of patients without (HR, 1.26 [95% CI, 0.57 to 2.80]; P=0.566). Events of CNS progression were observed in 10 patients (22.2%) before discontinuation of osimertinib or death, iTTP was not reached.

Osimertinib Activity In Patients With Co-Occurring Mutation

Of 77 patients with detectable T790M mutation prior to osimertinib, 5 patients harbored co-occurring mutations at baseline including ROS1 exon36 mutation, EGFR exon7 mutation, EGFR exon18 G719C mutation, EGFR exon19 V742I mutation, KRAS mutation and MET amplification. Of these patients co-occurring with uncommon mutations, 2 achieved PR, 3 achieved SD, and events of progression were observed in 2 patients at data cutoff, PFS ranged from 2.7 months to 7.5 months (Table 4).

Safety

The most common AEs were rash (30.9%), fatigue (29.8%), and stomatitis (25.5%), and majority of AEs were grade 1 or 2 in severity. Eleven events of grade 3 were occurred, no grade 4 AEs or AE-related death were reported. AEs leading to dose reduction or drug discontinuation were not observed, but 4 (4.3%) patients experienced a dose interruption (one because of grade 3 stomatitis, one because of grade 2 dizziness, one because of grade 3 platelet count decreased, and one because of grade 3 alanine aminotransferase [ALT] increased and

aspartate aminotransferase [AST] increased). Summary of AEs were listed in Table 5.

Discussion

In our study, osimertinib showed a high ORR (47.3%) and DCR (90.1%) with superior median time to response of 1.2 months and median PFS of 8.6 months in patients with pretreated NSCLC. Our data suggested that osimertinib is promising in patients with advanced NSCLC, especially for those harboring T790M mutation or those with CNS metastases.

The primary endpoint DCR was 90.1% of our results in overall population and range from 80.0% to 100.0% in subgroups. This finding was consistent with prospective clinical trials of AURA,8 AURA extension,12 AURA 213 and AURA 39 studies. While ORR and PFS were inferior than those randomized studies. AURA 213 and AURA extension¹² phase II study demonstrated ORR of 60–70% and median PFS of 9.9-12.3 months. The AURA 3 phase III study⁹ reported a similar ORR of 71% and PFS of 10.1 months. The lower ORR and PFS in our study possibly were attributed to the reasons below: Our study population had characteristics that were different from the global population of patients. 38/94 (40.4%) patients in our study treated with osimertinib as ≥3rd line therapy and 17/94 (18.1%) without detectable T790M mutation or unknown, while osimertinib act as second line treatment in AURA 3 trial⁹ and study population were T790Mpositive. Besides, patients with CNS metastases accounts for about 33-41% of overall population in AURA 2,¹³ AURA extension¹² and AURA 3, 9 and only patients with CNS metastases could be enrolled if the disease was asymptomatic, stable, and not requiring corticosteroids for at least 4 weeks before osimertinib in these clinical trials. In addition, it is possible that lacking of blinded

Table 4 Osimertinib Activity In T790M-Positive Patients With Co-Occurring Mutation

Patient	Co-Occurring Mutation	Treatment Line	Tumor Response	PFS (months)	Status At Last Follow-Up
1	ROSI exon36 mutation	4	PR	7.5	PR
2	EGFR exon7 mutation EGFR exon18 G719C mutation	2	SD	2.7	PD
3	KRAS mutation	2	SD	2.9	PD
4	EGFR exon 19 V7421 mutation	2	PR	3.6	PR
5	MET amplification	2	SD	6.1	SD

Abbreviations: PR, partial response; SD, stable disease; PD, progressive disease; PFS, progression-free survival.

Table 5 Adverse Events (n=94)

Type Of AE	AE Grade				Total Report	Grade≥ 3
	ı	2	3	4	No. Of Patients (%)	
Rash	26	3	0	0	29 (30.9)	0
Fatigue	22	5	1	0	28 (29.8)	1
Stomatitis	20	2	2	0	24 (25.5)	2
Dry skin	22	0	0	0	22 (23.4)	0
White blood cell decreased	16	6	0	0	22 (23.4)	0
Paronychia	16	4	0	0	20 (21.3)	0
Diarrhea	17	2	0	0	19 (20.2)	0
Anorexia	12	4	3	0	19 (20.2)	3
Constipation	15	2	1	0	18 (19.1)	1
Neutrophil count decreased	9	7	0	0	16 (17.0)	0
CRE increased	15	0	0	0	15 (16.0)	0
Hypertriglyceridemia	10	3	0	0	13 (13.8)	0
Nausea	11	1	0	0	12 (12.8)	0
Anemia	5	5	1	0	11 (11.7)	1
Vomiting	8	1	0	0	9 (9.6)	0
Hypoalbuminaemia	7	0	0	0	7 (7.4)	0
Cough	6	0	0	0	6 (6.4)	0
Headache	5	1	0	0	6 (6.4)	0
Dizziness	5	1	0	0	6 (6.4)	0
Cholesterol high	6	0	0	0	6 (6.4)	0
AST increased	4	1	1	0	6 (6.4)	1
ALT increased	4	0	1	0	5 (5.3)	1
Platelet count decreased	1	3	1	0	5 (5.3)	1
Palpitation	4	0	0	0	4 (4.3)	0
Blood bilirubin increased	3	0	0	0	3 (3.2)	0
Hyponatremia	3	0	0	0	3 (3.2)	0
Hypocalcaemia	2	0	0	0	2 (2.1)	0
Hypokalemia	2	0	0	0	2 (2.1)	0
BUN increased	1	0	0	0	1 (1.1)	0

Abbreviations: AE, adverse events; ALT, alanine aminotransferase; AST, aspartate aminotransferase; CRE, creatinine; BUN, blood urea nitrogen.

independent central review (BICR) and evaluation of efficacy of osimertimib by different treating oncologists in our retrospective study may have impacted ORR outcomes, and for this reason, we used DCR and PFS as our primary objectives.

Prior to osimertinib, the standard recommendation for patients who progress after first-line EGFR-TKIs is chemotherapy, alternative treatment strategies explorations included afatinib, afatinib plus single-agent chemotherapy, and afatinib in combination with cetuximab. Those studies were associated with ORR of 7–34% and PFS of 2–6 months, with a high rate of toxic effects. The ORR, PFS, and safety profile with osimertinib in both the clinical trials 1,12,13 and our retrospective study were superior compared with historical results. These data support osimertinib for the treatment of patients with T790M-positive

advanced NSCLC after progression of prior EGFR-TKI therapy.

In our study, we noted a superior outcome in patients with co-occurring T790M and exon19del mutations than those with co-occurring T790M and L858R mutations documented before the osimertinib initiation. This was in line with previous evidence for greater clinical benefits of early-generation EGFR-TKIs in patients harboring exon19del versus L858R mutations. ¹⁹ The reasons are not clear, one possibility is that exon19del are more efficiently inhibited by EGFR-TKIs. However, in vitro studies do not support this hypothesis. ²⁰ Continued analyses are needed to answer this question.

Our study investigated the efficacy of osimertinib, regardless of T790M status. Patients with non-T790M-mediated resistance counted for approximately 40% of

cancers. For patients without detectable T790M mutation, osimertinib was associated with relatively low response rate and PFS, especially for those had received an EGFR-TKI as the last treatment regimen before osimertinib.8 Other approaches to address cancers that are resistant to EGFR-TKIs with non-T790M-dependent resistance mechanisms was chemotherapy, with similarly limited efficacy. 14 Confirmation of T790M status was carried out mostly by using plasma ctDNA samples in our study. Previous studies have demonstrated the feasibility of detecting EGFR T790M from plasma ctDNA samples, the results were equivalent to patients with positive tissuebased outcomes. However, a biopsy sample for patients with a plasma T790M-negative result after PD of first-line EGFR-TKI was still needed considering the false negative rates with plasma ctDNA T790M testing, as knowledge of truly T790M status is important both for the clinical practice and prognosis prediction.

Patients with EGFR mutated NSCLC have a much higher risk of developing CNS metastases.²¹ Osimertinib had demonstrated greater penetration of the Blood Brain Barrier (BBB) than gefitinib or afatinib in preclinical studies,²² and promising intracranial efficacy in patients with advanced NSCLC.^{9,23,24} In our study, the benefit of osimertinib in the subgroup of patients with CNS metastases was not inferior than those without. We also reported an encouraging systemic PFS of 9.1 months of CNS metastases patients without local treatment to the brain before or during osimertinib therapy, which showed a potential that this strategy may avoid patients from the long-term complications of brain radiation.²⁵

Our study had several strengths. The data were relatively new, and we included elderly patients, patients with ECOG PS > 1 and patients with symptomatic CNS metastases, which were usually excluded from prospective clinical trials. The selection of early-generation EGFR-TKIs in the first-line setting and the available samples for confirmation of mutation status reflects actual current real-world medical practice. The limitations of the study included its single-center, retrospective design, relatively small sample size, the lack of BICR, and the relatively short follow-up time to obtain OS and iTTP. Thus, a long-term follow-up and multicenter study would be required. Additionally, AEs of osimertinib in our study were retrospectively extracted from the medical records, which may introduce potential documentation bias, especially for non-laboratory findings.

In conclusion, our study shows that osimertinib provides encouraging clinical activity with a manageable safety profile in patients with pretreated advanced NSCLC, especially for whom T790M-mediated drug resistance had developed.

Ethical Statement

The Ethics Committee of the National Cancer Center/ Cancer Hospital, Chinese Academy of Medical Sciences and Peking Union Medical College approved this study (approval no. 15-144/1071). This study was conducted compliance with the Declaration of Helsinki, and we ensure the confidentiality of patient data. Since it was a retrospective study, some patients had already died before this study, and we were not able to get their informed consent. Besides, we didn't use any specific information on patients in this study. We get informed consent exemption approved by the ethics committee.

Disclosure

The authors report no conflicts of interest in this work.

References

- Miller KD, Siegel RL, Lin CC, et al. Cancer treatment and survivorship statistics, 2016. CA Cancer J Clin. 2016;66(4):271–289.
- Midha A, Dearden S, McCormack R. EGFR mutation incidence in non-small-cell lung cancer of adenocarcinoma histology: a systematic review and global map by ethnicity (mutMapII). Am J Cancer Res. 2015;5(9):2892–2911.
- Mok TS, Wu YL, Thongprasert S, et al. Gefitinib or carboplatinpaclitaxel in pulmonary adenocarcinoma. N Engl J Med. 2009;361 (10):947–957. doi:10.1056/NEJMoa0810699
- Rosell R, Carcereny E, Gervais R, et al. Erlotinib versus standard chemotherapy as first-line treatment for European patients with advanced EGFR mutation-positive non-small-cell lung cancer (EURTAC): a multicentre, open-label, randomised phase 3 trial. *Lancet Oncol.* 2012;13(3):239–246. doi:10.1016/S1470-2045(11) 70393-X
- Sequist LV, Yang JC, Yamamoto N, et al. Phase III study of afatinib or cisplatin plus pemetrexed in patients with metastatic lung adenocarcinoma with EGFR mutations. *J Clin Oncol*. 2013;31(27):3327–3334. doi:10.1200/JCO.2012.44.2806
- Yu HA, Arcila ME, Rekhtman N, et al. Analysis of tumor specimens at the time of acquired resistance to EGFR-TKI therapy in 155 patients with EGFR-mutant lung cancers. *Clin Cancer Res.* 2013;19(8):2240– 2247. doi:10.1158/1078-0432.CCR-12-2246
- Cross DA, Ashton SE, Ghiorghiu S, et al. AZD9291, an irreversible EGFR TKI, overcomes T790M-mediated resistance to EGFR inhibitors in lung cancer. *Cancer Discov.* 2014;4(9):1046–1061. doi:10.1158/2159-8290.CD-14-0337
- Janne PA, Yang JC, Kim DW, et al. AZD9291 in EGFR inhibitorresistant non-small-cell lung cancer. N Engl J Med. 2015;372 (18):1689–1699. doi:10.1056/NEJMoa1411817
- Mok TS, Wu Y, Ahn M, et al. Osimertinib or platinum-pemetrexed in EGFR T790M-positive lung cancer. N Engl J Med. 2017;376(7):629– 640. doi:10.1056/NEJMoa1612674

Ramalingam SS, Yang JC, Lee CK, et al. Osimertinib as first-line treatment of EGFR mutation-positive advanced non-small-cell lung cancer. *J Clin Oncol*. 2018;36(9):841–849. doi:10.1200/JCO.2017.74.7576

- Soria JC, Ohe Y, Vansteenkiste J, et al. Osimertinib in untreated EGFR-mutated advanced non-small-cell lung cancer. N Engl J Med. 2018;378(2):113–125. doi:10.1056/NEJMoa1713137
- Yang JC, Ahn MJ, Kim DW, et al. Osimertinib in pretreated T790M-positive advanced non-small-cell lung cancer: AURA study Phase II extension component. *J Clin Oncol*. 2017;35(12):1288–1296. doi:10.1200/JCO.2016.70.3223
- Goss G, Tsai CM, Shepherd FA, et al. Osimertinib for pretreated EGFR Thr790Met-positive advanced non-small-cell lung cancer (AURA2): a multicentre, open-label, single-arm, phase 2 study. *Lancet Oncol*. 2016;17(12):1643–1652. doi:10.1016/S1470-2045(16)30508-3
- Yoshida T, Kuroda H, Oya Y, et al. Clinical outcomes of platinum-based chemotherapy according to T790M mutation status in EGFR-positive non-small cell lung cancer patients after initial EGFR-TKI failure. *Lung Cancer*. 2017;109:89–91. doi:10.1016/j.lungcan.2017.05.001
- Janjigian YY, Smit EF, Groen HJ, et al. Dual inhibition of EGFR with afatinib and cetuximab in kinase inhibitor-resistant EGFR-mutant lung cancer with and without T790M mutations. *Cancer Discov*. 2014;4(9):1036–1045. doi:10.1158/2159-8290.CD-14-0326
- 16. Soria JC, Wu YL, Nakagawa K, et al. Gefitinib plus chemotherapy versus placebo plus chemotherapy in EGFR-mutation-positive non-small-cell lung cancer after progression on first-line gefitinib (IMPRESS): a phase 3 randomised trial. *Lancet Oncol.* 2015;16 (8):990–998. doi:10.1016/S1470-2045(15)00121-7
- 17. Miller VA, Hirsh V, Cadranel J, et al. Afatinib versus placebo for patients with advanced, metastatic non-small-cell lung cancer after failure of erlotinib, gefitinib, or both, and one or two lines of chemotherapy (LUX-Lung 1): a phase 2b/3 randomised trial. *Lancet Oncol.* 2012;13(5):528–538. doi:10.1016/S1470-2045(12)70087-6
- Schuler M, Yang JC, Park K, et al. Afatinib beyond progression in patients with non-small-cell lung cancer following chemotherapy, erlotinib/gefitinib and afatinib: phase III randomized LUX-Lung 5 trial. Ann Oncol. 2016;27(3):417–423. doi:10.1093/annonc/mdv597

- Jackman DM, Yeap BY, Sequist LV, et al. Exon 19 deletion mutations of epidermal growth factor receptor are associated with prolonged survival in non-small cell lung cancer patients treated with gefitinib or erlotinib. *Clin Cancer Res.* 2006;12(13):3908–3914. doi:10.1158/ 1078-0432.CCR-06-0462
- Paez JG, Janne PA, Lee JC, et al. EGFR mutations in lung cancer: correlation with clinical response to gefitinib therapy. *Science*. 2004;304(5676):1497–1500. doi:10.1126/science.1099314
- Rangachari D, Yamaguchi N, VanderLaan PA, et al. Brain metastases in patients with EGFR-mutated or ALK-rearranged non-small-cell lung cancers. *Lung Cancer*. 2015;88(1):108–111. doi:10.1016/j. lungcan.2015.01.020
- Ballard P, Yates JW, Yang Z, et al. 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. 2016;22(20):5130–5140. doi:10.1158/1078-0432.
 CCR-16-0399
- Reungwetwattana T, Nakagawa K, Cho BC, et al. CNS response to osimertinib versus standard epidermal growth factor receptor tyrosine kinase inhibitors in patients with untreated EGFR-mutated advanced non-small-cell lung cancer. J Clin Oncol. 2018;JCO2018783118.
- Xing P, Mu Y, Hao X, Wang Y, Li J. Data from real world to evaluate the efficacy of osimertinib in non-small cell lung cancer patients with central nervous system metastasis. *Clin Transl Oncol*. 2019;21:1424– 1431. doi:10.1007/s12094-019-02071-5
- Brown PD, Jaeckle K, Ballman KV, et al. Effect of radiosurgery alone vs radiosurgery with whole brain radiation therapy on cognitive function in patients with 1 to 3 brain metastases: a randomized clinical trial. *JAMA*. 2016;316(4):401–409. doi:10.1001/jama.2016. 9839

Cancer Management and Research

Publish your work in this journal

Cancer Management and Research is an international, peer-reviewed open access journal focusing on cancer research and the optimal use of preventative and integrated treatment interventions to achieve improved outcomes, enhanced survival and quality of life for the cancer patient.

The manuscript management system is completely online and includes a very quick and fair peer-review system, which is all easy to use. Visit http://www.dovepress.com/testimonials.php to read real quotes from published authors.

Submit your manuscript here: https://www.dovepress.com/cancer-management-and-research-journa

Dovepress