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■ Research Article

Background of the need for targeted therapy options and platinum-based therapy responses in EGFR and ALK-mutated lung adenocarcinoma

EGFR ve ALK mutasyonu taşıyan akciğer adenokarsinomlarında platin bazlı tedavi yanitlari ve hedefe yönelik tedavi ihtiyacinin arka plani

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Abstract

Aim: To present our experience in EGFR and EML-4/ALK-mutated lung adenocarcinoma patients.

Material and Methods: 2580 patients were retrospectively evaluated. Only stage-4 lung adenocarcinoma patients who treated with at least 2-cycles of platinum-based regimens at frontline were included.

Results: Among 105 eligible patients, EGFR and EML-4/ALK mutations was detected in 14 and 4 patients. 75 were wild-type for both mutations. The median age and age of diagnose was 61 and 58.5, respectively. 81% was male and 78% was smoker. EGFR and EML-4/ALK-mutant patients were predominantly female and non-smoker (EGFR; $p=0.025$ and 0.002 , EML-4/ALK; $p=0.003$ and 0.012 , respectively). EML-4/ALK- mutant patients were significantly younger than EML-4/ALK wild-type ($p=0.02$) (Table 1). EGFR exon-19, 20 and 21 mutations were associated with liver, bone and pleural metastases, respectively ($p=0.046$, 0.05 and 0.035 , respectively). After firstline platinum-based chemotherapy, complete remission (CR) and partial remission (PR) rates were 4.7% and 24.6%, respectively. Concurrent radiotherapy and absence of bone metastases at diagnosis were significant factors influencing firstline platinum-based therapy responses ($p=0.004$ and $p=0.046$, respectively). EGFR or EML-4/ALK mutation status didn't show significant difference in terms of platinum-based treatment response ($p=0.933$ and 0.184 , respectively). Median progression-free survival (PFS) was 10 months. The observed effect of concurrent radiotherapy and the presence of bone metastases on treatment response didn't reflected in the PFS results ($p=0.079$ and 0.285 , respectively).

Conclusion: The presence of EGFR and ALK mutations does not effect the treatment response of platinum-based regimens. The association of EGFR exon subsets with metastasis points is worth investigating.

Keywords: EGFR, EML-4/ALK, ALK, platinum, lung adenocarcinoma

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Öz

Amaç: EGFR ve EML-4/ALK mutasyonlu akciğer adenokarsinomu hastalarındaki deneyimlerimizi sunmak.

Gereç ve Yöntemler: 2580 hasta retrospektif olarak değerlendirildi. Çalışmaya yalnızca evre 4 akciğer adenokarsinomu olup ilk sıra en az 2 siklus platin bazlı rejimlerle tedavi edilen hastalar dahil edilmiştir.

Bulgular: Çalışmaya uygun 105 vakanın 14'ü EGFR, 4'ü EML-4/ALK mutant iken 75 vaka her iki mutasyonu da taşıyordu. Medyan yaş ve tanı yaşı sırasıyla 61 ve 58.5 idi. %81'i erkekti ve %78'i sigara içiyordu. EGFR ve EML-4/ALK-mutant hastalar ağırlıklı olarak kadındı ve sigara içmiyordu (sırasıyla EGFR; $p=0.025$ ve 0.002 , EML-4/ALK; $p=0.003$ ve 0.012). EML-4/ALK-mutant hastalar, bu mutasyonu taşımayanlara göre daha gençti ($p=0,02$) (Tablo 1). EGFR ekson-19, 20 ve 21 mutasyonları sırasıyla karaciğer, kemik ve plevral metastazlarla ilişkiliydi (sırasıyla $p=0.046$, 0.05 ve 0.035). Birinci basamak platin bazlı kemoterapiden sonra tam remisyon ve kısmi yanıt oranları sırasıyla %4,7 ve %24,6 idi. Eşzamanlı radyoterapi ve tanı sırasında kemik metastazlarının olmaması birinci basamak platin bazlı tedavi yanıtını etkileyen faktörlerdi (sırasıyla $p=0.004$ ve $p=0.046$). EGFR veya EML-4/ALK mutasyon durumu platin bazlı tedavi yanıtı açısından anlamlı fark göstermemiştir (sırasıyla $p=0,933$ ve $0,184$). Medyan progresyonsuz sağkalım 10 ay iken eşzamanlı radyoterapi ve kemik metastazının tedavi yanıtı üzerinde gözlenen etkisi PFS sonuçlarına yansımamıştır (sırasıyla $p=0,079$ ve $0,285$).

Sonuçlar: EGFR ve ALK mutasyonlarının varlığı, platin bazlı rejimlerin tedavi yanıtını etkilememektedir. EGFR ekson alt gruplarının metastaz noktaları ile ilişkisi araştırılması gereken bir nokta olarak saptanmıştır.

Anahtar Kelimeler: EGFR, EML-4/ALK, ALK, platin, akciğer adenokarsinomu.

Introduction

Primary lung cancer is the second most common malignancy after non-melanoma skin cancer and is the most common cause of malignancy related death. Non-small cell lung cancers constitute 80-90% of primary lung cancers and due to the decrease in tobacco use in recent years, the frequency of squamous cell type has decreased while the adenocarcinoma type has become dominant [1]. Frontline platinum-based therapy response rates for lung adenocarcinoma have been reported to range from 30 to 40%. [2]. To increase these low response rates, EGFR and ALK gene mutations have emerged as important focal points in the development of targeted therapies.

Increased tyrosine kinase activity resulting from mutations effecting the tyrosine kinase domain of EGFR and inversion of the short arm of the chromosome 2, which causes the fusion of the ALK gene with the EML-4 gene, play an important role in the etiopathogenesis of non-small cell lung cancer. Therefore, current guidelines recommend routine screening of these genes regardless of the clinical status [1,3].

In this study, based on the rarity of these mutations and studies with limited number of patients, we aimed to present our single center experience and make a contribution to the literature.

Material and Methods

2580 lung cancer patients who applied to our clinic between 2007 – 2016 were retrospectively evaluated for the study. Only stage 4 lung adenocarcinoma patients who treated with at least 2 cycles of platinum-based regimens as firstline therapy and investigated for EGFR Exon 18-19-20-21 mutations and/or the EML-4/ALK translocation gene were included in the study. Patients treated with regimens other than platinum-based or without interim and/or end-of-treatment PET-CT imaging results were excluded. 12 patients were negative for EGFR mutations but their ALK mutation status was unknown. These patients also excluded from EGFR and EML-4/ALK wild-type group during statistical analysis.

The presence of activating mutations of the EGFR gene was investigated with the Enterogen EGFR Mutation Analysis Kit in the ABI7500 Real-time PCR device after DNA isolation from the slides of the tumor tissue. The exon 18 codon 719 region, exon 19 deletions, exon 20 insertions and codon 768 and 790 regions, exon 21 codon 858 and 861 regions of the EGFR gene were analyzed. The presence of EML-4/ALK gene translocation was examined immunohistochemically using CST D5F3 antibody. FISH analysis were also performed using the FDA-approved Abbott-Vysis LSI ALK Break Apart Rearrangement probe.

In evaluation of treatment response, evaluation was made according to RECIST v1.1 (Response Evaluation Criteria in Solid Tumors) criteria for computed tomography. PERCIST (Positron Emission Tomography Response Criteria In Solid Tumors) criteria were used for response evaluations with PET-CT. In the light of these criteria, patients who showed complete remission and partial remission after firstline treatment were considered to have responded to treatment. Patients with stable disease and progression were considered unresponsive to treatment.

While evaluating the findings obtained in the study, SPSS 21.0 statistical package program was used for statistical analysis. Pearson Chi-Square test and Fisher Exact test were used to compare qualitative data. The effect of risk factors on survival was analyzed by Kaplan-Meier and Log-rank tests. The results were evaluated at the 95% confidence interval, at the $p < 0.05$ significance level.

This study is approved by Istanbul University Clinical Research Ethics Committee with the decision numbered 83045809/604.01/02.381409 dated 3 December 2015 and conducted in accordance with the Helsinki Principles Declaration.

Results

Among 105 eligible patients, 14 patients harbored EGFR gene mutations while EML-4/ALK transfection was detected in 4 patients. 75 patients were wild-type for both mutations. Median age of entire cohort was 61 while median age of diagnose was 58.5. 81% was male and 78% was smoker. The most frequent metastasis sites were bone (38%), brain (35%) and contralateral lung (32%) (Table 1).

Table 1. Demographic and clinical results (WT/WT: Wild-type for EGFR and EML4/ALK mutations). All p values belong to the comparisons of the cases carrying gene mutations with the WT/WT group.

	Total Cohort (n=105)	EGFR (n=14)		ALK (n=4)		WT/WT (n=75)
			p		p	
Age (y)	61	60	0,79	48	0,026	61
Age at Diagnosis (y)	58,5	58,5	0,93	47	0,028	60
Gender (n)	F	20	6	3	0,020	10
	M	85	8	1	0,020	65
Tobacco Use (n)	82	6	0,002	1	0,037	65
Metastasis Sites (n)						
Brain	37	2	0,06	2	0,40	28
Bone	40	7	0,24	3	0,13	25
Liver	8	2	0,29	1	0,31	5
Pleura	12	3	0,2	0	0,56	9
Contralateral Lung	34	7	0,11	1	0,57	23
Soft Tissue	4	2	0,51	0	0,87	2
Adrenal Gland	19	1	0,44	1	0,56	14

EGFR and EML-4/ALK mutant patients were predominantly female and non-smoker compared to wild-type patients (EGFR; $p=0.025$ and 0.002 , EML-4/ALK; $p=0.003$ and 0.012 , respectively). Additionally, EML-4/ALK mutant patients were

significantly younger than EML-4/ALK wild-type ($p=0.02$) (Table 1). Although metastasis sites did not differ according to EGFR mutation status, when subgroup analysis was performed, exon 19, 20 and 21 mutations significantly appeared to be associated with liver, bone and pleural metastases, respectively ($p=0.046$, 0.05 and 0.035 , respectively).

After firstline platinum-based chemotherapy, complete remission (CR) and partial remission (PR) rates were 4.7% and 24.6%, respectively. Stable disease status was observed in 4.7% while 66% of patients showed progression. Concurrent radiotherapy (RT) and absence of bone metastases at diagnosis were statistically significant factors influencing firstline platinum-based therapy responses. Concurrent radiotherapy with platinum based chemotherapy ($n=35$) showed significantly superior treatment response rates comparing patients who did not receive radiotherapy ($p=0.004$). Relaps rates of patients with and without bone metastasis were %77.5 and %58.5 after platinum based therapy, respectively ($p=0.046$). EGFR or EML-4/ALK gene mutation status did not able to show any significant difference in terms of platinum-based treatment response ($p=0.933$ and 0.184 , respectively). The effect of EGFR exon mutations on treatment response or survival results did not performed due to low numbered subgroups.

Median progression-free survival (PFS) duration of entire cohort was 10 months. Median PFS durations of EGFR, EML-4/ALK and WT/WT groups were also comparable which were 10.8, 12 and 10.2 months, respectively ($p=0.506$) (Figure 1). The observed effect of concurrent radiotherapy and the presence of bone metastases on treatment response is not reflected in the PFS results ($p=0.079$ and 0.285 , respectively). Brain metastasectomy was performed in 8 (21.6%) of 37 cases with brain metastases and no positive effect was detected in terms of PFS and treatment response rates ($p=0.127$ and $p=0.465$). Estimated overall survival (OS) rate of entire cohort was 17% at first year.

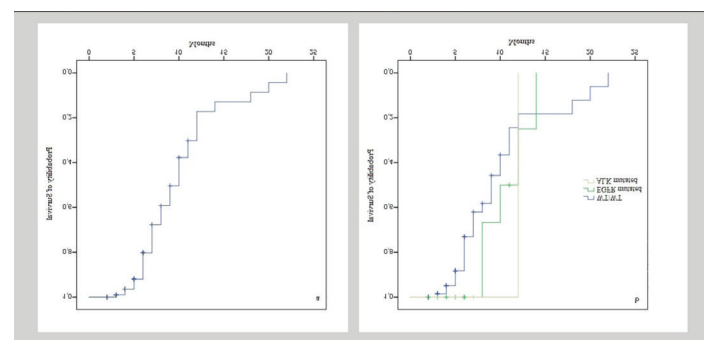


Figure 1. Cumulative PFS of entire cohort (a) and mutation groups (b).



Discussion

Lung cancer is still among the most common cancers but with the decrease in tobacco use and the addition of targeted therapy agents to conventional chemotherapy regimens, a halving of incidence and mortality rates has been observed in the last two decades [12]. As the details of the cancer development process are clarified, new possibilities for treatments emerge. Point mutations in exon 18 and 21, Exon 19 deletion and Exon 20 insertion of the EGFR gene affect the receptor tyrosine kinase domain of the transmembrane cell receptor protein. Similarly, translocation of the ALK gene on the short arm of chromosome 2 with the EML-4 gene causes an increase in intracellular tyrosine kinase activity and this pathway plays a critical role in cancer pathogenesis. Targeted therapies for EGFR and ALK genes, which have been shown to be important in lung non-small cell cancer, are among the most important developments in this regard.

Frequencies of EGFR and EML-4/ALK translocation gene mutations in lung adenocarcinoma were reported as 10-20% and 2-5%, respectively [1]. Patients harbouring these mutations are presenting a different clinical profile compared to wild types. Most

studies showed that, EGFR and EML-4/ALK-mutated lung adenocarcinoma patients are mostly non-smoker females and also EML-4/ALK positive patients are significantly younger [4-6,7]. Results of our cohort were consistent with these findings. In our study, EGFR mutant patients were unable to present significant difference regarding response to platinum-based chemotherapy, PFS or OS comparing EGFR wild-type patients. The only significant factors improving platinum-based treatment results were concomitant RT and absence of bone metastasis at diagnosis ($p=0.004$ and 0.046 , respectively). Also some other studies evaluating platinum-based chemotherapy response from the perspective of EGFR gene mutation also declared ~30% treatment response, but no statistically significant difference was found in treatment response, PFS and OS between mutated and wild-type patients [13-15].

In the study published by Capuzzo et al. [2], 185 cases were examined and EGFR gene mutations were found in 24 cases (15 Exon 19 mutated, 2 Exon 20 mutated and 7 Exon 21 mutated). In patients who received platinum-based chemotherapy as frontline therapy, the response rates were 37% vs. 32.8% in EGFR-mutated and wild-type group, respectively ($p=0.6$). PFS and OS durations of EGFR-mutated and wild-type groups

were also comparable (PFS: 8.1 vs. 4.1 mo., $p=0.1$; OS: 28.5 vs. 14.8 mo., $p=0.07$, respectively). But when EGFR subgroups were examined, it was observed that only patients with Exon 19 mutations responded to platinum-based chemotherapy (46.6%), while no treatment response was obtained in Exon 20 and 21 groups ($p=0.02$) [2].

In another article evaluating 162 lung adenocarcinoma patients [8], of which 40 were EGFR-mutated, platinum-based frontline therapy responses were comparable between EGFR- mutated and wild-type groups (43.5% vs. 23.9%, $p=0.072$). PFS and OS durations of EGFR- mutated and wild-type patients were also did not able to show significant difference between groups ($p=0.69$ and $p=0.069$, respectively). In this study only 9 patients were carrying classical EGFR mutations (exon 18, 19 and 21 mutations) while remaining 31 patients were positive for other EGFR mutations. Subanalysis of classical EGFR-mutated patients showed similar PFS duration ($p=0.81$) but better platinum-based therapy response ($p=0.021$) and improved OS duration ($p=0.028$) comparing wild-type patients. Although the small number of patients, the presence of EGFR gene mutation was declared as an independent favorable prognostic factor for platinum-based treatment response and overall survival [8].

EGFR gene mutation was found to be associated with brain, bone and pleural metastases in the REASON study [5] from Germany, which consisted of 432 EGFR gene mutation-positive cases. We also found a possible correlation between Exon 19, 20 and 21 mutations with liver, bone and pleural metastases, respectively. However, it was impossible to make a definitive interpretation due to the insufficient number of patients.

The presence of EML-4/ALK translocation stands out as a negative factor for patients. In the study of Mayo Clinic [6], 266 EML-4/ALK negative non-small cell lung cancer cases compared to 34 EML-4/ALK-mutated patients and EML-4/ALK positivity declared as a negative predictive factor for progression/relapse-free survival [6]. Koh et al. [9] examined 221 lung adenocarcinoma patients, of which 45 were harboring EML-4/ALK translocation and 46 were EGFR mutants, 170 cases were treated with platinum-based chemotherapy in frontline therapy. The treatment response rates were 18.8% in the EML-4/ALK group, 37.5% in the EGFR group and 40.4% in the EGFR and EML-4/ALK wild-type group ($p=0.091$). Progression-free survival durations were 6.2, 5.4 and 7.3 months, respectively ($p=0.348$). In patients treated with tyrosine kinase inhibitors (TKIs), the treatment response rate reached to 50% and PFS duration improved to 19.6 months in EGFR group. Due to low response rate to

platinum-based chemotherapy, EML-4/ALK transfection gene positivity was declared as a poor prognostic factor [9].

Shaw et al. [7] obtained comparable results in response rates to platinum-based treatment in a cohort consisted of 19 EML-4/ALK positive, 31 EGFR positive and 91 WT/WT patients. But as expected, EGFR-mutated group's frontline TKI response rate was significantly superior than EML-4/ALK and WT/WT groups ($p < 0.001$). PFS duration of EGFR group after TKI treatment was 16 months which was significantly longer than EML-4/ALK-mutated (5 months) and WT/WT patients (6 months) ($p = 0.004$). The presence of EML-4/ALK transfection gene was strongly related to TKI resistance [7].

Current guidelines recommend TKIs and ALK inhibitors as first-line therapy for patients with these mutations [1,3]. Although additional 20% increase in firstline therapy responses and approximately 12 months of PFS with targeted therapies for EGFR and ALK mutations, this improvement was not reflected in overall survival results.

First generation TKIs, gefitinib and erlotinib, were unable to show significant OS benefit with over platinum-based regimens but OR and PFS rates were significantly improved [1]. In the light of these results, and with their safe use even in patients with low performance scores, TKIs have become the backbone of EGFR-mutated lung adenocarcinoma patients. Following these agents, afatinib and dacomitinib emerged to the market as second generation TKIs. Osimertinib, third-generation TKI, proved itself in the FLAURA study [11] with its PFS contribution against 1st generation TKIs and its superior response results in cases with CNS metastasis. Moreover, with the statistically significant contribution to OS (38.6 vs 31.8 months, $p = 0.046$), the current guidelines are recommending osimertinib as preferred therapy in firstline treatment of patients with sensitizing EGFR mutations [1,3]. Crizotinib was the first targeted therapy improving OR and PFS rates of EML-4/ALK rearranged patients comparing platinum-based therapies. Ensuing ALK inhibitors, ceritinib, alectinib and brigatinib took these results to an even better point. In the ALTA-1L study [10], the estimated 1-year PFS rate for brigatinib was a remarkable success compared to crizotinib (67% vs 43%, $p < 0.001$). Ensartinib and lorlatinib also stand out as ALK inhibitors whose phase 3 studies are ongoing and are expected to be included in daily clinical practice in the near future [1].

Although targeted therapies have considerably increased the OR and PFS results compared to platinum-based therapies, there is still a need for new agents to prolong the duration of OS.

Conclusion

The presence of EGFR and ALK mutations does not effect the treatment response of platinum-based regimens according to current literature and our study. Retrospective study design and low numbered subgroups in EGFR and EML-4/ALK mutated patients are the limitations of our study. Although these limitations, the association of EGFR exon subsets with metastasis points is worth investigating, as demonstrated in REASON study [5].

Disclosure

The study was not supported by any individual, institution or organization. It does not contain any conflict of interest.

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