Detection of rare targetable EGFR variant in Metastatic Non-small Cell Lung Carcinoma by next Generation Sequencing: A Case Report

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ABSTRACT

Genetics Section

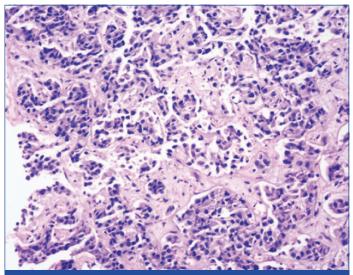
Single gene assays for variants in Epidermal Growth Factor Receptor (EGFR) demonstrate actionable and sensitising mutations in majority of the cases. However, the emergence of next generation sequencing platforms has facilitated the detection of newer variants in the already known drivers of lung carcinomas, which may be clinically actionable. Mutations in exons 18 to 21 of EGFR are widely characterised in literature, and rare unusual mutations in these regions are constantly being demonstrated recently. This report describes a rare exon 18 insertion variant in EGFR gene in a 61-year-old patient of non-small cell lung carcinoma, which is potentially actionable, thus highlighting the need of next generation sequencing based platforms in this era of precision medicine.

Keywords: E709 codon, Exon18, Epidermal growth factor receptor, Molecular testing

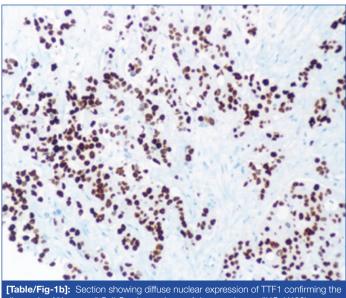
CASE REPORT

A 61-year-old male patient presented with the chief complaints of productive cough and occasional bouts of haemoptysis of 3-4 months' duration. There was associated fatigue and dyspnoea on exertion. On physical examination, the patient had an obvious conjunctival pallor. However, there were no palpable lymph nodes, and the organ systems were unremarkable. There was no history of associated fever or any medically relevant history. Contrast Enhanced Computed Tomography (CECT) of thorax (evaluated elsewhere) revealed an approximately 5 cm × 5 cm mass in the lower lobe of right lung, along with tiny nodules in the upper lobe and a few small mediastinal lymph nodes. Routine hematologic and biochemical investigations were unremarkable. Bronchoscopy performed revealed infiltrates and bulges over right bronchus intermedius, suspicious of malignancy. A provisional diagnosis of a lung primary cancer. Histopathological and Immunohistochemical (IHC) examination of a core of the biopsy revealed a non-small cell lung carcinoma, favouring adenocarcinoma with a predominantly acinar pattern, composed of malignant cells, with moderate cytoplasm, irregular nuclei with dispersed chromatin, along with negative PDL1 (Programmed Death Ligand-1) staining. IHC was positive for Thyroid Transcription Factor 1 (TTF1), favouring the adenocarcinoma histology [Table/Fig-1a,b]. Molecular testing performed at this diagnostic time point was negative for sensitising mutations in EGFR {by real-time Polymerase Chain Reaction (PCR)}, using an FDA (Food and Drug Administration) approved kit by QiagenTM, USA), Anaplastic Lymphoma Kinase (ALK) (IHC) and ROS1(c-ros oncogene 1) rearrangements (by fluorescence insitu hybridisation). Positron Emission Tomography (PET-CT) scan revealed a metabolically active right lung lesion with lymph node and bony involvement, mild right sided pleural effusion, pleural and bony (rib and ischium) deposits. The patient was started on Pemetrexed (500 mg/m²) and Cisplatin (75 mg/m² in divided doses) based chemotherapy and 6 cycles of the same were administered.

Post-six cycles PET CT re-evaluation revealed good partial response to treatment with residual metabolically active right lung lesion with bony involvements and right pleural deposits with pre-dominantly metabolically inactive mediastinal lymph nodal abnormalities. Magnetic Resonance Imaging (MRI) brain showed no metabolically active or enhancing parenchymal/meningeal lesion in brain. The patient's performance status dipped at this follow-up.



[Table/Fig-1a]: Tissue biopsy of lung mass demonstrating acinar pattern of growth. H&E Section (X400).



diagnosis of Non-small Cell Carcinoma Lung- Adenocarcinoma (IHC, X400).

At this point, the original tumour was subjected to next generation sequencing using Oncomine Focus Assay Kit (ThermoFischer

Scientific, Lifetechnologies Inc., USA) and the libraries were sequenced on Ion Torrent S5 platform. DNA based assay revealed a mutation in exon 18 of the EGFR gene which was p.(E709_T710delinsD):c.2127_2129delAAC [Table/Fig-2], and a mutation in TP53 gene: p.R248Q [Table/Fig-3] while the real-time PCR for EGFR was negative for any mutations. In view of the EGFR mutation detected, the patient was started on oral tyrosine kinase inhibitor therapy: Afatinib which was continued for seven months, during which the patient's general condition was fair.



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However, subsequently, the patient presented with increasing cough, and a re-evaluation done using PET CT revealed increase in extent and metabolic activity of the right lung lesion causing complete distal collapse of lower lobe. Metabolically, active right pleural deposit was noted which was not seen earlier. Lymph nodes were unchanged. Left rib lesion showed increase in extent with persistent metabolic activity. Right ischial lesion showed significant increase in metabolic activity; new lesions were noted in the pleura on the right side, left rib, and right ischium bone, indicating progressive disease. The patient is now being treated with second line (Gemcitabine and Carboplatin) based on chemotherapy.

DISCUSSION

Tier 1 mutations including EGFR, ALK, ROS1, NTRK and BRAF V600E form the first line testing as per the NCCN guidelines for lung malignancies. They are described in ~40% cases of NSCLC,

with EGFR occurring in 25% cases of NSCLC [1,2]. The most common somatic mutations in the EGFR gene include inframe deletion of exon 19 reported in ~45% cases [3], and single nucleotide variants in exon 21; L858R in ~40% cases [3], which have shown dramatic response rates and better Progression Free Survival (PFS) when treated with oral EGFR TKIs (Gefitinib, Erlotinib). Rare EGFR mutations, such as G719X (~3%) and the L861Q (~2%) show higher rates of response and a prolonged PFS [4-6]. Other less common EGFR mutations have not been completely characterised, for example, the exon 18 deletion/ insertion mutation delE709_T710insD described here. This mutation has been reported in the Catalogue Of Somatic Mutations In Cancer (COSMIC) database [7] in 14 NSCLC cases as well as in My Cancer Genome (2 cases) [6]. In a report on rare EGFR mutations, Sousa AC et al., reported alterations in the E709 codon with one patient harbouring the same mutation and was not treated with first line TKI [7]. They reported an overall survival of 17 months in their case. On the second follow-up, the patient was in clinical stage IVB and he was administered Erlotinib, subsequent to which he showed a PFS of 10 months. However, unlike the present scenario, this case did not harbour any other concurrent mutations, which may explain the better PFS in their patient.

The overall reported frequency of this mutation is ~0.1% [8,9] in previous reports, and almost all of them have shown increased sensitivity and response to Erlotinib [5,9]. This patient, however, was treated with afatinib, and showed only partial response, eventually culminating in a progressive disease. The plausible explanation for the same can also be attributed to the co-existent pathogenic variant in the TP53 gene which may induce resistance, as has been well documented in literature. The TP53 mutation detected in the present patient has been reported pathogenic and confirmed somatic in the International Agency for Research on Cancer (IARC) TP53 database [10].

CONCLUSION(S)

This variant is very rare and is usually missed on single gene testing, which is widely done in NSCLC cases. This report highlights the importance of next generation sequencing based assays, in the detection of newer mutations which may be potentially actionable and can benefit the treatment and overall outcomes of a patients. However, the confirmation of the same requires in-vitro studies.

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PLAGIARISM CHECKING METHODS: [Jain H et al.]

• iThenticate Software: May 15, 2020 (13%)

• Plagiarism X-checker: Mar 09, 2020

Manual Googling: May 01, 2020

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