

A Case Report of a Metastatic Adenocarcinoma of Lung with Dual Positivity for EGFR Mutation and ALK Fusion

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Abstract

Non-small cell lung cancer ranks among the most lethal cancers worldwide. The rate of epidermal growth factor receptor (EGFR) mutations and echinoderm microtubule-associated protein-like 4-anaplastic lymphoma kinase (EML4-ALK) gene fusion is the most common in younger age, non-smoking Asian adenocarcinoma lung cancer patients. EGFR mutations and ALK gene rearrangements are known to be mutually exclusive and as mutual causes of resistance to EGFR-tyrosine kinase inhibitors (TKIs) or ALK-TKIs. However, rarely such co-alterations do co-exist in some clinical cases. Here we report a 62 year old male, heavy smoker with cough, hemoptysis and fatigue. Histopathological examination of bronchoscopic guided biopsy showed adenocarcinoma histology. Staging evaluation, he was found to have stage IV disease on positron emission tomography-computed tomography scan. The biopsy blocks tested positive for both EGFR mutation and ALK fusion. Patient was initiated on tablet Gefitinib 250 mg once daily. To the best of our knowledge, there has been no report of dual EGFR mutation and ALK fusion positivity from India.

Keywords: Adenocarcinoma lung; Tyrosine kinase inhibitors; EGFR; Anaplastic lymphoma kinase; EGFR/ALK Co-alteration

Introduction

Lung cancer ranks among one of the most common and most lethal malignancies. Adenocarcinoma is the most commonly diagnosed histological subtype of non-small cell lung cancer (NSCLC) worldwide. Epidermal growth factor receptor (EGFR) mutations and echinoderm microtubule-associated protein-like 4-anaplastic lymphoma kinase (EML4-ALK) gene rearrangements are usually seen in adenocarcinoma histological subtype of lung. EGFR mutations and ALK rearrangements are mutually exclusive i.e ALK rearrangement is not seen in adenocarcinoma with EGFR mutation and vice-versa. Simultaneously, ALK rearrangements and EGFR mutations are known to be mutual causes of resistance to EGFR and ALK targeted therapies respectively.

To the best of our knowledge, there is no reported case of coexistent EGFR and ALK positive metastatic NSCLC from India. Here we report a case of a metastatic NSCLC found to be positive for both EGFR mutation and ALK rearrangement.

Case Presentation

A 62 year old male, working as a mechanic presented with cough with expectoration, intermittent hemoptysis and fatigue of 6 months duration. He was a heavy smoker for more than 40 years and had a history of alcohol abuse for more than 30 years. He had undergone coronary artery bypass grafting for ischemic heart disease 7 years ago after which he quit smoking and alcohol. On examination, he was moderately built and nourished with an Eastern co-operative oncology group (ECOG) performance status 1. On respiratory system

examination, had a midline sternal healed scar, rhonchi in the right supraclavicular, infra-clavicular and axillary region. Other systems examination was unremarkable. Routine Haemogram and biochemistries were normal. Chest X-ray showed right upper zone non-homogenous opacity. Bronchoscopy showed a large right endobronchial ulcero-proliferative growth with right upper lobe collapse.

Bronchoscopic biopsy showed features conclusive of adenocarcinoma histology; the neoplastic cells were positive by Immunohistochemistry (IHC) staining for Cytokeratin 7 (CK 7), Thyroid transcription factor 1 (TTF1) and Napsin A antibodies. A positron emission tomography-computed tomography (PET-CT) scan showed a perihilar right upper lobe mass lesion (3.8 × 3.5 cm; standardized uptake value (SUV)max 24.3), metastatic mediastinal lymph nodes and multiple bilateral fluorodeoxyglucose (FDG) avid hypo-dense lesions in both the kidneys (Figures 1 and 2).

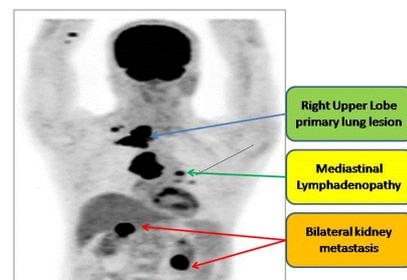


Figure 1: Baseline PET CT scan showing FDG avid right upper lobe lung lesions, mediastinal lymphadenopathy and bilateral kidney metastasis.

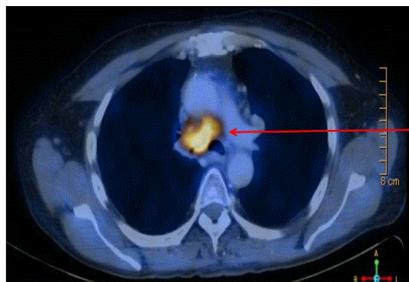


Figure 2: Baseline PETCT scan showing FDG avid right upper lobe perihilar lesions, mediastinal lymphadenopathy.

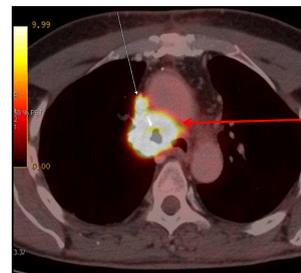


Figure 5: Re-assessment PETCT scan after 3 months of tablet Gefitinib showing the progression of the FDG avid right upper lobe perihilar lesion and mediastinal lymphadenopathy.

The paraffin blocks tested positive for EGFR mutation for G719X exon 18 (performed by amplification-refractory mutation system (ARMS)[™] and Scorpions Real time PCR method) (Figure 3) and EML4-ALK fusion (done by IHC for ALK D5F3 and confirmed by fluorescent in situ hybridization (FISH) analysis) (Figure 4). Patient was started on Tablet Gefitinib 250 mg once daily; he was initially better symptomatically for 1 month, but PET-CT scan after 3 months of starting Tablet Gefitinib, showed progressive disease (Figure 5).

Patient was recommended but was not affording for ALK-TKI (Tablet Crizotinib) treatment. He was treated with 3 weekly chemotherapy with intravenous pemetrexed 500 mg/m² and carboplatin area under curve (AUC) 5, both given on day 1 of each cycle. After 4 cycles of chemotherapy, patient was asymptomatic and reassessment with contrast enhanced computed tomography (CECT) scan showed a partial response. He was planned to continue on maintenance pemetrexed.

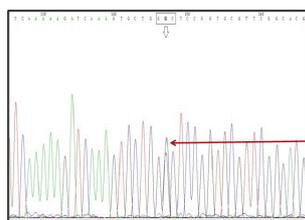


Figure 3: Tumour cells tested positive for EGFR mutation at exon 18 by the ARMS scorpion method.

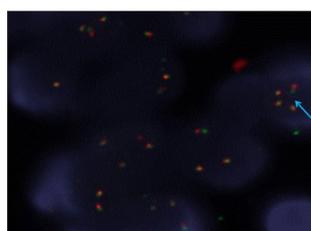


Figure 4: Tumour cells staining positive for EML4-ALK translocation by fluorescent in situ hybridization (FISH) method.

Discussion

NSCLC with adenocarcinoma histology have been known to be genetically diverse. The understanding of its genetic diversity has led to the discovery of new molecular-targeted approaches. In particular, *EGFR* mutations and *ALK* gene rearrangements are the two most studied targets for the treatment of adenocarcinoma lung.

Similar to *EGFR* mutations, *ALK* rearrangements occur almost exclusively in young, Asian adenocarcinoma patients, who are either never or former smokers [1]. *ALK* rearrangements are not usually found in tumors with mutations in *EGFR* or Kirsten rat sarcoma viral oncogene homolog (*K-RAS*) mutated patients [2,3]. *ALK* rearrangements and *EGFR* mutations have been largely reported to be mutually exclusive [1] and as mutual causes of resistance to *EGFR* tyrosine kinase inhibitors (TKIs) and *ALK* TKI [4,5]. There is a possibility that similar to the prevalence of *EGFR* mutations, the co-existence of *ALK* rearrangement may be higher in East-Asian lung adenocarcinoma patients as compared to the Caucasians [6-8]. Recently, rare case reports have emerged noting the coexistence of *EGFR* mutation in *EML4-ALK* fusion positive patients [1,7-12]. However, the frequency of such co-alterations is yet to be fully described [9-11]. Yang et al. in a case series showed the frequency of dual *EGFR* and *ALK* co-alteration to be ~1.3% and it was possibly higher in Chinese patients compared to the Caucasians [7]. Advanced molecular genetic technologies such as target next generation sequencing (NGS) with sensitivity to detect <1% of *EGFR* mutation cells and FISH study for *ALK* rearrangements have enabled the detection of increasing numbers of cases with dual co-alteration [13]. It will be the key to detect many such new cases in the future and possibly alter the treatment strategies for such cases. Two hypotheses have been made to explain the existence of this concomitant alteration-one of them being the differential genetic alteration occurring in the heterogenous tumor cell population and the other being the presence of a co-activating *EGFR* mutation in a homogenous cell population which has a pre-existing *ALK* gene rearrangement [12]. The most common co-existing *EGFR* mutations reported earlier included the Exon 19, L858R, L747P, and E868K [5]. In our patient, G719X exon 18 mutation was observed which is known to be sensitive to *EGFR* TKIs.

Knowing the possibility of Dual *EGFR/ALK* co-alteration may be of clinical relevance in terms of treatment strategies. First-line *EGFR*-TKIs in TKI sensitive-*EGFR* mutant NSCLC have been shown to be superior to chemotherapy in terms of response rate, progression-free survival (PFS), quality of life and even have better survival rates

[14,15]. Patients with ALK rearrangement benefit from the TKI Crizotinib [16]. However, for patients with concomitant EGFR mutations and ALK rearrangement, few reports are available regarding the clinical activity of EGFR-TKIs and ALK-TKIs [9-11]. Differential phosphorylation of EGFR or ALK which can be measured by IHC or the low burden of EGFR mutation in these cases might contribute to differences in sensitivity to EGFR-TKIs or Crizotinib. In a study by Won et al. concomitant EGFR and ALK gene alteration rates were noted in 0.3% patients [17]. When they studied the response rates in the dual positive cases with either of the TKIs, they found that the response rates with Gefitinib was disappointing while ALK inhibitors showed promising benefit in terms of response rate and progression free survival, similar to exclusive ALK rearrangement patients [16]. Our patient was treated with Gefitinib; Crizotinib was not used in view of financial constraints. Similar to the study by Won et al. we also observed a poor response with Gefitinib in our patient. This suggests that the dual positive patients are resistant to EGFR TKIs while retaining sensitivity to ALK inhibitors and that the clinical outcome of these patients with first line ALK inhibitors may be substantially better than with first line EGFR TKIs [17]. But the question of whether missing the concomitant EGFR mutation in ALK rearrangement positive patients has any impact on treatment outcome and prognosis is yet to be fully answered. The prevalence and clinical relevance of co-alterations in these two driver genes require detailed investigation especially in the Asian countries which has high prevalence of EGFR.

Conclusion

EGFR/ALK co-alterations could define a specific subgroup that has diverse, although mostly favorable, responses to first-line TKIs. Testing of the relative phosphorylation levels of EGFR/ALK and EGFR mutations might help to guide the selection of TKIs in clinical practice. Molecular mechanisms underlying responsiveness and resistance to EGFR-TKIs and ALK-TKIs, and potential combination or sequential treatment require further investigation in this specific subgroup of co-alterations. The true incidence, biology and treatment strategies of concomitant EGFR/ALK alteration need further evaluation in a large group of patients and for determining the effective management of such patients.

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