ORIGINAL ARTICLE



The Clinical Impact of Re-biopsies in Lung Adenocarcinoma: a Retrospective Multicenter Study

Pınar Akın Kabalak¹ • Derya Kızılgöz¹ • Suna Kavurgacı¹ • Nilgün Yılmaz Demirci² • Şenay Yılmaz³ • Güntülü Ak³ • Selma Metintaş^{3,4} • Muzaffer Metintaş³ • Funda Demirağ⁵ • Ülkü Yılmaz¹

Received: 3 April 2020 / Accepted: 9 July 2020 / Published online: 13 July 2020 \odot Arányi Lajos Foundation 2020

Abstract

If a patient's cancer progresses while undergoing targeted therapy, a re-biopsy is not mandatory. But when evaluating the benefits and risks on a case-by-case basis (transformation to small cell, assessing for a clinical trial), physicians should inform patients about the possible need for a re-biopsy (5). This was a retrospective and multicentre study. A total of 644 patients with lung adenocarcinoma were reviewed, 625 of whom were ruled eligible. From them, 399 were found to show disease progression, and 126 re-biopsies were performed. Progression status, re-biopsy sites, success of obtaining adequate tissue, molecular patterns after re-biopsy and subsequent treatments were analysed. Survival differences among patients with disease progression were then examined according to re-biopsy status. Overall, 625 patients with adenocarcinoma and a median age of 61.4 were evaluated. Initial tyrosine kinase inhibitor (TKI) usage numbered 37 patients (5.9%). Progression was diagnosed in 399 (63.8%) patients, out of which 26 (31.6%) underwent re-biopsies. The successful number of re-biopsies was 103 (81.7%). No complications were observed after any of the biopsy procedures. Subsequent treatments were changed in 15 patients (11.9%), who began new TKI treatments. Poor performance status was the most common reason for not performing a biopsy (n = 65; 23.8%), followed by the physician's decision (n = 40; 14.6%). Re-biopsies can demonstrate the new characteristics of a tumour and can detect the activation of pre-existing clones, making possible new treatment opportunities for patients. According to the performance status of the patient and the availability of the progressive lesion, we should increase the rate of re-biopsies before the decision to follow up with the best supportive care.

Keywords Non-small cell lung cancer · Re-biopsy · Targeted therapy

Background

Recent molecular developments in cancer biology have yielded a new era of targeted therapy. Among advanced stage

Pınar Akın Kabalak pinarakinn@yahoo.com

- ¹ Department of Chest Disease, Atatürk Chest Disease and Thoracic Surgery Teaching and Research Hospital, 06290 Ankara, Turkey
- ² Department of Chest Disease, Gazi University School of Medicine, Ankara, Turkey
- ³ Lung and Pleural Cancers Research and Clinical Center, Eskisehir Osmangazi University, Eskisehir, Turkey
- ⁴ School of Medicine, Department of Public Health, Eskisehir Osmangazi University, Eskisehir, Turkey
- ⁵ Department of Pathology, Atatürk Chest Disease and Thoracic Surgery Teaching and Research Hospital, Ankara, Turkey

non-small cell lung cancer (NSCLC) patients, the existence of targeted mutations, such as epidermal growth factor receptor (EGFR), anaplastic lymphoma kinase (ALK) and ROS protooncogene 1 (ROS-1), should be investigated at the beginning of their therapy (1). A re-biopsy for patients receiving EGFRtargeted tyrosine kinase inhibitors is strongly recommended when the disease progresses. Detecting any of the secondary resistance mechanisms can guide the selection of second- and third-generation EGFR tyrosine kinase inhibitors (EGFR-TKIs) for treatment in the advanced stages of lung adenocarcinoma [1].

ALK rearrangement is identified in 3–7% of advanced stage NSCLC cases. Until recently, there wasn't sufficient evidence concerning re-biopsies for ALK-rearranged NSCLC patients [2]. Some secondary point mutations have been described as causing acquired resistance to first-line ALK inhibitors (*crizotinib*) e.g., L1152R, C1156Y, F1174L, L1196M, L1198P, D1203N and G1269A [3]. However, many

studies have reported a wide variety of mutation types; unlike the EGFR mutant NSCLC (T790M), there is no predominant mutation for secondary ALK resistance. Acquired resistance mechanisms need to be better understood for recommending informed therapeutic guidance. Even if there is a good initial response to *crizotinib*, the median duration of response is only 11.3 months, and the transition to the central nervous system (CNS) is insufficient [4]. Due to an inadequate suppression of ALK and an overexpression of alternative point mutations, second-generation ALK inhibitors (alectinib, ceritinib, brigatinib and lorlatinib) were needed. The developed inhibitors were approved by the U.S. Food and Drug Administration (FDA) for the treatment of crizotinib-refractory ALKrearranged NSCLC patients with no need for testing for secondary mutations [2, 4].

The Second ESMO Consensus Conference on Lung Cancer reported the recommendation (with C grade strength) that performing a re-biopsy is not mandatory when there is a secondary resistance to TKIs and that patient-based risks and benefits should be considered [5]. Therefore, we aimed to investigate attitudes about re-biopsies and the procedure's success rates and safety after disease progression among Turkish patients with lung adenocarcinoma.

Materials and Methods

We retrospectively reviewed 644 consecutive patients who were diagnosed with adenocarcinoma between January 2011 and October 2018. The observational study included three centres: the Atatürk Chest Disease and Thoracic Surgery Training and Research Hospital, the Gazi University School of Medicine Department of Pulmonology and the Eskisehir Osmangazi University School of Medicine Department of Pulmonology. The study was approved by an institutional review board with the ethical committee number 509-24.11.2017.

Patients' baseline characteristics of age, gender, initial tumour, lymph node and metastasis classifications (TNM staging), treatment modalities, and mutational status were recorded. Their responses to first-line therapy were examined by computed tomography (CT) and evaluated using the Response Evaluation Criteria for Solid Tumours V.1.0 (RESIST V.1.0) [6]. When a progressive disease was identified, the most suitable biopsy procedure according to the lesion and patient's performance status was performed after informed consent was obtained from the patient. If a lesion was deemed to be eligible, various procedures, including computer-guided biopsy, pleural fluid cyto-block investigation, endobronchial ultrasound (EBUS), bronchoscopy (mucosal biopsy/transbronchial biopsy), liquid biopsy and biopsies from extra-pulmonary metastatic sites were performed. Statistical analyses were performed using IBM SPSS Statistics (IBM Corp., Armonk, NY) 15.0 program. The demographic characteristics of the study group were reported using descriptive statistics (frequencies, proportions, and means or medians). The appropriate chi-squared analysis was performed for comparisons between groups, while differences in time distributions between groups were tested for statistical significance using a log-rank test.

Results

A total of 644 patients were evaluated in this study. Nineteen patients were excluded due to insufficient data. A total of 625 eligible patients with lung adenocarcinoma, mean age $61.4 \pm$ 9.6 years, were included (Fig. 1). There was a male predominance (n = 480; 76.8%), and most patients were diagnosed to be in an advanced stage (n = 462; 73.9%). There were 37 (5.9%) patients being treated with TKIs. Progression of the disease was identified in 399 (63.8%) patients. Their baseline characteristics are summarized in Table 1.

From the patients diagnosed with progressive disease (PD) (n = 399), 126 (31.6%) underwent re-biopsies. The primary thoracic lesion was the most common site for performing the biopsies, including pleural cyto-block analyses (n = 95; 75.4%). Distant metastatic lesions were sampled in 25 (19.8%) patients, and 6 (4.8%) had liquid biopsies. No complications were observed after any of the biopsy procedures. A total of 103 (81.7%) specimens were suitable for molecular analyses. In 15 patients (11.9%), molecular profiles were found to have changed after their first re-biopsy, making them suitable for new TKI therapy (Table 2). Subsequent treatments for patients with progressive disease according to re-biopsy status are described in Table 2.

There was no difference between the re-biopsy and non-rebiopsy groups in terms of mean age (p = .145) or TNM staging distribution (p = .640). In the non-re-biopsy group, the most common reason for not performing a re-biopsy was patients' poor performance status (n = 65; 23.8%) followed by the physician's decision (n = 40; 14.6%) (Fig. 1). After the first progression, both in the re-biopsy and non-re-biopsy groups, second-line systemic chemotherapy and best supportive care (BSC) comprised most of the subsequent treatments (Table 3). Although the number of patients having subsequent treatment with TKIs was higher than the number of re-biopsy patients, some were continuing initial TKI treatment, while in a small number of patients, first-generation TKI was initiated without any targetable EGFR mutation. The distribution of secondline therapies did not show a significant difference between the two groups (Table 3).

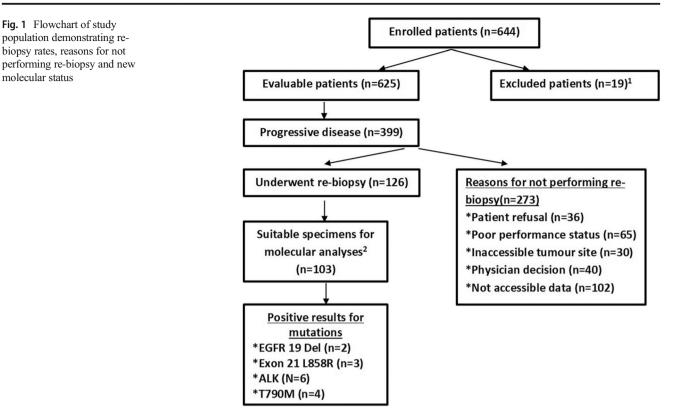


 Table 1
 General characteristics of study population

Characteristics	n = 625
Age, mean \pm SD, years	61.4±9.6
Gender, n (%)	145 (23.2)
- Female	480 (76.8)
- Male	
Stage, n (%)	192 (30.8)
I – II – III	433 (69.2)
IV	
Initial treatment, n (%)	462 (73.9)
- Chemotherapy	46 (7.4)
- Chemoradiotherapy	5 (0.8)
- Palliative radiotherapy	1 (0.2)
- Curative radiotherapy	37 (5.9)
- TKIs	24 (3.8)
- Surgery	50 (8.0)
- BSC	
Initial mutational status, n(%)	36 (5.8)
EGFR Exon Del19	19 (3.0)
EGFR Exon 21 L858R	34 (5.4)
ALK	0
ROS1	
Progressive disease, n (%)	399 (63.8)

SD: Standart deviation

TKIs: Thyrosine kinase inhibitors

BSC: Best supportive care

EGFR: Epidermal growth factor receptor

ALK: Anaplastic lymphoma kinase

ROS1: c-ros oncogene 1

Among all patients (n = 625), the frequency of targetable EGFR mutation (Exon Del19 and EGFR Exon 21 L858R) was 8.8% (n = 55) at the beginning of therapy. Among these cases, 26 patients received EGFR-TKI as a first-line treatment according to TNM stages and performance status. As described in Table 2, only 4 patients harbouring a targetable EGFR mutation revealed the T790M mutation after rebiopsy and continued with second-generation TKI.

The initial ALK positivity was 5.4% (n = 34 patients); 11 patients were using ALK inhibitors as a first-line treatment. Twenty-two of the ALK + patients had progressive disease, and only 1 patient underwent a re-biopsy resulting in showing ALK positivity again. The patient continued to receive first-generation ALK inhibitor treatment. The most common reasons for not performing re-biopsies in the remaining 21 patients was physician's decision (n = 13). In 3 patients, the lesion was not accessible, 4 patients had poor Eastern Cooperative Oncology Group (ECOG) performance status and 1 patient refused the biopsy procedure. Subsequent treatments for these 21 patients were chemotherapy (n = 8), BSC (n = 8) and palliative radiotherapy (n = 5).

Discussion

Currently, morphological and immune-histochemical diagnoses of lung adenocarcinoma can define the predominant Table 2

Stage	First-line treatment	Initial mutational status	Biopsy localisation	Mutational status after re-biopsy	Treatment after re-biopsy
IV	Chemotherapy	EGFR 21 exon (+)	primary lesion	T790M (+)	TKI
IV	Chemotherapy	Negative	primary lesion	ALK (+)	TKI
IV	Chemotherapy	Negative	primary lesion	ALK (+)	TKI
IV	Chemotherapy	Negative	primary lesion	EGFR exon 21(+)	TKI
III	Chemo-radiotherapy	Negative	primary lesion	ALK (+)	TKI
IV	Chemotherapy	Negative	primary lesion	ALK (+)	TKI
IV	Chemotherapy	Negative	primary lesion	EGFR 19 deletion	TKI
III	Chemo-radiotherapy	Negative	primary lesion	ALK (+)	TKI
IV	Chemotherapy	Negative	primary lesion	ALK (+)	TKI
IV	TKI	EGFR 19 deletion (+)	primary lesion	T790M (+)	TKI
IV	Chemotherapy	Negative	Distant metastasis	EGFR 21 exon (+)	TKI
III	TKI	EGFR 19 deletion (+)	primary lesion	T790M (+)	TKI
IV	TKI	EGFR 19 deletion (+)	primary lesion	T790M (+)	TKI
IV	Chemotherapy	Negative	primary lesion	EGFR 21 exon (+)	TKI
IV	Chemotherapy	Negative	primary lesion	EGFR 21 exon (+)	TKI

TKI: Thyrosine kinase inhibitor

EGFR: Epidermal growth factor receptor

ALK: Anaplastic lymphoma kinase

Crisotinib was the subsequent TKI for ALK positive patients

pattern according to differentiation (lepidic growth, acinar, papillary and micropapillary patterns and solid pattern with mucin); hence, detailed molecular analyses should be conducted at the beginning of lung cancer treatment [7] for utilizing adequate small biopsy samples to choose the most suitable therapeutic options for patients. Such analyses are all the more important because different morphological patterns produce differences in survival rates. For example, TKIs are novel therapeutic agents that are widely used in personalized medicine in lung adenocarcinoma with positive driver mutation [8].

Resistance to targeted therapy is the leading problem in adenocarcinomas. Unresponsiveness to TKIs is expected to occur approximately 1 year after starting them [9], at which time major problems begin. Even though there are some pathways explaining acquired resistance to first-line EGFR-TKIs, to demonstrate the new histopathological and molecular status of a tumour by acquiring a new biopsy specimen is a novel proposal in thoracic oncology. Among 139 lung cancer patients diagnosed with progressive disease while receiving EGFR-TKI treatment, re-biopsies were performed on 75 [9]. Their continuing rates of therapy with third-generation EGFR-TKI or other approved TKIs were significantly higher than for patients who did not have a re-biopsy. The most common reasons for not undergoing a repeat biopsy were poor medical condition, ineligible tumour site or physician's decision [9]. Similarly, in current study continuing rates of therapy with

Table 3	Subsequent treatments	for patients with	progressive disease	according to re-biopsy status

Characteristic of patients with positive targeted mutation after first re-biopsy

Subsequent regimen	Re-biopsy group $(n = 126)$	Non re-biopsy group $(n = 273)$	P value
Second line systemic chemotherapy	69 (54.7%)	141 (51.6%)	0.563
Third generation TKIs*	3 (2.3%)	-	-
TKIs (gefitinib, erlotinib, afatinib, crizotinib)	15 (11.9%)	19 (6.9%)	0.10
Best supportive care	37 (29.3%)	100 (36.6%)	0.155
Immunotherapy ¹	2 (1.5%)	1 (0.3%)	0.236
Palliative thorax radiotherapy*	-	12 (4.3%)	-

¹ Drugs were available within international-multicentre study

TKIs: Thyrosine kinase inhibitors

*: not suitable for statistical analyses

TKIs was higher in re-biopsy group and poor performance status and patient refusal were the common reasons not to perform biopsy procedure.

Re-biopsies for the ALK-positive patients did not reveal any clinically significant histopathological results. If at that time second-generation ALK inhibitors were available, it would have been a great opportunity for the ALK-resistant patients.

Only 4 EGFR-mutant patients revealed the T790M mutation and were subsequently treated with third-generation TKI. Overall, 11.9% (n = 15) patients continued with new targeted drugs in re-biopsy group. Similarly, a study including 65 lung adenocarcinoma patients demonstrated that, by having re-biopsies, 15% of the patients were given the chance to receive third-generation TKI, emphasizing the importance of performing re-biopsies and preserving extra frozen samples from the first diagnosis [10].

Not only for patients using first-line TKIs, but also for patients with a prior history of systemic chemotherapy (without any ALK or EGFR mutations), re-biopsies can offer guidance for new drug choices [11]. In our study, 11 patients with prior histories of no targetable mutations were subsequently treated with new targeted therapies after having re-biopsies (Table 2).

Due to its retrospective methodology, the absence of reasons for not performing re-biopsies in 102 patients should be considered a limitation of our study. In some cases, although the patients' performance status was good, the clinician's decision may have been that a biopsy was not required.

Tumour heterogeneity is a major point to consider. In a small specimen with a low mutated signal and higher tumour burden, the detection of targetable mutations can be difficult. Good clinical responses to EGFR-TKIs are possible if patients have targetable EGFR mutations in the bulk of the tumour cells. Additionally, DNA quality is another important factor when detecting mutations [12]. There are reports demonstrating a heterogeneity of EGFR mutations within the primary tumour and between primary tumour and metastases [13]. Regardless of the initial treatment, acquired resistance to first-line therapy should be re-assessed by re-sampling from either the primary tumour or an accessible metastatic site to determine the genotype of NSCLCs [12].

Although re-biopsies are not supported by a high level of evidence in guidelines, the repeat biopsy rate in this study, 126 patients (31.5%), is lower than expected. Designing the study retrospectively, starting to perform molecular analyses in the beginning of 2013, difficulties in accessing new drugs and different attitudes of health centres are the major factors for the low rate. According to a literature review article on rebiopsies in lung cancer, only 14 studies were available in the PubMed database. Despite that sometimes a patient with lung cancer is unsuitable for undergoing multiple re-biopsies, there is an obvious need for more research on re-biopsies in lung cancer [14].

Clinicians possess a variety of diagnostic tools to sample lung, pleura or mediastinal lymph nodes under current conditions. For example, a study comparing re-biopsy procedures in pre-treated lung cancer patients found an overall success rate of 90% for thoracic and extra-thoracic biopsies without any major complications [15].

Conclusions

This study demonstrated that re-biopsy is feasible and acceptable in NSCLC patients with advanced or metastatic disease. It is important to obtain adequate tissue to improve the success rates of molecular re-analyses. Demonstrating changed tumour behaviour and new molecular stratification can direct clinicians to initiate treatment with second- and thirdgeneration TKIs.

Acknowledgements Conception and design: Ülkü Yılmaz, MuzafferMetintaş, Güntülü Ak, Funda Demirağ

Data collection: Derya Kızılgöz, SunaKavurgacı, Şenay Yılmaz, Nilgün Yılmaz Demirci

Analysis: Selma Metintaş, Pınar Akın Kabalak,Nilgün Yılmaz Demirci

Writing: Pınar Akın Kabalak, Güntülü Ak, Şenay Yılmaz Guarantor: Ülkü Yılmaz Pathology: Funda Demirağ

Compliance with Ethical Standards

Conflicts of Interest The authors they have no conflicts of interest or received financial support for this study.

References

- Ettinger DS, Aisner DL, Wood DE, Akerley W, Bauman J, Chang JY et al (2018 Jul) NCCN Guidelines Insights: Non-Small Cell Lung Cancer, Version 5.2018. J Natl Compr Canc Netw 16(7): 807–821
- Lindeman NI, Cagle PT, Aisner DL, Arcila ME, Beasley MB, Bernicker EH et al (2018 Mar) Updated Molecular Testing Guideline for the Selection of Lung Cancer Patients for Treatment With Targeted Tyrosine Kinase Inhibitors: Guideline From the College of American Pathologists, the International Association for the Study of Lung Cancer, and the Association for Molecular Pathology. Arch Pathol Lab Med 142(3):321–346
- Huang D, Kim DW, Kotsakis A, Deng S, Lira P, Ho SN et al (2013) Multiplexed deep sequencing analysis of ALK kinase domain identifies resistance mutations in relapsed patients following crizotinib treatment. Genomics 102(3):157–162
- Vavalà T, Novello S (2018 Aug) Alectinib in the treatment of ALKpositive non-small cell lung cancer: an update on its properties, efficacy, safety and place in therapy. Ther Adv Med Oncol 3:10: 1758835918789364
- Besse B, Adjei A, Baas P, Meldgaard P, Nicolson M, Paz-Ares L et al (2014) 2nd ESMO Consensus Conference on Lung Cancer: non-small-cell lung cancer first-line/second and further lines of treatment in advanced disease. Ann Oncol 25(8):1475-84

- Therasse P, Arbuck SG, Eisenhauer EA, Wanders J, Kaplan RS, Rubinstein L et al (2000) New guidelines to evaluate the response to treatment in solid tumors. European Organization for Research and Treatment of Cancer, National Cancer Institute of the United States, National Cancer Institute of Canada. J Natl Cancer Inst 92: 205–216
- Thunnissen E (2012 Dec) Pulmonary adenocarcinoma histology. Transl Lung Cancer Res 1(4):276–279
- Rossi G, Pelosi G, Graziano P, Barbareschi M, Papotti M (2009) A reevaluation of the clinical significance of histological subtyping of non–small-cell lung carcinoma: Diagnostic algorithms in the era of personalized treatments. Int J Surg Pathol 17(3):206–18
- Kawamura T, Kenmotsu H, Taira T, Omori S, Nakashima K, Wakuda K et al (2016 Jul) Rebiopsy for patients with non-smallcell lung cancer after epidermal growth factor receptor-tyrosine kinase inhibitor failure. Cancer Sci 107(7):1001–1005
- Nosaki K, Inamasu E, Shimamatsu S, Yoshida T, Toyokawa G, Hirai F et al. Re-biopsy in advanced non-small cell lung cancer patients after the development of 3rd generation EGFR-TKI and new targeted therapies. https://doi.org/10.1200/jco.2015.33.15_ suppl.e19081

- Yoon HJ, Lee HY, Lee KS, Choi YL, Ahn MJ, Park K et al (2012 Dec) Repeat biopsy for mutational analysis of non-small cell lung cancers resistant to previous chemotherapy: adequacy and complications. Radiology 265(3):939–948
- Thunnissen E, Kerr KM, Herth FJV, Lantuejoul S, Papotti M, Rintoul RC et al (2012) The challenge of NSCLC diagnosis and predictive analysis on small samples. Practical approach of a working group. Lung Cancer 76:1–18
- 13. Yatabe Y (2010) EGFR mutations and the terminal respiratory unit. Cancer Metastasis Rev 29:23–36
- Jekunen AP (2015) Role of rebiopsy in relapsed non-small cell lung cancer for directing oncology treatments. J Oncol 2015:809835
- Hata A, Katakami N, Nanjo S, Okuda C, Kaji R, Imai Y. Rebiopsy of Histological Samples in Pretreated Non-small Cell Lung Cancer: Comparison Among Rebiopsy Procedures. In Vivo. 2017 31(3): 475–479

Publisher's Note Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.