Prognostic impact of initial maximum standardized uptake value of $^{18}$F-FDG PET/CT on treatment response in patients with metastatic lung adenocarcinoma treated with erlotinib

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Purpose: To investigate whether the initial maximum standardized uptake value (SUV\textsubscript{max}) on fluorine-18 fluorodeoxyglucose positron emission tomography/computed tomography ($^{18}$F-FDG PET/CT) has a prognostic significance in metastatic lung adenocarcinoma.

Patients and methods: Sixty patients (24 females, mean age: 57.9±12 years) with metastatic stage lung adenocarcinoma who used erlotinib and underwent $^{18}$F-FDG PET/CT at the time of diagnosis between May 2010 and May 2014 were enrolled in this retrospective study. The patients were stratified according to the median SUV\textsubscript{max} value, which was found as 11. Progression-free survival (PFS) rates for 3, 6, and 12 months were examined for SUV\textsubscript{max} values and epidermal growth factor receptor (EGFR) mutation status.

Results: The number of EGFR-sensitizing mutation positive/negative/unknown was 26/17/17, respectively, and the number of patients using erlotinib at first-line, second-line, and third-line therapy was 15, 31, and 14 consecutively. The PFS rates of EGFR mutation positive, negative, and unknown patients for 3 months were 73.1%, 35.3%, and 41.2% ($P=0.026$, odds ratio [OR]=4.39; 95% confidence interval [CI]: 1.45–13.26), respectively. The PFS rates of EGFR positive, negative, and unknown patients for 6 months were 50%, 29.4%, and 29.4% ($P=0.267$, OR: 2.4; 95% CI: 0.82–6.96), respectively. The PFS rates of EGFR positive, negative, and unknown patients for 12 months were 42.3%, 29.4%, 23.5% ($P=0.408$, OR: 2.0; 95% CI: 0.42–5.26), respectively. Thirty-one of 60 patients had SUV\textsubscript{max} values ≤11. The PFS rates for 3, 6, and 12 months were 70.5%/28% ($P=0.001$, OR=9.0; 95% CI: 2.79–29.04), 61.7%/8% (P<0.001, OR=28.35; 95% CI: 5.5–143), and 52.9%/8% ($P<0.001$, OR=18.69; 95% CI: 3.76–92.9) for low SUV\textsubscript{max} (≤11) group/high SUV\textsubscript{max} (>11) group, respectively.

Conclusion: Initial SUV\textsubscript{max} value on $^{18}$F-FDG PET/CT is found to be a prognostic factor anticipating the response to erlotinib for 3, 6, and 12-month rates of PFS in both EGFR-sensitizing mutation and wild-type tumor group.

Keywords: lung adenocarcinoma, erlotinib, $^{18}$F-fluorodeoxyglucose positron emission tomography/computed tomography, maximum standardized uptake value, treatment response, prognosis

Introduction

Predictive factors may help choose the priority for the treatment and, eventually, more and more to develop individualized therapeutic strategies in patients with cancer. In advanced non-small cell lung cancer (NSCLC), several pretherapeutic prognostic factors have been identified, such as performance status (PS), age, sex, weight loss, neutrophil counts, C-reactive protein, albumin, serum calcium level.\textsuperscript{1–3}
Moreover, genetic mutations having predictive and prognostic significance led to changes in the treatment algorithm of advanced NSCLC in consequence of developed targeted therapies. Sensitization mutation within the kinase domain of the epidermal growth factor receptor (EGFR) predicts greatly improved the clinical outcome in NSCLC treated with the EGFR-tyrosine kinase inhibitors (TKIs) including erlotinib, gefitinib, and afatinib.6,7 The incidence of EGFR-sensitizing mutations is approximately 16.6% and the mostly detected and prognostic significant types of EGFR kinase mutations are exon-19 deletions and L858R mutation in exon 21.6,7 EGFR-sensitizing mutations of exons 19 and 21 are more frequent in females, non-smokers, patients with adenocarcinoma, and Asian origin patients. Those features have been associated with positive responses to TKIs.8,9

Positron emission tomography/computed tomography with fluorine-18 fluorodeoxyglucose ([18F]-FDG PET/CT) is used to visualize the metabolic activity of tumors. In early stage NSCLC, the maximum standardized uptake value (SUV\textsubscript{max}) of FDG on PET was found to be the strongest prognostic factor among the patients treated with curative surgery or radiotherapy.10 However, there was no evidence proving prognostic value of baseline FDG PET uptake in patients with advanced NSCLC who were treated with standard chemotherapy.11 In this regard, we aimed to assess the prognostic significance of SUV\textsubscript{max} value of [18F]-FDG PET/CT in patients with metastatic lung adenocarcinoma treated with erlotinib for the first-line therapy, after chemotherapy failure, or maintenance therapy.

Patients and methods

Study design

This retrospective-observational study was conducted in single center Gaziantep University Hospital in Turkey. The study and informed consent documentation were reviewed and approved by the Independent Ethics Committee of Gaziantep University, and the participants all provided informed, written consent. This study was conducted in compliance with the ethical principles according to the Declaration of Helsinki. This trial was designed to assess whether [18F]-FDG PET pretreatment SUV\textsubscript{max} is prognostic for the response to erlotinib treatment.

Patient selection

Between May 2010 and May 2014, 60 patients with histologically confirmed metastatic lung adenocarcinoma who underwent [18F]-FDG PET/CT prior to systemic therapy were analyzed. The following parameters were assessed prior to the beginning of therapy: age, sex, Eastern Cooperative Oncology Group PS (0–2), EGFR mutation status, smoking habit, treatment line, and the presence of cranial metastasis. The patients over 18 years old with an Eastern Cooperative Oncology Group PS 0–2 and normal blood glucose level (150 mg/dL) were allowed. Patients with brain and/or bone metastasis requiring further local treatment prior to [18F]-FDG PET/CT were included in the study. Highest SUV\textsubscript{max} in the primary tumor or metastatic lesion on [18F]-FDG PET/CT was recorded. In patients using erlotinib at second line or more, [18F]-FDG PET/CT SUV\textsubscript{max} values after the detection of progression were noted. For the patients on maintenance treatment, initial [18F]-FDG PET/CT SUV\textsubscript{max} values were noted. The study population was divided into two groups according to the median SUV\textsubscript{max} value as 11. Three, 6, and 12-month progression-free survival (PFS) rates were investigated according to both SUV\textsubscript{max} values and EGFR mutation status.

[18F]-FDG PET/CT imaging

All patients fasted for at least 6 hours before FDG injection and their blood glucose levels were less than 150 mg/dL before radiotracer injection. Eight to 15 mCi (296–555 MBq) [18F]-FDG was administered intravenously. All patients received an oral contrast (a solution containing sodium amidotrizoate and meglumine amidotrizoate) before PET/CT imaging. Whole-body [18F]-FDG PET/CT images were acquired from vertex to proximal thigh 60 minutes after FDG injection, using a Biograph Duo LSO PET/CT system (Siemens Medical Solutions, Hoffman Estates, IL, USA). After the CT topogram, a spiral CT scan and subsequent PET scan with an acquisition time of 2–3 minutes for each bed position according to patient weight were performed. After CT-based attenuation correction, PET images were reconstructed with an ordered subset expectation maximization iterative reconstruction algorithm. A circular region of interest was drawn manually on hypermetabolic lesions on axial fused PET/CT images. The SUV\textsubscript{max} value, a semiquantitative index of FDG uptake in tissue, was calculated as the maximum measured activity concentration divided by injected activity divided by body weight of the subject as in the following formula: SUV\textsubscript{max} = decay corrected selected region activity (mCi/mL)/(injected dose [mCi]/body weight [kg]).

Molecular analysis

DNA was extracted from 1 cm\textsuperscript{2} of 10 mm thick paraffin-embedded tissue containing at least 20% tumor tissue. Genomic DNA was extracted using a QIAamp DNA formalin-fixed, paraffin-embedded tissue kit (Qiagen NV,
Venlo, the Netherlands) according to the manufacturer’s instructions with overnight proteinase K digestion and eluting in 50 µL of water. Pyrosequencing for pyrosequencing analysis, the protocol was followed according to the manufacturer’s instructions (EGFR Pyro Assay; Qiagen). The pyrosequencing results were analyzed using the PyroMark Q24 version 2.0.6 software (Qiagen), which identifies the presence of a specific mutation and its percentage. Manufacturer-supplied logarithm of the odds (LOD) thresholds were used to call a mutation for LOD studies (≥% LOD is positive).

Treatment

All patients were allowed to receive erlotinib 150 mg/day for at least 6 weeks. Erlotinib treatment was continued until disease progression or toxicity. It was either given as a first-line treatment or as a second-line treatment after platinum-doublet failure or as a maintenance treatment.

Response evaluation

The response was assessed using the response evaluation criteria in solid tumors version 1.1. The first CT scan and 18F-FDG PET/CT were performed prior to treatment and follow-up CT scans or 18F-FDG PET/CT were performed every 12 weeks or in case of clinically suspected progression.

Statistical analysis

First, univariate analyses were performed to compare baseline characteristics. To compare the two groups, chi-squared tests (for categorical variables) were used. The Mann–Whitney U test was used to show the difference between independent groups according to non-normally distributed numerical variables. Multivariate binary logistic regression analyses were performed to determine odds ratio (OR) and 95% confidence intervals (CIs). All univariate analyses were performed in SPSS for Windows version 22.0 (SPSS Inc., Chicago, IL, USA).

Result

Patients

Sixty patients who underwent 18F-FDG PET scans prior to erlotinib treatment were enrolled in this study. The mean age was 57.9 years (SD 12; range: 36–84 years) and 24 (40%) were women. Forty-three (71.6%) patients had tumor tissue available for EGFR mutation analysis. Twenty-six patients (12.5%) had EGFR-sensitizing mutations (mostly; deletions in exon 19, L858R mutation in exon 21), 17 patients had wild-type tumor, and 17 patients had unknown EGFR status. In this population of 60 patients, SUVmax ranged between 3.8 and 24.6, with a mean of 11.5 (SD 4.4) and a median of 11. SUVmax of 31 patients were ≤11 and 29 patients were >11. Demographic features of the patients were recorded (Table 1).

Association of initial SUVmax and PFS according to EGFR mutation analysis

The 3-month PFS rates were 73.1% in the EGFR-sensitizing mutation group, 35.3% in the EGFR negative group, and 41.2% in the rEGFR unknown group (P=0.026); the 6-month PFS rates were 50%, 29%, and 29.4% (P=0.267), respectively; and the 12-month PFS rates were 42.3%, 29.4%, 23.5%, respectively (P=0.408). Using Mann–Whitney U test, OR for 3, 6, and 12-month PFS rates based on EGFR positivity were 4.39 (95% CI: 1.45–13.26, P=0.017), 2.4 (95% CI: 0.82–6.96, P=0.017), and 2.0 (95% CI: 0.42–5.26, P=0.20), respectively (Table 2). Thirty-one of the patients had SUVmax values ≤11 and 29 patients had SUVmax values >11. The 3-month PFS rate was 77.4% in patients who have SUVmax ≤11 and was 27.6 in patients who have SUVmax >11 (OR=9.0; 95% CI: 2.79–29.04, P<0.001). The 6-month PFS rates were 67.7% vs 6.9% (OR=28.35; 95% CI: 5.5–143, P<0.001) and the 12-month PFS rates were 58.1% vs 6.9% (OR=18.69; 95% CI: 3.76–92.9, P<0.001) (Table 2 and Figure 1).

### Table 1 Demographic characteristics of patients

<table>
<thead>
<tr>
<th>Characteristics</th>
<th>SUVmax &lt; 11, n (%)</th>
<th>SUVmax &gt; 11, n (%)</th>
<th>P-value</th>
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<tbody>
<tr>
<td>Age (years)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>≤65</td>
<td>22 (71)</td>
<td>23 (79.3)</td>
<td>0.456</td>
</tr>
<tr>
<td>&gt;65</td>
<td>9 (29)</td>
<td>6 (20.7)</td>
<td></td>
</tr>
<tr>
<td>Sex</td>
<td></td>
<td></td>
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<tr>
<td>Female</td>
<td>15 (48.4)</td>
<td>9 (31)</td>
<td>0.170</td>
</tr>
<tr>
<td>Male</td>
<td>16 (51.6)</td>
<td>20 (69)</td>
<td></td>
</tr>
<tr>
<td>Smoking</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Smoking</td>
<td>15 (48.4)</td>
<td>14 (48.3)</td>
<td>0.993</td>
</tr>
<tr>
<td>No smoking or light smoking</td>
<td>16 (51.6)</td>
<td>15 (51.7)</td>
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<tr>
<td>Therapy line</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>First</td>
<td>5 (16.1)</td>
<td>11 (37.9)</td>
<td>0.153</td>
</tr>
<tr>
<td>Second</td>
<td>16 (51.6)</td>
<td>12 (41.4)</td>
<td></td>
</tr>
<tr>
<td>Third or higher</td>
<td>10 (32.3)</td>
<td>6 (20.7)</td>
<td></td>
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<tr>
<td>EGFR status</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Mutant</td>
<td>11 (35.5)</td>
<td>15 (51.7)</td>
<td>0.447</td>
</tr>
<tr>
<td>Wild-type</td>
<td>10 (32.3)</td>
<td>7 (24.1)</td>
<td></td>
</tr>
<tr>
<td>Insufficient/not examined</td>
<td>10 (32.3)</td>
<td>7 (24.1)</td>
<td></td>
</tr>
<tr>
<td>Use indication</td>
<td></td>
<td></td>
<td></td>
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<tr>
<td>EGFR-mutant first line</td>
<td>5 (16.1)</td>
<td>11 (37.9)</td>
<td>0.073</td>
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<tr>
<td>After progression</td>
<td>19 (61.3)</td>
<td>16 (55.2)</td>
<td></td>
</tr>
<tr>
<td>Maintenance</td>
<td>7 (22.6)</td>
<td>2 (22.2)</td>
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<tr>
<td>Cranial metastasis</td>
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</tr>
<tr>
<td>Present</td>
<td>9 (29)</td>
<td>6 (20.7)</td>
<td>0.456</td>
</tr>
<tr>
<td>Absent</td>
<td>22 (71)</td>
<td>23 (79.3)</td>
<td></td>
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</tbody>
</table>

Abbreviations: EGFR, epidermal growth factor receptor; SUVmax, maximum standardized uptake value.
Table 2 Odds ratio (OR) and P-values using Mann–Whitney U test and multivariate binary logistic regression

<table>
<thead>
<tr>
<th>Survival time, n (%)</th>
<th>OR (95% CI)</th>
<th>P-value</th>
<th>Non-smoker</th>
<th>OR (95% CI)</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>≥11</td>
<td>24 (77.4)</td>
<td>7 (22.6)</td>
<td>9.00 (2.79–29.04)</td>
<td>0.001</td>
<td>18.3 (3.66–91.3)</td>
</tr>
<tr>
<td>&gt;11</td>
<td>8 (25.0)</td>
<td>21 (75.0)</td>
<td>1 (Reference)</td>
<td>1 (Reference)</td>
<td></td>
</tr>
<tr>
<td>Mutation</td>
<td>Positive</td>
<td>19 (73.1)</td>
<td>7 (26.9)</td>
<td>4.39 (1.45–13.29)</td>
<td>0.009</td>
</tr>
<tr>
<td>Negative + unknown</td>
<td>13 (38.2)</td>
<td>21 (61.8)</td>
<td>1 (Reference)</td>
<td>1 (Reference)</td>
<td></td>
</tr>
<tr>
<td>≥6 months</td>
<td>21 (67.7)</td>
<td>10 (32.3)</td>
<td>28.3 (5.50–143)</td>
<td>&lt;0.001</td>
<td>57.7 (7.98–417)</td>
</tr>
<tr>
<td>&gt;12 months</td>
<td>2 (6.6)</td>
<td>27 (93.1)</td>
<td>1 (Reference)</td>
<td>1 (Reference)</td>
<td></td>
</tr>
<tr>
<td>≥11</td>
<td>18 (58.1)</td>
<td>13 (41.9)</td>
<td>18.7 (3.76–92.9)</td>
<td>0.001</td>
<td>22.2 (4.17–118)</td>
</tr>
<tr>
<td>&gt;11</td>
<td>2 (6.9)</td>
<td>27 (93.1)</td>
<td>1 (Reference)</td>
<td>1 (Reference)</td>
<td></td>
</tr>
<tr>
<td>Mutation</td>
<td>Positive</td>
<td>11 (42.3)</td>
<td>15 (57.7)</td>
<td>2.03 (0.68–6.05)</td>
<td>0.20</td>
</tr>
<tr>
<td>Negative + unknown</td>
<td>9 (26.5)</td>
<td>25 (73.5)</td>
<td>1 (Reference)</td>
<td>1 (Reference)</td>
<td></td>
</tr>
</tbody>
</table>

Abbreviations: CI, confidence interval; OR, odds ratio; SUV_{max}, maximum standardized uptake value.

Subgroup analysis

In EGFR-sensitizing mutation group, eleven of the patients had SUV_{max} values ≤11 and 15 of the patients had SUV_{max} values >11. Accordingly, the 3-month PFS rates were 100% vs 53.3% (P = 0.08), 6-month PFS rates were 100% vs 13.3% (P < 0.001), and 12-month PFS rates were 81.3% vs 13.3% (P < 0.001), favoring the group having SUV_{max} ≤11.

In the EGFR negative group, ten of the patients had SUV_{max} values ≤11 and seven of the patients had SUV_{max} values >11. Accordingly, the 3-month PFS rates were 60% vs 0% (P = 0.01), 6-month PFS rates were 50% vs 0% (P = 0.026), and 12-month PFS rates were 50% vs 0% (P = 0.026), favoring the group having SUV_{max} ≤11.

In the EGFR unknown group, ten of the patients had SUV_{max} values ≤11 and seven of the patients had SUV_{max} values >11. Accordingly, 3-month PFS rates were 70% vs 0% (P = 0.004), 6-month PFS rates were 50% vs 0% (P = 0.026), and 12-month PFS rates were 40% vs 0% (P = 0.056), favoring the group having SUV_{max} ≤11.

Subgroup analysis according to smoking habit

Eleven of the patients were smokers and 15 of the patients were non-smokers in patients who had SUV_{max} ≤11. Using multivariate binary logistic regression analyses, OR for 3, 6, and 12-month PFS rates were 18.29 (95% CI: 3.66–91.3, P = 0.001), 57.7 (95% CI: 7.98–417, P < 0.001), and 22.2 (95% CI: 4.17–118, P < 0.001), respectively, in patients who had SUV_{max} ≤11 and were non-smokers (Table 2). Fourteen of the patients were smokers and 15 of the patients were non-smokers in patients who had SUV_{max} >11. The 3-month PFS rates of these patient groups were 0% and 53.3% (P = 0.002), 6-month PFS rates were 0% and 13.3% (P = 0.483), and 12-month PFS rates were 0% and 13.3% (P = 0.483).

According to EGFR mutation status, there were five smokers and 21 non-smokers in the EGFR-sensitizing group. The 3-month PFS rates were 20% and 85.7% (P = 0.03), 6-month PFS rates were 20% and 57.1% (P = 0.322), and 12-month PFS rates were 20% and 47.6% (P = 0.356) in the smoker and non-smoker groups, respectively. There were 13 smokers and four non-smokers in the EGFR negative group. The 3-month PFS rates were 30% and 50% (P = 0.482), 6-month PFS rates were 23.1% and 50% (P = 0.301), and 12-month PFS rates were 23.1% and 50% (P = 0.301) in the smoker and non-smoker groups. There were eleven smokers and six non-smokers in EGFR unknown group. Three-month PFS rates were 45.5% and 33.3% (P = 0.627), 6-month PFS rates were 27.3% and 33.3% (P = 0.793), and 12-month PFS rates were 37.3% and 16.7% (P = 0.622) in the smoker and non-smoker groups, respectively. Using multivariate binary logistic regression analyses based on EGFR-sensitizing mutation and non-smoking, OR was 2.64 (95% CI: 0.75–921, P = 0.127).
for 3 months PFS rate, OR was 1.46 (95% CI: 0.42–5.03, $P=0.54$) for 6 months PFS rate, and OR was 1.5 (95% CI: 0.42–5.26, $P=0.52$) for 12 months PFS rate (Table 2).

**Discussion**

New diagnostic and treatment strategies provide more successful survival rates in patients with lung cancer. For the first-line treatment of NSCLC, use of platinum-doublet chemotherapy for patients with good PS has been widely accepted. According to one meta-analysis, the 1-year survival for the platinum-containing regimens was 34% (95% CI: 33%–36%) and 29% (95% CI: 27%–30%) for the nonplatinum therapies.12 The standard first-line treatment for patients with EGFR-sensitizing mutations is EGFR TKIs.13 Mok et al showed that the 12-month rates of PFS were 24.9% with gefitinib and 6.7% with carboplatin–paclitaxel in patients having EGFR mutation in the first-line treatment of advanced NSCLC patients selected on the basis of clinical characteristics that included a history of no smoking or light smoking as well as histologic evidence of adenocarcinoma. The gefitinib group had a significantly longer median PFS (10.8 vs 5.4 months in the chemotherapy group; hazard ratio [HR] 0.30; 95% CI: 0.22–0.41; $P<0.001$) and the median overall survival was 30.5 vs 23.6 months, respectively ($P=0.31$).14 And also it was shown that afatinib is associated with prolongation of PFS when compared with standard doublet chemotherapy in patients with advanced lung adenocarcinoma with EGFR mutations (median PFS was 11.1 months for afatinib and 6.9 months for chemotherapy (HR 0.58; 95% CI: 0.43–0.78; $P=0.001$).15 Similar finding was achieved with erlotinib in ENSURE study.16 Although EGFR-sentitizing mutations are the most important predictive markers of clinical outcome for EGFR-TKIs treatment in advanced NSCLC, according to DELTA trial there was not statistically differences after the first-line treatment. DELTA trial which assessed the efficacy of erlotinib versus docetaxel in second-line and third-line therapy in 255 patients with EGFR-mutant and wild-type tumors. A total of 51 EGFR-mutant patients assessed and

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**Figure 1** The (A) 3, (B) 6, and (C) 12-month rates of progression-free survival according to $\text{SUV}_{\text{max}}$ values.

**Note:** $\text{SUV}_{\text{max}} = \text{decay corrected selected region activity (mCi/mL)/injected dose (mCi)/body weight (kg)}$.

**Abbreviations:** $\text{SUV}_{\text{max}}$, maximum standardized uptake value; PFS, progression-free survival.
PFS was 9.3 vs 7 months in erlotinib vs docetaxel group (HR 0.96; 95% CI: 0.51–1.79; \( P = 0.91 \)) and it was 1.3 vs 2.9 months in patients with EGFR wild-type tumor (HR 1.45; 95% CI: 1.09–1.94; \( P = 0.01 \)).

Thus, the presence of EGFR-sensitizing mutation was insufficient to predict the outcome of treatment after the first-line therapy.

Despite the EGFR-sentizing mutations, drug resistance may unavoidably appear, and the disease eventually progresses. The mechanisms of secondary resistance to EGFR TKIs include the acquisition of the T790M gatekeeper mutation, mesenchymal–epithelial transition amplification, and transformation into small-cell lung cancer. And also, approximately 30% of the patients with EGFR-sensitizing mutations do not exhibit objective responses to EGFR TKIs within 3 months. Although possible mechanisms have been investigated in several preclinical and retrospective studies, the mechanism of primary resistance to EGFR TKIs in EGFR-mutant NSCLC has not been clearly understood.

V-Ki-ras2 Kirsten rat sarcoma viral oncogene homolog (KRAS) mutations, mesenchymal-epithelial transition amplification, and phosphatase and tensin homolog loss could be related to the response to TKIs in these groups.

Erlotinib is registered for the treatment of all patients with advanced NSCLC and prolongs survival after first-line, second-line chemotherapy, and maintenance therapy in patients having an EGFR mutation and also wild-type tumors. The survival benefit of erlotinib is unlikely to be solely due to the EGFR mutations and other molecular mechanisms probably also contribute to the effect. The SATURN trial showed that patients with wild-type tumors who had been treated with first-line platinum-doublet chemotherapy also benefit from maintenance treatment with erlotinib and have prolonged PFS when compared with placebo in patients with stable disease and complete response/partial response. The HR for erlotinib benefit was 0.68 in the stable disease group (95% CI: 0.56–0.83; \( P < 0.0001 \)) with a median PFS of 12.1 vs 11.3 weeks (2.8 vs 2.6 months), respectively. The HR in the complete response/partial response group was 0.74 (0.60–0.92; \( P = 0.0059 \)) with a median PFS of 12.4 vs 11.1 weeks (2.9 vs 2.6 months), respectively.

One placebo-controlled phase III study investigated the effect of gefitinib on survival as second-line or third-line treatment for patients with advanced NSCLC with unknown EGFR status. Subgroup analyses showed significantly longer survival in the gefitinib group than the placebo group for never-smokers (\( P = 0.012 \); median survival 8.9 vs 6.1 months) and patients of Asian origin (\( P = 0.01 \); median survival 9.5 vs 5.5 months).

TKIs, docetaxel, and pemetrexed are approved for the second-line treatment of NSCLC. TITAN trials were conducted to show that there were no significant differences in efficacy between patients treated with erlotinib and those treated with docetaxel or pemetrexed in second-line therapy. Median PFS in the erlotinib group was 6.3 weeks (95% CI: 6.1–6.9) vs 8.6 weeks (7.1–12.1) in the chemotherapy group. There was no statistically significant difference in PFS between the two treatment groups (HR: 1.19, 95% CI: 0.97–1.46; \( P = 0.089 \)).

In this study, the detected rate of EGFR-sensitizing mutation was only 3% and 2% in erlotinib group and chemotherapy group, respectively, and smoking status was not a predictive factor.

Hellenic Oncology Research Group study assesses the efficacy of pemetrexed versus erlotinib in pretreated patients with advanced NSCLC. The median PFS was 2.9 months (range: 0.4–27.3 months) and 3.6 months (range: 0.2–47.8 months) for the pemetrexed and erlotinib arms, respectively (\( P = 0.136 \)). No difference was observed in treatment line or smoking status between the treatment arms. However, the presence of KRAS mutation was reported to be a possible negative predictive factor for response to EGFR TKIs.

In contrast, TAILOR study, which compared erlotinib and docetaxel in the treatment of NSCLC showed that KRAS mutational status and smoking status were not predictive for response.

In this context, new instruments are needed to predict the response to TKIs, which have a wide indication of use in first-line and sequential therapies. This study specifically focused on the role of pretreatment \(^{18}\)F-FDG PET in providing prognostic information in response to erlotinib for patients with metastatic-stage lung adenocarcinoma at first-line and after first-line therapy. Nevertheless, EGFR-sensitizing mutation is an absolute predictor for TKIs selection at first-line therapy, TKIs were not statistically superior to chemotherapy after first-line treatment. Studies investigating the predictive value of smoking habit and K-RAS mutation showed controversial results as mentioned earlier. Along with various studies done on many genetic mutations, these tests are both costly and also cannot explain the majority of cases who have primary and secondary resistance in EGFR-sensitizing patient and patients with wild-type tumors.

Scheffler et al showed that the identification of the lesion with the highest metabolic activity in FDG PET has significant prognostic relevance before the initiation of erlotinib therapy independent to EGFR mutation status in patients with advanced NSCLC. Accordingly, patients with an \(^{18}\)F-FDG SUV\(_{\text{max}}\) value less than 6.6 had a significantly better overall survival (16.3 months, 95% CI 7.1–25.4 months) compared to patients with an \(^{18}\)F-FDG SUV\(_{\text{max}}\) value more than 6.6 (3.1 months, 95%
Metastatic lung adenocarcinoma treated with erlotinib

Conclusion

Despite the predictive value of the EGFR-sensitizing mutation, erlotinib performed a low efficiency in patients who have SUVmax values \( > 11 \). Additionally, erlotinib had no efficiency in patients who have SUVmax values \( > 11 \) in wild-type and EGFR-unknown groups. This study indicated that initial 18F-FDG PET/CT SUVmax value is a prognostic instrument for lung cancer patients who treated with erlotinib, which is an approved drug in the sequential treatment of lung adenocarcinoma independent to EGFR mutation status.

Disclosure

The authors report no conflicts of interest in this work.

References


