



Predictors of Acquired T790M Mutation in Patients Failing First- or Second-Generation Epidermal Growth Factor Receptor-Tyrosine Kinase Inhibitors

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Chee-Shee Chai¹
Chong-Kin Liam²
Mau-Ern Poh²
Diana Bee-Lan Ong³
Yong-Kek Pang²
Phaik-Leng Cheah³
Gwo-Fuang Ho⁴
Adlinda Alip⁴

¹Department of Medicine, Faculty of Medicine and Health Science, University Malaysia Sarawak, Kota Samarahan, Sarawak, Malaysia; ²Department of Medicine, Faculty of Medicine, University of Malaya, Kuala Lumpur, Malaysia; ³Department of Pathology, Faculty of Medicine, University of Malaya, Kuala Lumpur, Malaysia; ⁴Department of Clinical Oncology, Faculty of Medicine, University of Malaya, Kuala Lumpur, Malaysia

Background: This study aims to determine the predictors of acquired *exon 20* T790M mutation in advanced non-small cell lung cancer (NSCLC) patients harbouring sensitizing epidermal growth factor receptor (EGFR) mutation following the failure of first- or second-generation EGFR tyrosine kinase inhibitor (TKI).

Methods: This is a retrospective observational study of NSCLC patients with sensitizing EGFR mutation experiencing disease progression (PD) whilst on first- or second-generation EGFR-TKIs with subsequent investigations to detect acquired T790M mutation at the University of Malaya Medical Centre from 1st January 2015 to 31st December 2017.

Results: A total of 87 patients were included. Upon PD, acquired T790M mutation was found in 55 (63.2%) patients and was significantly more common in patients who achieved partial response (PR) whilst on the EGFR-TKIs ($p = 0.008$) or had new lung metastasis upon PD ($p = 0.048$). It was less frequent in patients who developed new symptomatic brain lesions ($p = 0.021$). Patients with *exon 19* deletion were more likely to acquire T790M mutation compared to those with *exon 21* L858R point mutation ($p = 0.077$). Multivariate analysis revealed PR whilst on EGFR-TKI treatment was an independent predictor of acquiring T790M mutation ($p = 0.021$), whereas development of new symptomatic brain lesions ($p = 0.034$) or new lymph node metastases ($p = 0.038$) upon PD was independently against acquiring T790M mutation. Patients with *exon 19* deletion were more likely to acquire T790M mutation compared to those with *exon 21* L858R point mutation (odds ratio: 2.3, 95% confidence interval: 0.84–6.25, $p = 0.104$).

Conclusion: The best tumour response of PR to first- or second-generation EGFR TKI treatment independently predicts acquired T790M mutation. Patients with *exon 19* deletion are likely to acquire T790M mutation. This would prove useful for clinicians to prognosticate and plan subsequent treatments for patients with advanced NSCLC harbouring EGFR mutations.

Keywords: non-small cell lung cancer, epidermal growth factor receptor, acquired T790M mutation, independent predictor, tyrosine kinase inhibitor

Introduction

Lung cancer, 85% of which are non-small cell lung cancer (NSCLC), remains the leading cause of cancer mortality globally.¹ Upon diagnosis, the majority of NSCLC patients have locally advanced or metastatic disease. Conventional first-line chemotherapy in these patients confers a dismal median overall survival of 8–10 months and a 2-year survival rate of 11%.^{2,3}

The discovery of mutations of the epidermal growth factor receptor (EGFR) has completely revolutionized the management of patients with advanced NSCLC.

Correspondence: Mau-Ern Poh
Department of Medicine, Faculty of
Medicine, University of Malaya, Kuala
Lumpur 50603, Malaysia
Tel +60 3 7949 4422
Fax +60 3 7955 2253
Email ernepoh@gmail.com