




CONSENSUS STATEMENT

The American Cancer Society National Lung Cancer Roundtable strategic plan: Methods for improving turnaround time of comprehensive biomarker testing in non-small cell lung cancer

Sinchita Roy-Chowdhuri MD, PhD¹  | Haresh Mani MD² | Adam H. Fox MD³ | Anne Tsao MD⁴ | Lynette M. Sholl MD⁵ | Farhood Farjah MD⁶  | Bruce E. Johnson MD⁷ | Raymond U. Osarogiagbon MBBS⁸ | M. Patricia Rivera MD⁹ | Gerard A. Silvestri MD³ | Robert A. Smith PhD¹⁰  | Ignacio I. Wistuba MD¹

¹Division of Pathology and Laboratory Medicine, The University of Texas MD Anderson Cancer Center, Houston, Texas, USA

²Department of Pathology, Inova Fairfax Hospital, Falls Church, Virginia, USA

³Division of Pulmonary Critical Care, Allergy and Sleep Medicine, Medical University of South Carolina, Charleston, South Carolina, USA

⁴Department of Thoracic/Head and Neck Medical Oncology, The University of Texas MD Anderson Cancer Center, Houston, Texas, USA

⁵Brigham and Women's Hospital and Department of Pathology, Harvard Medical School, Boston, Massachusetts, USA

⁶Department of Surgery, University of Washington, Seattle, Washington, USA

⁷Department of Medical Oncology, Dana-Farber Cancer Institute, Harvard Medical School, Boston, Massachusetts, USA

⁸Multidisciplinary Thoracic Oncology Program, Baptist Cancer Center, Memphis, Tennessee, USA

⁹Department of Medicine, Division of Pulmonary and Critical Care Medicine, Wilmet Cancer Institute, The University of Rochester Medical Center, Rochester, New York, USA

¹⁰Center for Early Cancer Detection Science, American Cancer Society, Atlanta, Georgia, USA

Correspondence

Sinchita Roy-Chowdhuri, Division of Pathology and Laboratory Medicine, The University of Texas MD Anderson Cancer Center, 1515 Holcombe Boulevard, Houston, TX 77030, USA.

Email: sroy2@mdanderson.org

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Abstract

Comprehensive biomarker testing for patients with non-small cell lung cancer is critical for selecting appropriate targeted therapy or immunotherapy. Ensuring timely ordering, processing, and reporting is key to optimizing patient outcomes. However, various factors can prevent or delay patients from being offered the option of treatment selection based on comprehensive biomarker testing. These factors include problems with access to testing, tissue adequacy, turnaround time, and health insurance coverage and billing practices. Turnaround time depends on several logistical and tissue handling factors, which involve institutional policies, processes, resources, testing methodology, and testing algorithms that vary across different practices. In this article, the authors identify key factors that prolong biomarker testing turnaround time, propose strategies to reduce it, and present a

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process map to aid physicians and key organizational stakeholders in improving testing efficiency.

KEYWORDS

comprehensive biomarker testing, non-small cell lung cancer, process map, quality assessment, quality improvement, quality metrics, testing algorithms, testing efficiency, tissue adequacy, turnaround time

INTRODUCTION

Optimal treatment options for patients with non-small cell lung cancer (NSCLC) require comprehensive biomarker testing to guide clinical decision making.¹ Clinical practice guidelines from professional societies, including the National Comprehensive Cancer Network, the College of American Pathologists (CAP)/International Association for the Study of Lung Cancer (IASLC)/Association of Molecular Pathologists (AMP), the American Society of Clinical Oncology, and the European Society for Medical Oncology, all recommend biomarker testing for patients with newly diagnosed NSCLC.¹⁻⁵

In addition to identifying targetable genomic biomarkers, evaluating programmed death-ligand 1 (PD-L1) expression by immunohistochemistry (IHC) can identify patients who may benefit from immune checkpoint inhibition as initial monotherapy or in the adjuvant setting.⁶⁻¹⁰ Therefore, obtaining both genomic and immunohistochemical biomarkers is essential to determine the optimal treatment for patients with NSCLC. Further highlighting the need for comprehensive biomarker testing, patients with *EGFR*, *ALK*, and *ROS1* alterations are unlikely to respond to frontline immunotherapies.¹¹ Administration of immunotherapy before some tyrosine kinase inhibitor therapies exposes patients to increased and avoidable toxicity.^{8,12,13}

Despite the growing evidence of clinical benefit from biomarker-driven treatment selection, many patients with metastatic NSCLC do not receive testing for biomarkers as recommended by existing guidelines.¹⁴⁻¹⁸ A survey by the IASLC reported higher testing rates for biomarkers with the longest standing approvals for targeted therapies, such as *EGFR* and *ALK* (ranging between 83% and 94%); however, biomarkers like *ROS1*, *BRAF*, *KRAS*, *MET*, *RET*, and *ERBB2* (*HER2*), which guide the selection of specific targeted therapies that were approved after 2013, had much lower rates of testing (ranging from 20% to 69%).¹⁹ With the increasing number of biomarkers needed to guide therapeutic decisions for patients who have NSCLC, *sequential* testing strategies for single gene abnormalities now run the risk of depleting available tissue and increasing the turnaround time (TAT) from initial test requisition to availability of actionable results.^{14,15,20} Therefore, a strategy of timely *comprehensive* biomarker testing, in which all biomarkers that test eligibility for approved therapies are performed simultaneously, is increasingly advocated.^{21,22} We support the practice of comprehensive biomarker testing, whenever possible, to identify optimal treatments for patients. It is worth clarifying because there is variation in actual practice for assays used and the

comprehensiveness of biomarkers assessed across different institutions. Over time, the term *biomarker testing* throughout this statement refers to assessing any treatment-predictive biomarkers using available assays.

In addition to the importance of comprehensive biomarker testing and addressing other factors that delay the time from diagnosis to treatment, the TAT of biomarker results is needed to capitalize on the improved outcomes offered by effective targeted and immune therapies. Evidence from single centers shows that TAT varies and may span from within the CAP/IASLC/AMP recommended 10 working days to well over 1 month.^{17,23-27} Evidence also suggests that delays in TAT are a barrier to optimal care.^{14,20,24} One study indicated that approximately 20% of patients with metastatic NSCLC undergo chemotherapy before genotyping results for *EGFR* or *ALK* are available.²⁰ Another nationwide biomarker survey in France demonstrated that long TATs caused local tumor boards to recommend conventional chemotherapy rather than waiting for biomarker test results.²⁴ To better define the factors that influence TAT, it is essential to recognize the series of interconnected steps occurring across multiple disciplines, each of which may contribute to lengthy TAT through poor coordination, inefficiencies, and lack of communication. This article focuses on the key issues related to prolonged TAT, from tissue delivery to the pathology department to reporting biomarker test results.

The tissue journey: From diagnosis to biomarker testing

A primary care clinician (physician or nurse practitioner) or specialist (commonly pulmonologist and thoracic surgeon) frequently initiates the initial evaluation of suspected NSCLC after symptoms, a radiologic abnormality found incidentally, or by low-dose computed tomography screening, which leads to tissue procurement by an interventional radiologist, pulmonologist, or thoracic surgeon. The procured tissue is then evaluated by a pathologist who confirms the diagnosis of lung cancer and subtypes the tumor (e.g., small cell lung cancer, NSCLC, etc.). For patients who have biomarker testing performed, testing is usually initiated at some point after a diagnosis of NSCLC is made. Some health care physicians participating in the diagnostic process may receive pathology results and order biomarker testing. However, this process is not initiated in most instances until the patient sees a medical oncologist, which may be several days to weeks after the initial diagnosis. In many cases,

patients are not offered biomarker testing and proceed immediately to conventional chemotherapy; other patients offered biomarker testing might reject it to avoid treatment delay or unreimbursed expenses; finally, some patients may not receive comprehensive biomarker testing, and their cancer may harbor an actionable mutation that is not assessed.²⁸ Each instance is a missed opportunity to offer the patient a more informed and potentially superior treatment option.

When biomarker testing is ordered, a pathologist evaluates whether there is adequate tissue. In cases where the tissue is deemed adequate, different laboratory processes are required before it is either directed to in-house biomarker assays or sent to a commercial or reference laboratory for testing. However, if the available tissue is deemed inadequate, physicians and patients must consider the impact on treatment planning and the risks associated with a repeat biopsy to complete biomarker testing. They may consider plasma-based testing as an alternative.

Once biomarker testing is initiated, PD-L1 IHC results are usually available within days (approximately 2–5 days); however, molecular test results for the remaining biomarkers may become available individually over several days (for single-gene assays) or weeks (for comprehensive, panel-based molecular profiling). The CAP/IASLC/AMP guideline for tissue-based molecular testing recommends that, once a diagnosis of NSCLC is rendered, tissue should be sent to the testing laboratory within 3 working days of receiving test requests for reference laboratories and within 24 hours for in-house laboratories, with a TAT of 10 working days or less for test results to be available (from sample receipt in the molecular laboratory to reporting of all results).^{2,29}

The challenge of timely and comprehensive biomarker testing

Timely ordering, completion, and reporting of comprehensive biomarker testing results have important implications for patient care. Several studies demonstrate that prolonged TAT (≥ 2 weeks) for biomarker test results can adversely influence clinical decision making; physicians may choose potentially inferior therapy instead of waiting for biomarker test results.^{14,20,24} The decision to proceed with systemic therapy without biomarker guidance may be influenced by patients' or physicians' anxiety. Still, it may indicate clinical urgency to begin treatment, such as declining performance status. Long TAT of genotyping results may prevent up to 50% of patients with metastatic NSCLC from eligibility for enrollment in clinical trials driven by the biomarker results.^{30,31} With the expanding list of biomarker-selected treatments, shorter TAT of comprehensive panel-based biomarker testing is urgently needed. The National Comprehensive Cancer Network and CAP/IASLC/AMP guidelines recommend panel-based testing, such as next-generation sequencing (NGS) assays.^{1,2} However, there is an urgent need to determine optimal and timely testing strategies for different practice settings.

Barriers to large-scale adoption of comprehensive biomarker testing include unavailability of expanded panel-based testing, especially in low-resource settings; variability in adequate tissue acquisition; variable reimbursement for NGS-based assays; poor physician awareness of clinically relevant biomarkers; limited physician knowledge or expertise in interpreting results; and, in some cases, physician preference for highly reimbursable infusion therapies.^{17,18,32,33} These result in inconsistent use of comprehensive biomarker testing across oncology practices.

This article highlights the important timepoints in the workflow from tissue procurement to biomarker reporting and their potential impact on TAT, which varies between institutions and practices. We describe preanalytical and analytical factors that can adversely affect TAT and recommend mitigating strategies. We also propose a process map for physicians and key organizational stakeholders to support greater efficiency and new strategies to reduce TAT.

Factors affecting TAT

TAT for comprehensive biomarker testing should be measured from the day tumor tissue becomes available (typically, once the pathologic diagnosis is finalized) to the day the treating physician receives the molecular test results. For patients with a high pretest probability of NSCLC, the *perceived* TAT often starts from when the clinical diagnosis is made, even prior to tissue procurement and a pathology diagnosis. The following *key factors* affect the TAT of lung cancer biomarker tests (Table 1):

Preanalytical factors

- Timeliness of test ordering
- Availability and adequacy of tissue for testing

Analytical factors

- Choice of assay
- Type of testing laboratory and tissue-handling processes

PREANALYTICAL FACTORS

Timeliness of test ordering

Streamlining biomarker test requests

Delay in initiating biomarker testing is one of the most important preanalytical variables. A high level of local interdisciplinary communication and coordination is central to consistent and timely test ordering. There is no consensus for a single subspecialty to act as the responsible physician. The optimal strategies may vary based on expertise and logistics at the particular practice or institution.¹⁶

TABLE 1 Factors affecting turnaround time of biomarker test results and possible strategies to optimize testing.

Factors affecting TAT	Strategy	Challenges and limitations
Timeliness of test ordering	<p>Health care physician orders molecular testing at the time of initial clinical diagnosis</p> <ul style="list-style-type: none"> • Prospective order in anticipation of NSCLC diagnosis • Electronic health record order based on rapid on-site evaluation • Electronic health record order triggered by final pathology diagnosis (e.g. NSCLC) <p>Pathologist-directed reflex testing based on stage and histology</p>	<ul style="list-style-type: none"> • Requires close communication and education of clinicians (physicians [MD] and nurse practitioners [NP]) • Requires standardized protocols for test ordering within electronic health record • May result in inappropriate or redundant testing • May not know the stage or prior testing history • May result in inappropriate or redundant testing • Institutional or state-mandated restrictions
Availability and adequacy of tissue for testing	<ul style="list-style-type: none"> • Ensure adequate tissue acquisition and provide feedback to proceduralists by using rapid on-site evaluation • Create <i>molecular only</i> block by processing core biopsies as separate blocks • Optimize cytology cell-block preparation by minimizing material for smears or liquid-based cytology • Precutting unstained sections to minimize tissue wastage through block refacing • Minimize unnecessary diagnostic immunohistochemistry use • Preselect adequate samples at the time of finalizing pathology report 	<ul style="list-style-type: none"> • Education of proceduralists necessary to ensure adequate collection of tissue • Performance feedback systems require resources • Requires education of pathologists and laboratory personnel for judicious triaging of tumor tissue • Standardized protocols in histology and cytology laboratories needed for optimal processing
Choice of assay	<ul style="list-style-type: none"> • Limiting the use of sequential and single-gene testing to an <i>as-needed</i> basis • Preference for panel-based, comprehensive testing when feasible and appropriate 	<ul style="list-style-type: none"> • Requires standardized workflows for test send-outs • Requires institutional infrastructure and appropriate resources and access to testing • Financial impact analysis (of in-house testing vs. sending out to a reference laboratory) may be necessary
Alternate or complementary testing	<ul style="list-style-type: none"> • Plasma-based genotyping at diagnosis 	<ul style="list-style-type: none"> • Insurance and billing issues • Reduced detection rates in patients with modest tumor burden • May require parallel tissue-based testing
Quality metrics	<ul style="list-style-type: none"> • Developing process improvement tools to identify areas for improvement • Creating testing algorithms for choosing the optimal biomarker testing strategy • Creating dashboards that can provide real-time monitoring of biomarker test ordering and reporting • Monitor time from diagnosis to biomarker test results 	<ul style="list-style-type: none"> • Requires education of clinicians (MD, NP) • Requires institutional commitment and resources to develop monitoring and reporting tools • Requires individual and institutional commitment to continuous quality improvement

Abbreviations: NSCLC, non-small cell lung carcinoma; TAT, turnaround time.

Potential challenges experienced by each subspecialty include the following.

1. *Proceduralists*: Many pulmonologists, interventional radiologists, and thoracic surgeons who perform the initial diagnostic biopsy do not initiate testing because they may not be comfortable with the testing requirements (for instance, the stage of the lung cancer, the histology appropriate for biomarker testing, the specific biomarkers to order, the testing algorithm to follow, the kind of specimen preparation required for testing), interpretation of tests, or responding to patient inquiries about subsequent management.^{33,34} For many nonsurgical proceduralists, there is no ongoing physician-patient relationship, and the ordering of any further testing beyond routine diagnostic pathology may be commonly viewed as the treating physician's responsibility.
2. *Pathologists*: Pathology departments often wait for biomarker test requests from other physicians to initiate testing, partly because of the current stage-limited indication for biomarker testing.¹ Pathologists are often uncertain of the clinical stage, so there has been a reluctance to take on the responsibility of test ordering when testing was mostly restricted to patients with stage IV disease. Waiting for a test order request from another treating physician is also a means of compliance with statutes against self-referral.³⁵
3. *Oncologists*: It often falls on oncologists to order biomarker testing because the results drive their treatment decisions. However, patients often establish care with an oncologist days to weeks after the initial diagnosis of NSCLC. Waiting for the oncologist to order biomarker testing delays the process of informed treatment decision making. Ideally, biomarker results should be available to help formulate a treatment plan when the patient first meets the oncologist. In addition, some oncologists

may be reluctant to order biomarker testing until established care is set up or they meet the patient to ensure appropriate follow-up of test results.

In addition, patients who self-refer to a tertiary care cancer center often do not have a complete set of reports that outline biomarker testing performed at the initial diagnosis. Possible approaches to address these problems are summarized below in Table 1.

Ordering within the electronic health record

Ordering through an electronic health record (EHR) can help prevent delays in biomarker testing. A templated biomarker panel order-set request created using the EHR to be initiated within 1 day of the finalized pathology report can improve efficiency. When the pathology report populates the EHR, an automatic notification can alert the appropriate health care physician to place the order. The order may be as simple as “lung cancer comprehensive biomarker testing,” and appropriate biomarkers can be updated using a dynamic interface based on evolving testing guidelines. Thus the physician can be prompted through the EHR to initiate test orders within 1 day of the finalized pathologic diagnosis.

Although EHR optimization has the potential to improve testing, this approach still relies on various physicians ordering testing promptly. For admitted patients, an EHR prompt or message could be sent to the physician responsible for their care; however, this may be less practical for outpatients. It may be unclear which outpatient clinician (primary care physician, interventional radiologist, pulmonologist, thoracic surgeon, nurse practitioner, or someone else) will be responsible for initiating the test order. However, a triage system compatible with the organization could be established to designate a clinician under appropriate circumstances and then alert the designated clinician to initiate biomarker testing for either inpatients or outpatients. One barrier to timely test ordering for inpatients is the Medicare 14-day rule. This rule precludes independent billing for biomarker testing within 14 days of hospitalization, thus disincentivizing medical systems from providing timely testing for current or recently hospitalized patients. Paradoxically, these patients are even more likely to need timely treatment and thus efficient TAT.³⁶

Another drawback is that automated notification within the EHR to trigger the biomarker order may not be feasible when practices use different EHRs or hospital EHRs without a closed loop. In some cases, the anatomic pathology laboratory information management system may communicate in only one direction with the hospital EHR, allowing reports to cross over but not accepting incoming orders. Thus testing may rely on less fail-safe processes, such as faxed or emailed orders. These factors delay biomarker testing, diminishing access to timely testing and effective targeted therapy.

Reflex biomarker testing

One of the most efficient ways to reduce a delay in ordering is to have reflex testing protocols, in which biomarker testing is initiated automatically based on predetermined criteria and institutional policy.^{20,25,37} Reflex testing will become more feasible as the indications for biomarker testing extend across the clinical stage and may eventually be akin to the American Society of Clinical Oncology/CAP guidelines for breast cancer (i.e., reflex fluorescence in situ hybridization [FISH] testing must be performed when ERBB2 [HER2] IHC results are equivocal).³⁸ Reflex testing could be based on predetermined institutional qualification criteria initiated when a final pathologic diagnosis has been rendered. Several centers have published their experience implementing a reflex testing strategy and have demonstrated improved testing rates and TATs.^{25,39-41}

The potential impact of a reflex strategy on TAT depends on preintervention ordering delays. One center improved the average TAT (from diagnostic pathology report to biomarker report) from 26 days to 15 days in 1 year.²⁵ To avoid regulatory or reimbursement problems, institutions and practices should establish consensus processes, including clear expectations and communication (ideally through electronic orders in the her, as discussed in the previous section), to ensure timely and appropriate testing. Given the complex logistics involved in the timely diagnosis, staging, and treatment of lung cancer, interdisciplinary communication and coordination are crucial. Pathologists should work closely with the key subspecialists involved in diagnosis and treatment, including interventional radiologists, pulmonologists, medical oncologists, and thoracic surgeons, to establish the rules, roles, and relationships to streamline their institution's workflow. Such processes should consider and communicate the appropriate stage of NSCLC for reflex testing. Similarly, proceduralists working within institutions that adopt reflexive testing should be made aware of such testing so that they routinely aim to procure sufficient tissue for comprehensive biomarker testing to avoid unnecessary additional procedures. Institutional policies for reflexive biomarker testing should consider the current level of clinical evidence, insurance policies, billing requirements, other contractual relationships, and available resources. These approaches to streamlining biomarker test requests can enable a more efficient workflow for timely testing and reporting.

Availability and adequacy of tissue for testing

A critical component of optimal biomarker testing is procuring an adequate tumor sample. Inadequate (*quantity not sufficient*) tissue samples lengthen TAT by necessitating a search for more suitable material. Often this means scheduling a repeat biopsy or treatment without information from comprehensive biomarker testing. Adequate samples should be allocated for routine tissue processing during the initial procedure. Although some laboratories may be able to use non-formalin-fixed paraffin-embedded (FFPE) specimens, such as cytology smear slides or liquid-based cytology preparations,⁴²⁻⁴⁵

most commercial laboratories require FFPE blocks with adequate tumor tissue to further process the sample. Given the differences in laboratory practices, assays, and strategies, it is crucial for those performing tissue sampling to know what *adequate tissue* means for their testing laboratory. In addition to the size, quality of tissue, and tumor cellularity needed, the feasibility of biomarker testing also depends on judicious specimen processing to ensure minimal tissue depletion.^{46–48} Some strategies to minimize *quantity not sufficient* results are outlined below and summarized in Table 1.

Rapid on-site evaluation and handling of cytology samples

Rapid on-site evaluation (ROSE) at the tissue acquisition increases the likelihood of specimen adequacy.⁴⁹ CAP guidelines recommend ROSE for all endobronchial ultrasound-guided, bronchoscopic, and transthoracic biopsies to ensure collection of an adequate sample for comprehensive biomarker testing when the clinical suspicion of lung cancer is high.⁵⁰ ROSE on specimens collected by fine-needle aspiration and touch-imprint cytology of core-biopsy samples can also facilitate preordering reflex biomarker testing within the EHR when diagnosing NSCLC. The primary limitation is having the necessary infrastructure, personnel, and resources to support ROSE and the perceived delays in procedure TAT when ROSE is included.⁵¹

With or without ROSE, cytology samples should be prioritized for FFPE cell-block preparation rather than split for cytospin or liquid-based cytology (e.g., ThinPrep) and cell block, especially where biomarker testing is performed from FFPE blocks only. A minimal volume of the aspirate (e.g., a drop of the sample) can be used to prepare a direct smear for adequacy assessment by ROSE. The remaining aspirate and at least three—but preferably five—passes can be directly submitted for preparing the FFPE cell block.⁵⁰ This would improve yield for diagnosis and comprehensive biomarker testing and potentially help practices that do not have adequate resources for ROSE on cytology samples. For patients who present with malignant pleural or pericardial effusions, the cytologic fluid may be the only specimen available for diagnosis and comprehensive biomarker testing. In these situations, CAP guidelines recommend prioritizing the FFPE cell-block preparation in anticipation of biomarker testing.⁵⁰ Incorporating ROSE during specimen collection to ensure adequacy can reduce delays resulting from insufficient biopsies for comprehensive biomarker testing.^{49,52,53}

Optimized handling of small biopsy samples

One option to conserve scanty biopsy specimens is to submit individual cores from different biopsy passes in separate cassettes for processing as individual FFPE tissue blocks.⁵⁴ This allows a tissue block with one core-biopsy sample to be processed with sections for hematoxylin and eosin staining (the first and last cut sections) for diagnosis and provides intervening unstained slides for diagnostic

IHCs. The remaining tissue blocks can be preserved for biomarker studies, with only a superficial section used for hematoxylin and eosin staining to allow an assessment of viable tumor adequacy in each core sample. Although this approach may use additional resources in the histology laboratory, it may ultimately be safer and more cost-effective by reducing the number of inadequate biopsies and the need for repeat procedures.⁵⁴

Minimizing diagnostic immunohistochemistry

IHC for diagnostic markers and subtyping specimens should be informed by the clinical context, following recommendations set forth by the IASLC.⁵⁵ For most purposes, when IHC is needed for the subtyping of NSCLC, TTF1 and p40 are considered sufficient in clinical practice if there are no morphologic features of neuroendocrine differentiation. If neuroendocrine differentiation is present or metastasis from other primary sites is suspected, additional IHC markers may be used for a more definitive diagnosis.⁵⁵ Minimizing the number of diagnostic IHCs performed can conserve tissue for biomarker testing and prevent delays caused by insufficient tissue.

Preidentifying tumor blocks for testing

Pathologists rendering a NSCLC diagnosis should indicate the suitability and availability of tissue blocks for biomarker testing at the time of diagnosis, ideally within the pathology report itself and, at a minimum, within the pathology laboratory information-management system. This can prevent delays by avoiding re-reviewing slides to identify the appropriate block for biomarker testing.

Providing tissue-qualification criteria

Biomarker testing laboratories can provide tissue-qualification criteria based on biomarker assay performance characteristics. This empowers the primary diagnostic pathologist to select an appropriate sample for biomarker testing that the testing laboratory will deem adequate. Key elements for the tissue-qualification criteria would include minimal tissue size (mm²); overall cellularity (especially for cytology specimens); tumor cellularity (percentage of tumor cells in the specimen); qualitative features, including the presence of necrosis and extracellular mucin; and a list of agents that can interfere with molecular testing, such as heavy metal fixatives and harsh acid decalcification.⁵⁶ These steps can identify specimens that may be suboptimal for accurate biomarker characterization. Documentation of these elements, either directly within the pathology report or in the laboratory information-management system, can minimize delays in selecting the best specimen for biomarker studies. Laboratory-specific workflows that adhere to tissue-qualification guidelines can ensure a streamlined process, minimize delays, and reduce TAT.

ANALYTICAL FACTORS

Choice of assay

Multiple methods are available for biomarker testing, including IHC (PD-L1, ALK, ROS1), FISH (ALK, ROS1, RET), single-gene assays (e.g., Sanger sequencing, real-time polymerase chain reaction, fully automated and integrated testing platforms for EGFR, KRAS, BRAF mutations), and multiplex, high-throughput sequencing platforms (e.g., NGS). Testing strategies include sequential assays (either methodology-based or gene-based) or panel-based assays. Multiplexed assays may be small panels (≤ 50 genes) or expanded panels (with several hundred genes that encompass most of the known somatic changes in solid tumors). It is worth noting that there is inherent conflict between comprehensive biomarker testing and the TAT of results. Various forms of noncomprehensive biomarker testing may be more easily performed than comprehensive biomarker testing. We advocate for comprehensive assessment of actionable mutations to direct patient treatment decisions whenever possible.

Sequential testing, in which a biomarker (gene-based) or panel (methodology-based) is tested stepwise before proceeding to other tests, has been judged by many practices to be convenient and cost-effective. Nevertheless, this strategy is falling out of favor because of the expanding number of biomarkers that need to be assessed promptly.⁵⁷ Some sequential testing strategies may be preferred based on their short TAT, making them alluring when there is clinical urgency to start treatment. For instance, IHC-based and FISH-based test results are typically available within a few days, unlike NGS test results, which frequently take 2 or more weeks (Table 2). A sequential testing approach requires individual practice-specific *standard operating procedures* that prioritize one particular gene or assay over another. However, a sequential single-gene testing approach may exhaust the tumor tissue because each time a tissue block is recut for sections for additional testing, the block needs to be refaced, thus losing precious tissue material and raising the risk of tissue exhaustion. One study evaluating the adequacy of small biopsy

specimens for lung cancer biomarker testing demonstrated that only 18% of patients had samples that were adequate for comprehensive testing of all guideline-recommended biomarkers when using a single-gene testing approach.¹⁵ In addition, waiting for each sequential test to be complete before moving to the next can consume valuable time.⁵⁸

The CAP/IASLC/AMP guidelines recommend that laboratories consider cutting multiple additional unstained sections *up-front* for biomarker testing to minimize the need to reface the tissue block and prevent tissue loss during processing in the histology laboratory.²⁹ Furthermore, having sections available to submit for biomarker studies when finalizing the diagnosis can reduce TAT because there is minimal delay in waiting for the block to be recut.

Historically, molecular laboratories have prioritized single-gene assays because billing for these assays and reimbursement for testing are more consistent. Furthermore, many IHC-based assays (PD-L1, ALK, ROS1) are performed in-house and are preferred for this reason. Panel-based testing provides results for all guideline-recommended biomarkers and a more predictable TAT for all, including relatively uncommon mutations. Therefore, the lower cost and relative ease of using single-gene testing may be an attractive approach to some practices (especially when comprehensive NGS assays require integration of laboratory processes with an outside vendor, seem cost-prohibitive, or raise concerns about reimbursement). In reality, primary reliance on a single-gene testing approach poses a higher risk of depleting limited tissue material and failing to provide results for less prevalent biomarkers that are lower on the sequential testing priority list.

When choosing a testing platform, the goal for all laboratories should be to adhere to the CAP/IASLC/AMP guidelines, which recommend a TAT of ≤ 10 working days for tissue-based molecular testing results, while comprehensively testing all guideline-recommended biomarkers for individual patients.² Ultimately, the choice of a testing platform that is optimal for patient care within an individual practice and provides biomarker test results with an acceptable TAT will depend on institutional infrastructure, access to

TABLE 2 The types of biomarker testing available that guide the choice of the testing platform.

	Testing method	Testing biomarkers	Turnaround time	Testing priority
Single-gene assays	Molecular (e.g., Sanger sequencing, real-time PCR)	EGFR, KRAS, BRAF, ERBB2, ALK, ROS1, RET, MET, NTRK1-3	5-10 days	If performed sequentially, increases the overall turnaround time of test results and depletes tissue
	FISH	ALK, ROS1, RET, MET, ^a NTRK1-3	5-7 days	
	IHC	ALK, ROS1, ^b MET, ^c PD-L1	2-5 days	
Panel-based, multiplexed assays	Molecular e.g. NGS	EGFR, KRAS, BRAF, ERBB2, ALK, ROS1, RET, MET, NTRK1-3	≤ 10 days (Lindeman 2013 ²⁹)	Minimizes tissue wastage and optimizes overall turnaround time of test results

Abbreviations: FISH, fluorescence in situ hybridization; IHC, immunohistochemistry; NGS, next-generation sequencing; PCR, polymerase chain reaction; PD-L1, programmed death-ligand 1.

^aFISH can detect MET copy number changes but not MET exon 14-skipping mutations.

^bROS1 fusions detected by IHC require confirmation by molecular testing/FISH.

^cIHC can detect MET amplification but not MET exon 14-skipping mutations.

comprehensive testing resources, and input from physicians involved in clinical decision making.

NGS-based testing requires multiple steps (nucleic acid extraction and purification, library preparation, sequencing, analysis using a bioinformatics pipeline) and typically takes 2 weeks or longer. The TAT can vary significantly within molecular testing platforms based on the method used for sample purification and sequencing. Also, laboratories need to decide on DNA-based versus RNA-based NGS assays for the detection of gene fusions because there are some inherent limitations related to DNA-based, amplicon-mediated testing. This may lead to adding days to the TAT if a sequential NGS strategy (RNA-based testing for gene fusions, if initial DNA-based NGS is negative for actionable targets) is adopted. The emergence of integrated NGS solutions can considerably reduce the TAT to a few days (approximately 3 days), even for large, expanded panels; however, cost and throughput may serve as barriers to widespread implementation of these solutions.^{59,60} However, no test is perfect. Examining a biomarker with multiple modalities (for instance, *ALK* detection by FISH, IHC, or NGS) may increase the likelihood of detecting common alterations, especially for cases in which molecular alterations may be missed because of low specimen tumor content.

Type of testing laboratory and tissue-handling processes

The choice of in-house testing versus send-out testing by a reference laboratory is an institutional decision. It is usually based on multiple factors, including test volume, direct and indirect costs, available resources, infrastructure for test performance, validation and operations, regulations related to testing, such as patents and licenses, and reimbursement. Specific workflows need to be established for material to be sent out for testing or used for in-house assays. If biomarker testing is being sent to different reference laboratories for separate assays (e.g., PD-L1 IHC and NGS for mutations and gene fusions), practices must consider specimen requirements and acceptance criteria for each assay to minimize delays in test reporting, including inability to test because of sample inadequacy. An efficient workflow to handle tissue requests for biomarker testing (molecular, FISH, and IHC) can reduce the time from specimen procurement and diagnosis to tissue submission for biomarker testing, thus improving overall TAT. Rapid molecular tests that provide results within hours to days may be considered for the more frequent gene alterations and expanded NGS assays for comprehensive testing.^{61–63} However, this approach needs to be evaluated within the context of tissue availability, resources, and reimbursement considerations before implementation.

Incorporating the plasma genotyping option

Contemporary management of metastatic NSCLC hinges on comprehensive biomarker testing. Yet, in multiple studies, 20% to

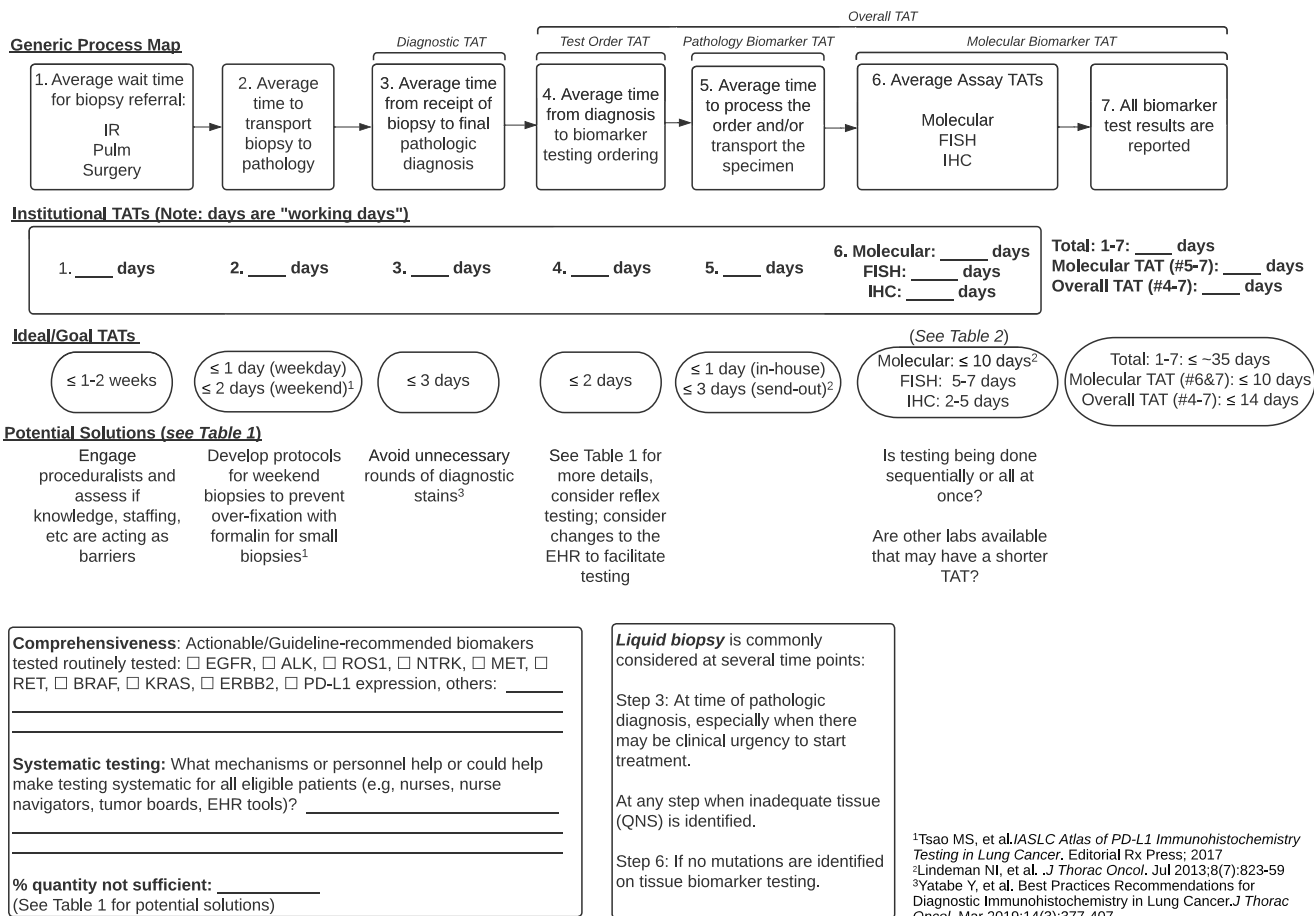
30% of patients with newly diagnosed NSCLC do not have *EGFR* or *ALK* genotyping results, let alone the full range of guideline-recommended biomarkers, before initiating therapy.¹⁵ The MYLUNG (Molecularly Informed Lung Cancer Treatment in a Community Cancer Network) consortium study, which evaluated real-world biomarker testing rates for five biomarkers (*EGFR*, *ALK*, *ROS1*, *BRAF*, and PD-L1), reported that, although most patients with metastatic NSCLC received at least one biomarker test before first-line systemic therapy, <50% received testing for all five biomarkers.⁶⁴ Plasma-based genotyping assays (also frequently referred to as *liquid biopsy*) have a high positive predictive value, may provide actionable results, and expedite clinical decisions if a targetable alteration is detected. However, the sensitivity of plasma-based genotyping assays may be suboptimal in patients with a lower metastatic tumor burden. Additional limitations include lack of assessment of PD-L1 expression to direct immunotherapy use and lower performance to detect certain targetable mutations (e.g., fusions and copy number variants). Nevertheless, these assays are useful when positive but should prompt further tissue testing when negative, and a more comprehensive approach to tumor genotyping is preferred.

Although CAP/IASLC/AMP molecular testing guidelines recommend plasma-based genotyping assays as complementary to (not a replacement for) tissue-based testing,² recent US Food and Drug Administration approvals of commercial assays support plasma-based genotyping for up-front tumor profiling. False-negative results remain a limitation, and practitioners must recognize the ongoing need for tissue-based testing. However, given the pressure for expeditious comprehensive genotyping results, plasma-based genotyping assays with shorter TAT (approximately 5–7 days) and easy access to specimens can be attractive.¹⁵ Therefore, a possible workflow could involve performing plasma-based genotyping on all patients with metastatic NSCLC at the time of their initial oncology appointment, with tissue testing either concurrently or subsequently if the *liquid biopsy* is negative.⁶⁵ This latter strategy can further delay access to actionable biomarker data. Unfortunately, at this time, most third-party payors will not cover both plasma-based genotyping and tumor tissue-based testing for patients with NSCLC, thereby limiting routine implementation of such an approach. This may change in time with the accumulation of evidence.

Assessing quality metrics and tools to incorporate feedback

Performance benchmarking is critical to improving TAT, closely monitoring quality metrics, and using process-improvement tools to identify areas that may benefit from additional resources (Figure 1).^{29,55,66} Testing algorithms and decision aids for choosing the optimal biomarker testing strategy can help reduce the time to test initiation and improve key metrics (Table 3). Developing evidence-based clinical decision tools that associate biomarker test results with specific treatment algorithms, dissemination efforts for educating physicians on testing strategies, and creating tools, such as

ACS NLCRT Biomarker Testing Quality Assessment Tool



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FIGURE 1 Non-small cell lung cancer biomarker testing quality-assessment tool illustrating the workflow for specimen diagnosis and biomarker testing. Included are steps to identifying delays in TAT and solutions that can help improve the process. EHR indicates electronic health record; FISH, fluorescence in situ hybridization; IHC, immunohistochemistry; IR, interventional radiology; ACS NLCRT, American Cancer Society National Lung Cancer Roundtable; TATs, turnaround times; PD-L1, programmed death-ligand 1; Pulm, pulmonary; QNS, quantity not sufficient.^{29,55,65}

TABLE 3 Key quality metrics that can guide quality-improvement efforts.

Proposed quality metric	90% compliance goal
Pathology diagnostic turnaround time (i.e., time from specimen received in pathology to final pathologic diagnosis)	≤3 working days
Biomarker test order turnaround time (i.e., time from final pathologic diagnosis to biomarker test ordered)	≤2 working days
Pathology biomarker turnaround time (i.e., time from final pathologic diagnosis and/or biomarker test ordered to specimen sent to molecular laboratory) for eligible patients with NSCLC	≤3 working days (Lindeman 2013 ²⁹)
Molecular biomarker turnaround time (i.e., time from specimen received in molecular testing laboratory to reporting of all biomarker results) for eligible patients with NSCLC	≤10 working days (Lindeman 2013 ²⁹)
Overall biomarker turnaround time (i.e., time from final pathologic diagnosis rendered to reporting of all biomarker results) for eligible patients with NSCLC	≤14 working days

Abbreviation: NSCLC, non-small cell lung cancer.

dashboards, that can provide real-time aggregate monitoring of biomarker test ordering and reporting may improve testing TAT and decision making.

CONCLUSION

Reducing TAT for comprehensive biomarker testing in NSCLC to fully inform treatment decisions is challenging but necessary. Given the diverse clinical scenarios, practice settings, and logistical barriers, there are different effective approaches to establishing timely access to biomarker information. The goal is to achieve timely, comprehensive biomarker testing for therapeutic decision making, given the rapid expansion of biomarker-driven, personalized NSCLC treatment. Minimizing test ordering delays should be a key institutional, practice, and physician objective. Frequent interdisciplinary communication and collaboration, leveraging tools available in the EHR to aid in ordering and notification of results, and developing standardized protocols for test orders are critical elements to consider.

In summary, reducing TAT needs to be a multidisciplinary effort. It cannot be accomplished unless there is consensus and coordination of standard operating procedures, roles, and relationships among all key specialties within the institution. Ensuring adequate tissue collection, optimizing specimen handling, and processing to maximize tissue preservation for biomarker testing can improve patient access to comprehensive biomarker testing. The transition from sequential, single-gene testing to reflexive, panel-based molecular testing, the complementary use of plasma-based genotyping, and the use of rapid molecular assays can also expand access to timely, comprehensive genotyping. Finally, a robust quality-assurance program is vital to monitor implementation success. Expanding access to timely, comprehensive biomarker testing will remain an iterative process given the rapid evolution in the field. Existing processes must be adaptable to new knowledge and clinical relevance, including evolving eligibility criteria and technological advances that change the optimal workflow. They should incorporate tools for assessing program implementation across the spectrum of health care delivery systems. The ultimate goal is to foster universal implementation of timely, comprehensive biomarker testing to promote universal access to precision cancer medicine for all patients with lung cancer.

AUTHOR CONTRIBUTIONS

Sinchita Roy-Chowdhuri: Conceptualization, formal analysis, project administration, writing—original draft preparation, and review and editing. **Haresh Mani:** Formal analysis, writing—original draft preparation, and review and editing. **Adam H. Fox:** Formal analysis, writing—original draft preparation, and review and editing. **Anne Tsao:** Conceptualization and writing—review and editing. **Lynette M. Sholl:** Conceptualization and writing—review and editing. **Farhood Farjah:** Writing—review and editing. **Bruce E. Johnson:** Writing—review and editing. **Raymond U. Osarogiagbon:** Writing—review and editing. **M. Patricia Rivera:** Writing—review and editing. **Gerard A.**

Silvestri: Writing—review and editing. **Robert A. Smith:** Resources and writing—review and editing. **Ignacio I. Wistuba:** Conceptualization, supervision, and writing—review and editing.

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CONFLICT OF INTEREST STATEMENT

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ORCID

Sinchita Roy-Chowdhuri  <https://orcid.org/0000-0002-9447-7701>

Farhood Farjah  <https://orcid.org/0000-0002-5205-3804>

Robert A. Smith  <https://orcid.org/0000-0003-3344-2238>

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