





# Cost-effectiveness of biomarker testing for treatment choices in advanced non-small cell lung carcinoma: impact of diagnostic strategies and their turnaround times

#### To the Editor:

Lung cancer remains the main cause of cancer death in the world, not only because of its high frequency but also because it is often diagnosed at poor-prognosis, advanced stages. A better understanding of the molecular mechanisms involved in tumour initiation and progression has permitted to develop new targeted therapies and immunotherapies and molecular testing has now become a standard in care for patients with advanced non-small cell lung cancer (NSCLC). In the era of personalised medicine, the study by LOUBIERE *et al.* [1] demonstrated that molecular testing prior to treatment initiation is a cost-effective approach in the management of patients with advanced NSCLC. Beside the calculation of global incremental cost-effectiveness ratios, a major interesting point of the study by LOUBIERE *et al.* [1] is their proposal of a step-by-step evaluation of the costs related to diagnosis, treatments and inpatient care. In this correspondence, I intend to further comment on the diagnosis-related costs and their potential impact on the total costs.

LOUBIÈRE et al. [1] used reference prices for EGFR, KRAS and ALK testing of 180.90, 213.30 and 110.70 EUR per test, respectively. Nevertheless, according to the prices provided by the French Ministry of Health, the costs of molecular analyses can highly vary from one diagnostic strategy to another. For example, to analyse EGFR, 180.90 EUR in fact indicates the price of two-exon analysis, whereas the price of four-exon EGFR analysis (i.e. exons 18-21, as mentioned in the study by LOUBIERE et al. [1]) is 315.90 EUR. A global quote for concomitant EGFR and KRAS analysis in NSCLC is 459.00 EUR but the use of next-generation sequencing (NGS) with a coverage of <20 kilobases, commonly used in NSCLC samples for EGFR and KRAS analyses, is set at 882.90 EUR [2, 3]. In this manner, it is very difficult to estimate the real costs of molecular analyses per patient in France. In their study, LOUBIÈRE et al. [1] highlighted that the diagnosis-related costs (i.e. histopathological diagnosis and biomarker testing) only represented <3% (i.e. 513.00-529.00 EUR, including biomarker testing) of the total cost for the management of a patient with NSCLC (i.e. between 17000 and 20000 EUR according to LOUBIERE et al. [1]), whereas inpatient care represented between 45% (biomarker testing strategy) and 68% (no testing strategy) of the total costs [1]. The diagnosis-related costs could, in this manner, vary from one diagnostic strategy to another but their contribution in the total cost seems to remain minimal, considering the costs of inpatient care and treatments.

Beyond the costs of the molecular tests themselves, various diagnostics methods nevertheless imply different turnaround times (TATs). The TATs from test request to result delivery could, in this manner, in my opinion, strongly influence the cost of the initial management of a patient. Indeed, although the College of American Pathologists, International Association for the Study of Lung Cancer, Association for Molecular Pathology and European external quality assessment schemes have established a benchmark TAT target of 10 working days for the delivery of molecular results to the oncologist, TATs exceeding this

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Reducing the turnaround times of molecular analyses in patients with advanced lung cancers could help to reduce the costs of the patients' management by permitting rapid treatment choices http://ow.ly/Zv7230jAQv4

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recommendation are not rare, with some biomarkers results being available only after several weeks for some patients [4–7]. During the time waiting for treatment initiation, the tumours continue to grow and the patients' deterioration could require additional inpatient care, causing increased costs [8]. In this manner, it could be valuable to use first-line fast molecular diagnosis methods able to provide *EGFR* and *KRAS* results in a single day (*e.g.* real-time PCR), as well as those for *ALK* and now *ROS1* and PD-L1 (using dedicated first-line immunohistochemistry) [9, 10]. Such methods could permit rapid treatment choices to optimise the therapeutic management of patients with NSCLC and to reduce inpatient care-related costs during the initial management of patients newly diagnosed with advanced NSCLC. This could be especially relevant for patients' ongoing acute deterioration.

Finally, I fully agree with LOUBIERE *et al.* [1] that it is likely that, despite the impressive analytical capacities of NGS, clinicians will keep on using a combination of both step-by-step and NGS strategies in the coming years to better match with the management of patients. Beyond expanding the number of molecular analyses (and, to date, expanding their related costs) accompanying the availability of new treatments, the reduction of biomarker analysis TATs could also represent progress for the patients' management, from a medical but also medicoeconomic point of view. This medicoeconomic evaluation of various methods and of the global impact of their TAT on the whole management of patients must be the subject of future studies.

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## From the authors:

We would like to thank A. Uguen for his correspondence with comments on our article [1]. In this prospective cohort study, the maximum cost for the diagnosis of advanced non-small cell lung cancer (NSCLC) was estimated to be 620 EUR, including EGFR/KRAS/ALK testing. In 52% of the patients in the "at least one biomarker status known", the diagnostic strategy included EGFR/KRAS and ALK testing

simultaneously, incurring a mean cost around 550 EUR. In 32% of the "at least one biomarker status known" group, the diagnostic strategy consisted first of a molecular analysis for KRAS with simultaneous immunohistochemistry (IHC) ALK testing, and second a targeted analysis for EGFR, followed by an additional validation by fluorescence *in situ* hybridisation assay for ALK IHC positive cases, for example at the time of second-line treatment initiation. For those patients, the total cost of the diagnostic strategy was estimated to be 616 EUR. All costs were based on reimbursed prices provided by the French Ministry of Health.

Depending on the point of view adopted, be it societal, healthcare payer or healthcare provider, the type and level of accuracy for cost estimates are not the same [2]. From the healthcare payer perspective we adopted in our study, using prices to assess diagnosis costs is the recommended approach [2, 3]. Nonetheless, from the point of view of the healthcare provider (i.e. hospital), we acknowledge the inadequacy of using prices rather than accurate costs, and as reported by A. Uguen as well, the difficulty in capturing such costs. For example, there is limited national data on the magnitude of each component of those biomarker testing costs, such as labour (including but not limited to technician/engineer mobilised during all the steps and biologist or pathologist time), consumables, and equipment and overhead costs. Thus, more reliable estimates of diagnosis costs would be useful inputs for economic evaluation of strategies based on specific biomarker testing with appropriate targeted or non-targeted therapies from the providers' perspective. Therefore, we are conducting a micro-costing analysis on the costs of each technology of testing used in the IFCT-PREDICT.amm study in order to generate accurate cost estimates for biomarker testing strategies through a detailed measurement and valuation of resource inputs. In addition, we are conducting the same analysis on next-generation sequencing (NGS) techniques performed in the same certified genetic centres. We favoured a micro-costing approach, which is particularly well-suited to capturing the variations between settings and patients, as the total costs of EGFR and KRAS mutations testing (including DNA extraction) are expected to vary across French National Cancer Institute-certified genetic centres according to the different mutation analysis strategies.

Considering the remark of A. Uguen on the turnaround time (TAT), we acknowledge that the risk inherent to a step-by-step approach for the NSCLC diagnosis would be an increased proportion of patients displaying an unknown oncogenic driver prior to starting first-line treatment, due to excessive TAT to obtain the second set of molecular analyses. The IFCT Biomarker France study showed that 23% of NSCLC patients had started treatment before they received the results of the molecular analysis [4]; the main reason being the too long TAT of biomarker results in those specific cases. In a retrospective study conducted in Canada between April 2010 and March 2013, the prevalence of biomarker testing rose from 46% in the first year to 60% during the last year of the study for EGFR testing, and from 5% to 40% for ALK testing, with 19% of patients having started chemotherapy before their biomarker results became available [5]. These results are very similar to those found in our study.

Although the small panel NGS recently implemented in France allows simultaneous analysis of several molecular targets (EGFR, HER2, BRAF, *etc.*), it does not allow identification of the presence of gene rearrangements, such as ALK or ROS [6]. Moreover, the new targetable alterations mainly concern rearrangements (RET, NTRK, NRG1, NTRK, *etc.*) or complex alterations such as the skip exon 14 mutation of MET which also require the concurrently use of several techniques. Finally, the small panel NGS currently used does not measure the mutation burden of the tumour, which appears to be an emerging biomarker for the prescription of immunotherapies. In the coming years, it is likely that several diagnostic techniques will still be required to examine all predictive biomarkers in thoracic oncology, always asking the question of the step-by-step approach and risking further lengthening the TAT. To conclude, it is hoped that in future, though the timeframe is uncertain, RNA or exome sequencing approaches with hybrid capture will make all predictive biomarkers immediately available before starting first-line treatment at reasonable cost. The Biomarkers France #2 study is being initiated precisely in order to capture the changes that occurred with the implementation of NGS and includes more than two-thirds of the advanced NSCLC patients tested in 2017.

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