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ORIGINAL RESEARCH

Epidermal growth factor receptor exon 20 mutation in lung cancer: types, incidence, clinical features and impact on treatment

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Background: There are limited data available on the treatment and outcome of epidermal growth factor receptor (EGFR) exon 20-mutated lung cancer patients. Hence, we planned an analysis of the demographic details, clinical profile and survival of lung cancer patients with exon 20 mutations. We compared our results to patients with EGFR tyrosine kinase inhibitor (TKI)-sensitizing activating and EGFR/anaplastic lymphoma kinase (ALK)-negative mutations.

Methods: This was a retrospective analysis of lung cancer patients who were treated at our center between January 2010 and August 2014. We reviewed the results of EGFR mutation testing by real-time polymerase chain reaction and Sanger sequencing. We also reviewed the data relating to baseline demographics, clinical profile, patient treatment and outcome measures in terms of response and overall survival (OS).

Results: A total of 580 patients fulfilled the selection criteria. In all, 227 (39.1%) patients had EGFR TKI-sensitizing activating mutations, 20 (3.4%) patients had exon 20 insertion mutations and 333 patients were EGFR/ALK mutation negative (57.5%). The median OS was 5 months (95% confidence interval [CI] 0.17–9.8 months) in exon 20 insertion mutations, 16.1 months (95% CI 12.8–19.5 months) in EGFR TKI-sensitizing activating mutations and 10 months (95% CI 7.9–12.1 months) in EGFR/ALK mutation-negative patients. The median OS was significantly better for the EGFR TKI-sensitizing activating mutation group (P=0.000, log-rank test) and for the EGFR/ALK-negative group (P=0.037, log-rank test) compared to the exon 20-mutated group.

Conclusion: Exon 20 mutation results in a poorer OS prognosis compared to EGFR- and ALK-negative patients and patients harboring EGFR TKI-sensitizing activating mutations. The incidence of de novo exon 20 insertions was 3.4%. Different types of exon mutations seem to have different outcomes.

Keywords: exon 20, lung cancer, EGFR mutation, TKI resistance, insertions

Introduction

Somatic mutations in epidermal growth factor receptor (EGFR) were identified in 2004, and this discovery has revolutionized the management of lung cancer. Different types of EGFR mutations have been identified; in-frame deletions in exon 19 and exon 18 mutations and point mutations in exon 21 are the most common.¹ These mutations account for 90% of EGFR mutations and are also termed as EGFR tyrosine kinase inhibitor (TKI)-sensitizing activating mutations. These mutations predict a favorable response to oral TKIs; the response rate to first-generation TKIs (erlotinib or gefitinib) ranges between 60% and 70%.²⁻⁴ In addition to these mutations, a rare

OncoTargets and Therapy 2017:10 2903-2908

© 0 0 2017 Noronha et al. This work is published and licensed by Dove Medical Press Limited. The full terms of this license are available at https://www.dovepress.com/terms.php and incorporate the Creative Commons Attribution – Non Commercial (unported, v3.0) License (http://creativecommons.org/license/by-nc/3.0/). By accessing the work you hereby accept the Terms. Non-commercial uses of the work are permitted without any further permission from Dove Medical Press Limited, provided the work is properly attributed. For permission for commercial use of this work, please see paragraphs 4.2 and 5 of our Terms (https://www.dovepress.com/terms.php). subset of mutations has been identified, which are associated with de novo resistance to first-generation and second-generation TKIs.⁵

Exon 20-mutated lung cancer is a rare subset of lung cancer (4%).⁵⁻⁷ There are limited data available on the treatment and outcome of these patients. Hence, we planned an analysis of the demographic details, clinical profile and survival of lung cancer patients with exon 20 mutations and then compared these results to EGFR TKI-sensitizing activating mutations and EGFR/anaplastic lymphoma kinase (ALK) mutation-negative patients.

Methods Ethics

This was a retrospective analysis of lung cancer patients who were treated at our center between January 2010 and August 2014. The institutional review board (IRB) and the ethics committee (EC) of Tata Memorial Center (TMC) – Advanced Centre for Treatment, Research and Education in Cancer (ACTREC; Mumbai, India) approved the project of lung cancer audit (No 108) during the 21st TMC-ACTREC IRB meeting. Since this was a retrospective analysis, the IRB and the EC waived the need for an informed consent. Patient records/information were anonymized and de-identified prior to analysis.

Selection criteria

From the lung cancer medical oncology database, patients were selected based on the following criteria:

- 1. Biopsy-positive adenocarcinoma lung
- 2. EGFR mutation positive or EGFR/ALK mutation negative
- 3. Underwent treatment at our center

The demographic factors, clinical profile and overall survival (OS) period were noted. The OS was calculated from the date of start of the first line of treatment till death. Patients who were alive till March 31, 2015, were censored at their last known follow-ups.

The testing for exon mutations was done by real-time polymerase chain reaction (PCR) and Sanger sequencing, the details of which have been published earlier.¹

Statistical analysis

SPSS version 16 was used for the analysis. The continuous variables were compared by using Student's *t*-test and categorical variables by chi-square test or Fisher's test. The Kaplan–Meier method was used for the estimation of OS. The log-rank test was used for univariate analysis. Cox-proportional hazard model was used for multivariate analysis. The factors considered for multivariate analysis were the presence of exon 20 mutations, EGFR TKI-sensitizing activating mutations, extrathoracic disease, brain metastasis, liver metastases and Eastern Cooperative Oncology Group (ECOG) performance status (PS).

Results Demographic profile

A total of 580 patients were selected for this study. There were 227 (39.1%) patients in the EGFR TKI-sensitizing activating mutation cohort, 20 patients in the exon 20 insertion cohort (3.4%) and 333 patients in the EGFR/ALK mutation-negative cohort (57.5%). The median age of the entire cohort was 56 years (interquartile range [IQR] 49-62 years). The gender distribution included 375 males (64.7%) and 205 females (35.3%). The EGFR TKI-sensitizing activating mutations involved exon 19, exon 21 and exon 18 in 143 (24.7%), 72 (12.4%) and 12 (2.1%) patients, respectively. Exon 20 mutation was seen in 20 patients (3.5%; 95% CI 2.2%–5.7%). The comparison of the demographic profile between the three subgroups is shown in Table 1. There was no difference in age or gender distribution between the two groups. However, there was a significantly higher proportion of nonsmokers in the EGFR TKI-sensitizing activating mutation group (74.0% vs 52.0%, P=0.000, chi-square) compared to the EGFR/ALK mutation-negative group. There was no statistical difference in the proportion of nonsmoker patients between exon 20 and EGFR and ALK mutation-negative groups (P=0.189).

Clinical profile

All patients had Stage IV disease with extrathoracic disease present in 222 patients (38.3%). Metastases to the brain, bone and liver were seen in 45 (7.8%), 148 (25.5%) and 70 (12.1%) patients, respectively. The ECOG PS was 0–1 in 382 patients (65.9%) and ≥ 2 in 198 (34.1%) patients.

There were a significantly higher proportion of patients with poor ECOG PS in the EGFR TKI-sensitizing activating mutation group compared to EGFR/ALK-negative group. (51.5% vs 21.9%, P=0.000 and P=0.146). There were also a higher number of patients with extrathoracic metastases (P=0.001) and brain metastases (P=0.000) in the EGFR TKI-sensitizing activating mutation group compared to the EGFR/ALK-negative group. There was no difference in PS distribution between the exon 20 group and EGFR/ALK mutation-negative group. There were more patients with extrathoracic metastasis (P=0.011) and brain metastasis

Variables	EGFR TKI-sensitizing	Exon 20 (n=20)	EGFR and ALK mutation	
	activating mutations (n=227)		negative (n=333)	
Median age, years	56 (IQR 50–63)	59 (IQR 47.8–65)	56 (IQR 49–62)	
Gender distribution, n (%)	Male: 141 (62.1)	Male: 12 (60.0)	Male: 222 (66.7)	
	Female: 86 (37.9)	Female: 8 (40.0)	Female: 111 (33.3)	
Nonsmokers, n (%)	168 (74.0)	13 (65.5)	168 (52.0)	
PS, n (%)	0–1: 110 (48.5)	0-1:12 (60.0)	0-1: 260 (78.5)	
	2 or >2: 117 (51.5)	2 or >2: 8 (40.0)	2 or >2: 73 (21.9)	
Extrathoracic metastasis, n (%)	105 (46.3)	11 (55.0)	106 (31.8)	
Brain metastasis, n (%)	29 (12.8)	06 (30.0)	10 (3.0)	
Bone metastasis, n (%)	59 (26.0)	05 (25.0)	84 (25.2)	
Liver metastasis, n (%)	39 (17.2)	02 (10.0)	29 (08.7)	
Multiple organ metastases, n (%)	19 (8.3)	02 (10.0)	17 (5.1)	

Table I Demographic and clinical profiles of the three patient cohorts

Abbreviations: EGFR, epidermal growth factor receptor; TKI, tyrosine kinase inhibitor; ALK, anaplastic lymphoma kinase; IQR, interquartile range; PS, performance status.

(P=0.000) in the exon 20 mutation group compared to the EGFR/ALK-negative group. There were no differences in bone metastasis or multiple organ involvement between the three groups.

Molecular characteristics of exon 20 mutation

The details about the types of exon 20 mutation are listed in Table 2. De novo S768I mutation was the most common type of exon 20 mutation, seen in seven patients (n=20, 35%).

Exon 20 treatment and outcome details

Our exon 20-mutated patients received the following therapies in first line: pemetrexed + platinum chemotherapy (nine patients), paclitaxel + platinum chemotherapy (three patients), oral TKI (seven patients) and no systemic treatment (one patient). The response to chemotherapy at 2 months was progressive disease in five patients, stable disease in three patients and partial response in three patients. In one patient, response evaluation was not done. Among the patients receiving oral TKIs in the exon 20 mutation subgroup, only one patient had stable disease, whereas the rest had progression. The second-line therapy was received by six patients. Five of these patients had received chemotherapy in first line and one had received oral TKI. The second-line treatment received was docetaxel in two patients, erlotinib in three patients and pemetrexed carboplatin in one patient.

There was no statistical difference in OS between different types of exon 20-mutated groups (Figure 1). However, the median survival of patients with CAC mutation was 12.0 months, which was numerically superior to the OS noted in patients with other types of exon 20 mutations.

Treatment details in EGFR-sensitizing mutation-positive patients

In all, 110 patients received TKI (gefitinib [82 patients] and erlotinib [28 patients]) and the rest of 117 patients received chemotherapy. The regimens received were pemetrexed and platinum in 84 patients, paclitaxel and platinum in 19 patients, gemcitabine and platinum in 13 patients and single-agent paclitaxel in one patient.

In patients receiving chemotherapy, 73 patients (62.4%, n=117) received second-line treatment of which 51 patients received gefitinib and 22 patients received erlotinib. Following progression on first-line TKI, 42 patients (38.2%, n=110)

Table 2 Distribution and median OS of exon 20-mutated patients

Type of exon 20 mutation	Point mutation/insertion sequence	Proportion of	Median OS in
		patients (n=20), n (%)	months
S768 I c. 2303 G>T	EILDEAYVMA S VDNPHVCR	7 (35)	6.0 (0-16.3)
H773_V774insH	EILDEAYVMASVD G NPHVCR	4 (20)	12.0 (0.2-23.98)
2319–2320 (CAC)			
V769_D770insASV	EILDEAYVMASV ASV DNPHVC	5 (25)	2.0 (1.1–2.9)
(2307–2308) GCCAGCGTG			
T790M c. 2369 C>T	EILDEAYVMASVDNPHVCRLLGICLTSTVQLI T Q	4 (20)	5.0 (1.2-8.8)

Abbreviation: OS, overall survival.

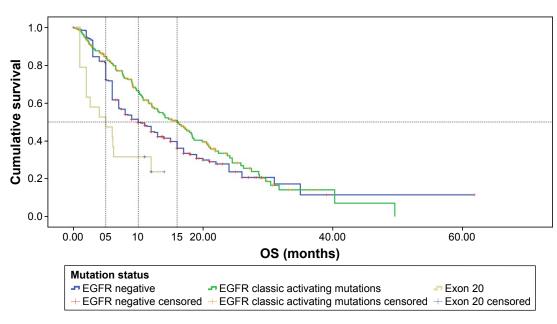


Figure I Graph of survival in different types of exon 20-mutated patients. Abbreviations: OS, overall survival; EGFR, epidermal growth factor receptor.

received second-line pemetrexed + platinum. Poor PS at progression precluded second-line therapy in the other patients.

Overall survival

The median OS in the three cohorts is shown in Figure 2. The median OS was 5 months in exon 20-mutated patients (95% CI 0.17–9.8 months), 16.1 months (95% CI 12.8–19.5 months) in activating EGFR mutated patients and 10 months (95% CI 7.9–12.1 months) in EGFR/ALK mutation-negative patients.

The median OS was significantly longer for the EGFR TKIsensitizing activating mutation group (P=0.000, log-rank test) and for the EGFR/ALK-negative group (P=0.037, log-rank test) compared to the exon 20-mutated group. On multivariate analysis, the presence of EGFR TKI-sensitizing activating mutations (hazard ratio [HR]: 0.74, P=0.013), the absence of exon 20 mutation (HR: 0.47, P=0.009) and the absence of extrathoracic disease (HR: 0.75, P=0.042) were associated with a favorable impact on survival (Table 3).

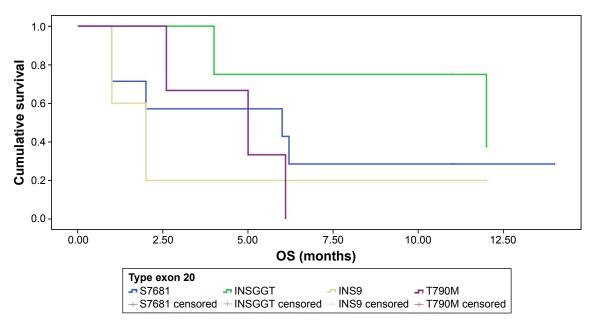


Figure 2 Graph of OS in all three cohorts. Abbreviation: OS, overall survival.

Table 3 Multivariate analysis results for OS

Variable		95% CI of HR	P-value
Presence of EGFR TKI-sensitizing	0.74	0.58–0.94	0.013
activating mutations compared to rest			
Presence of exon 20 mutations	2.10	1.20-3.65	0.009
compared to rest			
Presence of extrathoracic disease	1.34	1.01-1.77	0.042
Presence of multiple organ metastases	0.99	0.59–1.68	0.990
Presence of brain metastasis		0.85-2.33	0.178
Presence of liver metastasis		0.87–2.04	0.192
Presence of poor PS (PS \geq 2)		0.93–1.46	0.182

Note: Bold values are statistically significant.

Abbreviations: OS, overall survival; CI, confidence interval; HR, hazard ratio; EGFR, epidermal growth factor receptor; TKI, tyrosine kinase inhibitor; PS, performance status.

Discussion

Exon 20 insertions and mutations are considered a rare subset of EGFR mutations and are known to be associated with TKI resistance.5 Limited data regarding demographics, clinical features and outcomes are available from East Asian and Western populations, while we do not have much data from the Indian subcontinent.⁶⁻¹¹ The Indian population has a distinct biological makeup with respect to lung cancer. In particular, the frequency of EGFR TKI-sensitizing activating mutations differs between the East Asian and the Western world.1 The incidence of exon 20 mutation in the East Asian and Western populations ranges from 0% to 5%.67,10,11 In our study, we also found the incidence of exon 20 mutation to be 3.5%. The demographic profile of our population did not reveal an association between gender, smoking status and exon 20 mutations, although, in certain reports, a higher proportion of patients with exon 20 mutation was found to be nonsmokers.^{7,10} In general, the incidence of nonsmokers in Indian non-small cell lung cancer (NSCLC) patients is relatively high in the range of 40%–50%.¹ Our data too showed a similar trend with the proportion of nonsmokers at 60.1%.

The reported median OS in patients with exon 20 mutation is 16–26 months.⁶⁻⁹ In a report by Naidoo et al,⁶ the survival in exon 20-mutated patients was not statistically different from that in patients with exon 19 deletion. Contrary to this, in our study, exon 20 mutations were associated with very low median OS of 5 months, and on multivariate analysis, it stood out with an unfavorable HR of 2.1. Exon 20 patients seemed to harbor biologically more aggressive tumors as suggested by the fact that a higher proportion of our patients had extrathoracic metastases (55%), particularly brain metastasis (12.8%), compared to EGFR/ALK-negative patients. This may partly explain the reason for inferior outcomes. Extrathoracic disease itself

was an independent poor prognostic factor in multivariate analysis in our study. Interestingly, most of the reported series in literature have included patients who underwent surgery and had then relapsed, although these studies have calculated the OS from the date of recurrence. The biological behavior of upfront metastatic high-volume disease is different from postsurgical resection recurrent disease. These studies had a high proportion of patients, varying from 30% to 60%, who had surgery and then recurrence.⁷⁻⁹ The median OS reported from France in exon 20-mutated patients was 21 months, whereas it was only 14 months in patients with Stages IIIB and IV. The median OS in patients with Stages I-IIIA was not reached in this study.7 These data clearly indicate that survival is poorer in patients with a high-volume disease. The other reason for the poor outcome noted in our exon 20-mutated patients might be due to a significant proportion having poor PS. In all, 40% of our exon 20 mutation patients had ECOG PS 3-4, which may be a reflection of the high disease burden. Poor PS itself is a known independent prognostic factor for OS in NSCLC. Virtually, all case series reported in the literature on exon 20 mutations have only 10%-15% patients with poor PS.^{7–10,12} It is well known that exon 20-mutated patients are resistant to oral TKIs. Unfortunately, the poor PS hampered the delivery of intravenous chemotherapy. As the treatment details reveal, a majority of patients with poor PS (seven out of eight) had received oral TKI treatment. Unfortunately, all patients had progression on TKI and a dismal survival. Although exon 20 mutations are considered to be TKI resistant, certain mutations have shown to respond to TKI.13 However, no such mutation was seen in our patients. Another possible explanation for better outcomes reported in other series is the coexistence of EGFR TKI-sensitizing activating mutations with exon 20 mutations. In an article by Wu et al,¹⁴ ~40% of patients had a coexisting EGFR activating mutation. It is logical that a coexisting EGFR TKI-sensitizing activating mutation would contribute toward an improved response and improved progression-free survival with oral TKI.

Although exon 20 mutations are considered similar, in actuality, they consist of multiple mutations seen in the exon 20 region. The different types of mutations in our analysis seem to suggest a trend toward differential outcomes. The proportion of patients with different types of exon 20 mutations differs from the existing literature. In our series, the S768 I c. 2303 G>T was the most common mutation found. However, in other series (which have reported the mutation types in detail), the most common mutations found were D770_N771insSVD⁶ and S768_D770dupSVD (Wu) and V769_D770insASV.⁹ The poor outcomes seen in our series may be a reflection of the different biology of the mutation types. Similar conclusions were driven by Oxnard et al;¹⁵ they attributed an indolent behavior of exon 20-mutated cohort to T790 M mutation.

Conclusion

Exon 20 mutations in India are associated with a high-volume disease, poor PS, poor response to TKI and chemotherapy. More studies at the national and international levels are warranted for the better understanding of the biology of this disease.

Disclosure

The authors report no conflicts of interest in this work.

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