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REVIEW

# Testing for EGFR Mutations and ALK Rearrangements in Advanced Non-Small-Cell Lung Cancer: Considerations for Countries in Emerging Markets

Mercedes L Dalurzo (D<sup>1,\*</sup> Alejandro Avilés-Salas (D<sup>2,\*</sup> Fernando Augusto Soares (D<sup>3,\*</sup> Yingyong Hou (D<sup>4,\*</sup> Yuan Li (D<sup>5,\*</sup> Anna Stroganova (D<sup>6,\*</sup> Büge Öz (D<sup>7,\*</sup> Arif Abdillah (D<sup>8,\*</sup> Hui Wan<sup>8,\*</sup> Yoon-La Choi (D<sup>9,\*</sup>

<sup>1</sup>Department of Pathology, Hospital Italiano de Buenos Aires, Buenos Aires, Argentina; <sup>2</sup>Department of Pathology, Instituto Nacional de Cancerología, Mexico City, Mexico; <sup>3</sup>Department of Pathology, A.C. Camargo Cancer Center, São Paulo, Brazil; <sup>4</sup>Department of Pathology, Zhongshan Hospital, Fudan University, Shanghai, People's Republic of China; <sup>5</sup>Department of Pathology, Fudan University Shanghai Cancer Center, Shanghai, People's Republic of China; <sup>6</sup>N.N. Blokhin National Medical Research Centre of Oncology, Russian Academy of Medical Sciences, Moscow, Russia; <sup>7</sup>Cerrahpaşa School of Medicine, Istanbul University-Cerrahpaşa, Istanbul, Turkey; <sup>8</sup>Takeda Pharmaceuticals International AG - Singapore Branch, Singapore, Singapore; <sup>9</sup>Department of Pathology and Translational Genomics, Samsung Medical Centre, Sungkyunkwan University School of Medicine, Seoul, South Korea

\*These authors contributed equally to this work

Correspondence: Yoon-La Choi Department of Pathology and Translational Genomics, Samsung Medical Centre, Sungkyunkwan University School of Medicine, 81, Irwon-ro, Gangnam-gu, Seoul, 06351, South Korea Tel +82 2 3410 2797 Fax +82 2 3410 6396 Email ylachoi@skku.edu Abstract: The treatment of patients with advanced non-small-cell lung cancer (NSCLC) in recent years has been increasingly guided by biomarker testing. Testing has centered on driver genetic alterations involving the epidermal growth factor receptor (EGFR) and anaplastic lymphoma kinase (ALK) rearrangements. The presence of these mutations is predictive of response to targeted therapies such as EGFR tyrosine kinase inhibitors (TKIs) and ALK TKIs. However, there are substantial challenges for the implementation of biomarker testing, particularly in emerging countries. Understanding the barriers to testing in NSCLC will be key to improving molecular testing rates worldwide and patient outcomes as a result. In this article, we review EGFR mutations and ALK rearrangements as predictive biomarkers for NSCLC, discuss a selection of appropriate tests and review the literature with respect to the global uptake of EGFR and ALK testing. To help improve testing rates and unify procedures, we review our experiences with biomarker testing in China, South Korea, Russia, Turkey, Brazil, Argentina and Mexico, and propose a set of recommendations that pathologists from emerging countries can apply to assist with the diagnosis of NSCLC. Keywords: non-small-cell lung cancer, EGFR testing, ALK testing, immunohistochemistry, FISH, next-generation sequencing

## Introduction

Lung cancer continues to be the leading cause of cancer mortality worldwide due to late diagnoses and limited treatment interventions.<sup>1–3</sup> In 2020, lung cancer accounted for 11.4% of all newly diagnosed cancers (approximately 2.2 million) and was responsible for an estimated 1.8 million deaths (18% of all cancer-related deaths).<sup>4</sup> Despite ongoing small declines in lung cancer-related mortality in industrialized nations, such as the United States (US) and the United Kingdom (UK), mortality rates continue to rise in emerging nations, including Brazil, Russia, China, South Korea and Turkey.<sup>2</sup> The reasons for these patterns are complex and multifactorial but include cigarette smoking, unequal access to healthcare leading to delayed diagnosis and treatment, environmental contamination and sociocultural barriers.<sup>5</sup>

Non-small-cell lung cancer (NSCLC) accounts for 85% of all lung cancers and is a histologically and genetically heterogeneous disease.<sup>6</sup> NSCLC includes two major types: (1) non-squamous cell carcinoma (non-SCC), including adenocarcinoma

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# Biomarker Testing for NSCLC

Several biomarkers predictive of therapeutic efficacy have emerged for NSCLC, including epidermal growth factor receptor (*EGFR*) mutations, alterations in the anaplastic lymphoma kinase (*ALK*) gene, *ROS1* rearrangements, *BRAF* V600E point mutations and programmed cell death ligand-1 (PD-L1) expression levels.<sup>10</sup> The most established of these are *EGFR* mutations<sup>11</sup> and *ALK* rearrangements.<sup>12,13</sup> This review only focuses on the prognostic role of these two biomarkers in NSCLC and the challenges in global uptake of *EGFR* and *ALK* testing, especially in emerging markets. Other biomarkers are beyond the scope of this review.

In patients who harbor these biomarkers, targeted therapies with EGFR tyrosine kinase inhibitors (TKIs) and ALK inhibitors are now standard treatment, based on dramatic improvements observed in clinical trials.<sup>10,14,15</sup> In recent studies of ALK inhibitors (eg alectinib, brigatinib, ceritinib, and lorlatinib), tumors were demonstrated to respond positively to treatment. These newer ALK inhibitors offer greater potency against resistance mutations to crizotinib and improved central nervous system (CNS) penetration, which is crucial in the treatment of *ALK*-positive brain metastases.<sup>16</sup>

Similarly, first-line therapy with an EGFR TKI (eg erlotinib, gefitinib and afatinib) significantly prolongs progression-free survival (PFS) and is associated with a significantly higher tumor response rate than first-line standard chemotherapy for patients with *EGFR* exon 18 to 21 mutations. In general, patients with activating mutations, such as exon 19 deletions and exon 21 mutations (L858R), are more responsive to TKIs. Conversely, acquired resistance mutations, such as the exon 20 T790M substitution, which is detected using circulating tumor DNA (ctDNA),<sup>17</sup> is associated with poorer response to TKIs and disease progression within the first year of TKI treatment.<sup>18</sup> Osimertinib is a recent EGFR TKI which was developed in response to the resistant exon 20 T790M substitution.<sup>19</sup> Unlike newer EGFR TKIs that target specific mutations, older EGFR TKIs are less effective in treating emerging mutations, such as *EGFR* exon 20 insertions (ex20ins).<sup>20</sup>

Given the potential benefits of targeted therapy, timely and accurate classification of NSCLC subtypes has become fundamental in patients with advanced NSCLC. Indeed, most international guidelines recommend that all patients with advanced-stage non-SCC NSCLC should be tested for both EGFR mutations and ALK rearrangements before initiation of first-line treatment, with a maximum turnaround time (TAT) of 10 working days.<sup>10,14,15</sup> Despite this, diagnostic testing for predictive biomarkers in patients with NSCLC requires substantial resources and effort, and implementation of guideline recommendations in routine practice is not always applicable, especially in emerging countries where local health policies, drug approval, reimbursement issues, logistical constraints and lack of awareness can result in barriers to testing. The objective of this article is to review EGFR mutations and ALK rearrangements as predictive biomarkers for NSCLC diagnosis, discuss a selection of appropriate tests for these biomarkers and identify trends in the uptake of molecular testing for EGFR mutations and ALK rearrangements in both developed and emerging countries. Finally, we review our experiences in China, South Korea, Russia, Turkey, Brazil, Argentina and Mexico, and propose a set of recommendations that pathologists from emerging countries can apply to implement effective biomarker testing to assist with the diagnosis of NSCLC.

# EGFR Mutations and ALK Rearrangements in Advanced NSCLC

*EGFR* mutations and *ALK* rearrangements are involved in NSCLC pathogenesis by stimulation of downstream signal transduction that leads to cell proliferation and inhibition of apoptosis.<sup>21,22</sup> TKIs act by blocking cross-phosphorylation, leading to reduced activity of intracellular signaling pathways and cell proliferation.<sup>22,23</sup> Historically, *EGFR* mutations and *ALK* rearrangements were thought to be mutually exclusive.<sup>24</sup> However, several recent reports have described these events occur concomitantly in 0.1–2.4% of patients with NSCLC.<sup>25–30</sup> Cases with alterations

of two oncogenic drivers remain rare, and the best management approach is still unclear.

#### EGFR Mutations

The most common *EGFR* mutations in patients with NSCLC are deletions in exon 19 (Exon 19del in 45% of all patients with *EGFR* mutations) and a point mutation in exon 21 (L858R in 40% of all patients).<sup>14,31</sup> Because the presence of *EGFR* exon 19 deletions or exon 21 L858R mutations is predictive of treatment benefit from EGFR TKI therapy, these mutations are referred to as sensitizing *EGFR* mutations. Current guidelines suggest that patients without sensitizing *EGFR* mutations should not be treated with EGFR TKIs in any line of therapy.<sup>10,14</sup> Exon 18 and 20 insertion mutations are less common and comprise the remaining 10% of *EGFR* mutatis in NSCLC.<sup>31</sup> The exon 20 T790M point mutation, and most *EGFR* ex20ins mutations, are predictive of treatment resistance to first- and second-generation EGFR TKI therapies.<sup>14,19</sup>

Designed in response to acquired T790M resistance, osimertinib is a third-generation EGFR TKI that irreversibly binds to the tyrosine kinase domain with lower toxicity than second-generation EGFR TKIs.<sup>19</sup> Osimertinib has demonstrated significant activity in salvage treatment of T790M-positive Japanese patients (n=147), with an objective response rate (ORR) and median PFS of 55.6% and 17.2 months, respectively.<sup>32</sup> Using osimertinib to treat patients with *EGFR* ex20ins has yielded inconclusive efficacy results. A retrospective study reported an ORR of 5% and median PFS of 3.6 months,<sup>33</sup> whereas a Phase II study (where osimertinib was administered at double the approved dose) reported an ORR of 24% and median PFS of 9.6 months.<sup>34</sup>

Recently, the FDA granted approval for amivantamab,<sup>35</sup> a novel bispecific *EGFR-MET* antibody, administered via intravenous infusion for the treatment of patients with *EGFR* ex20ins NSCLC whose disease has progressed during or after treatment with platinum-based chemotherapy. Amivantamab showed an ORR of 36% per investigator assessment (IA) and 40% per independent review committee (IRC). Both IA and IRC assessments for median PFS were 8.3 months, median overall survival (OS) was 22.8 months and median duration of response per IRC was 11.1 months.<sup>36,37</sup> Other emerging treatments such as mobocertinib, an oral EGFR TKI with selective activity against *EGFR* ex20ins, demonstrated rapid and durable responses in platinum-pretreated patients with *EGFR* ex20ins metastatic NSCLC. In these patients, an ORR of 28%, median PFS

of 7.3 months, median OS of 24.0 months and median duration of response of 17.5 months, according to IRC assessment, were reported.<sup>38,39</sup>

The overall prevalence of sensitizing EGFR mutations varies significantly according to ethnicity.<sup>11,40–46</sup> Recently, Graham et al determined the frequency of sensitizing EGFR mutations detected in 170 clinical laboratories from 20 countries participating in the College of American Pathologists (CAP) proficiency testing program.<sup>43</sup> The highest activating EGFR mutation frequency was seen in southern Asia (4260/9337; 46%), and the lowest mutation frequencies in South and North America (113/1439; 8% and 7926/86,654; 9%, respectively) among patients tested for EGFR. However, interpretation of survey-led data should be treated with caution, as the information provided is limited to participating centers only.<sup>43</sup> For example, the frequency reported for South America in the study by Graham et al only examined cases in Brazil,43 while another dedicated study of EGFR mutations in 5738 patients with NSCLC in Latin America that included Argentina, Mexico, Colombia, Peru, Panama and Costa Rica, reported a frequency of 26% – a rate between that observed in Asian (40%) and Caucasian populations (15%).47 Similarly, a US-based cancer registry study (2009-2015) reported a higher frequency of EGFR mutations in black (35/98, 35.7%), and non-black patients (63/98, 64.2%) compared with figures observed by Graham et al highlighting the limitations of a small patient population.43,46 Interestingly, Cheng et al reported significantly shorter survival outcomes among black patients presenting with EGFR mutations, with 2year survival rates almost half those of non-black patients (p=0.001). No racial disparity in survival was observed among patients with wild-type EGFR (p=0.774), suggesting the need for improved, tailored management in this patient population.<sup>46</sup>

In general, patients with sensitizing EGFR mutations are more likely to be non-smokers or former light smokers with AdC histology. However, these characteristics alone should not be used for selecting patients for EGFRtesting.<sup>14</sup>

Most patients with *EGFR* sensitizing mutations treated with first- and second-generation TKI therapy experience disease progression after approximately 12 months.<sup>48,49</sup> A variety of mechanisms are involved in acquired resistance to EGFR TKIs; however, an estimated 50–60% of cases can be attributed to the *EGFR* T790M exon 20 substitution mutation.<sup>48,49</sup> Sensitive assays have detected the T790M mutation in patients prior to initiation of a first-generation TKI. This shows T790M could also be a *de novo* mutation and provides one explanation for intrinsic resistance.<sup>50–53</sup> It is critical to detect this mutation in patients who have developed acquired resistance against first- or second-line EGFR TKIs, as third-generation EGFR TKIs can effectively target T790M-positive cancers.<sup>54–58</sup> Observational medical record data from the US (n=308) suggest that 80% of patients receive a first-line TKI-based treatment with or without combination chemotherapy for *EGFR*-positive Stage IV NSCLC. Only 26% of patients received TKI therapy in the second line, whereas over half of patients stopped first-line TKI and presented no subsequent treatment record.<sup>59</sup>

## ALK Rearrangements

In NSCLC, the *ALK* gene rearrangement results in a fusion protein containing a dysregulated, constitutively active *ALK* kinase domain.<sup>24,60–64</sup> Although evidence of *ALK* rearrangement is present in only 2–12% of all NSCLC cases,<sup>42,65–67</sup> approximately 60% of patients will respond to targeted ALK inhibition<sup>17</sup> using therapies such as alectinib, brigatinib, ceritinib, crizotinib and lorlatinib.<sup>68–74</sup> Clinical characteristics associated with the *ALK* gene rearrangement include AdC histology, never-/light-smoking history and younger age.<sup>71,75–77</sup> However, these characteristics alone should not be used to determine the need for *ALK* testing; *ALK* fusion has also been detected in older patients (>70 years of age) with a smoking history and in patients with SCC.<sup>78,79</sup>

# Testing for EGFR Mutations and ALK Rearrangements

There is general consensus among international guidelines (National Comprehensive Cancer Network [NCCN], European Society for Medical Oncology [ESMO], CAP, International Association for the Study of Lung Cancer [IASLC], Association for Molecular Pathology [AMP]) that all patients with advanced non-SCC NSCLC, regardless of clinical characteristics (such as age, race or smoking status, and including some patients with SCC, such as non-smokers or those <40 years of age), should undergo, at a minimum, testing for *EGFR* mutation, *ALK* and *ROS1* rearrangements, *BRAF* mutation and *PD-L1* expression. Moreover, if next-generation sequencing (NGS) is available, additional alterations in genes such as *RET, MET, HER2* and *KRAS* should also be assessed.<sup>10,14,15,80,81</sup>

In practice, *EGFR* and *ALK* testing is usually performed upon request by the medical oncologist. However, several consensus statements and local policies advocate reflex molecular testing (by pathologists) upon diagnosis of non-SCC NSCLC, regardless of clinical stage.<sup>82,83</sup> This policy has been shown to increase the rate of molecular testing by approximately one-third in some settings<sup>84</sup> and reduce the time to initiating treatment.<sup>82,85</sup> In a recent study from Toronto, Canada, Cheema et al compared outcomes during routine and reflex biomarker testing among 306 patients with newly diagnosed NSCLC. Reflex *EGFR/ALK* testing was associated with a significant improvement in time to optimal systemic therapy, as defined by published guidelines (from 36 to 24 days).<sup>82</sup>

# EGFR Testing

Advanced polymerase chain reaction (PCR)-based methods, such as amplified refractory mutation system (ARMS)-PCR, real-time or quantitative PCR (qPCR), reverse transcriptase PCR (RT-PCR), Sanger sequencing (ideally paired with tumor enrichment) and NGS are the most common methodologies for examining *EGFR* mutation status (Tables 1 and 2).<sup>14,17</sup> Guidelines issued by CAP, IASLC and AMP strongly recommend against using total *EGFR* expression by immunohistochemistry (IHC) testing to select patients for EGFR-targeted TKI therapy.<sup>81</sup>

The gold standard for analyzing mutations is direct sequencing. However, direct DNA sequencing without a mutation enrichment step has a lower limit of detection of 10-25% of total DNA, meaning that the use of samples with low tumor cellularity can result in false-negative outcomes.<sup>86</sup> In contrast, qPCR-based methods show high sensitivity, and testing can be performed relatively quickly on small quantities of tissue (eg one day on average for one sample).<sup>81,87</sup> FDA-approved companion diagnostic tests (CDx) include the Roche Cobas® EGFR Mutation test V2 (a real-time PCR test for erlotinib and osimertinib) and the Qiagen therascreen EGFR rotor-gene Q (RGQ) PCR kit (for gefitinib and afatinib) (Table 3).88,89 However, several studies have shown that commercially available PCR kits may not detect 50% or more of the patients with ex20ins NSCLC identified by NGS<sup>89-92</sup> and may only detect known or a limited number of mutations.<sup>93,94</sup> This could mostly be due to the design of the primers in the PCR kits.95 This suggests NGS has greater sensitivity than PCR in detecting EGFR mutations.<sup>96</sup> However, newer PCR kits are designed to overcome this limitation and increase

| Assay  | Advantage  | Disadvantage   |
|--|--|--|
| Sanger sequencing<br>(direct DNA<br>sequencing)        | <ul> <li>Identification of all known and previously unknown mutations<br/>within the studied region</li> </ul>   | <ul> <li>High tumor content required (mutation detected<br/>when allele frequency &gt;25% total DNA [50%<br/>tumor content])</li> <li>Low sensitivity</li> </ul>   |
| Allele-specific real-<br>time PCR (targeted<br>assays) | <ul> <li>Allows rapid multiplex genotyping of specific known hotspot mutations</li> <li>More sensitive than Sanger (requires 5–10% of the starting tumor DNA)</li> </ul>   | <ul> <li>Designed only to detect most frequent mutations</li> <li>Unable to detect mutations different from those included in the assay (low frequency or novel mutations)</li> </ul>  |
| NGS (massive<br>parallel sequencing<br>technology)     | <ul> <li>Detection of multiple genetic alterations (mutations, gene fusions, CNV), allowing the sequencing of large regions of the genome with higher sensitivity</li> <li>Can be performed by FFPE extraction and freshly collected tissue specimens</li> </ul> | <ul> <li>Effective implementation of NGS requires good-<br/>quality DNA and RNA (not always present in<br/>FFPE samples)</li> <li>Validation of panels can be expensive and difficult<br/>for some laboratories in low- or mid-income<br/>countries</li> </ul> |

#### Table I Mutation-Detection Assays (Eg EGFR, BRAF)

Note: Data from Mok et al.<sup>98</sup>

Abbreviations: CNV, copy-number variation; FFPE, formalin-fixed paraffin-embedded; NGS, next-generation sequencing; PCR, polymerase chain reaction; TAT, turnaround time.

#### Table 2 Rearrangement-Detection Assays (Eg ALK, ROS1 Fusion)

| Assay      | Advantage  | Disadvantage   |
|------------|--|--|
| RT-<br>PCR | <ul> <li>High sensitivity</li> <li>High specificity</li> <li>Can be used on mRNA/cDNA to directly detect fusion genes</li> </ul>   | <ul> <li>Not applicable for unknown partners</li> <li>High-quality of RNA is required (difficult to apply in long-term stored tissue and FFPE samples)</li> <li>Multiplexed assays are required to cover the large variety of fusion transcripts<sup>185,186</sup></li> </ul>  |
| FISH       | <ul> <li>Sensitive and specific</li> <li>Detects fusions irrespective of the fusion partner</li> <li>Break Apart assay Vysis CDx was established as a 'gold standard' to detect <i>ALK</i> fusion in NSCLC</li> <li>Allows use of archived FFPE tissue samples and all cytology samples<sup>187–189</sup></li> </ul> | <ul> <li>Unable to identify the specific gene fusion partner</li> <li>Interpretation requires specialized training</li> <li>The 'break apart' can be difficult to identify due to small physical separation of <i>ALK</i> or <i>ROS1</i> fragments</li> <li>Testing is relatively costly and time-consuming</li> <li>Samples may not contain enough assessable cells to be properly interpreted</li> <li>The 15% cutoff and potential false-positive and false-negative signaling profiles may challenge interpretation</li> </ul> |
| IHC        | <ul> <li>Widely available, relatively inexpensive</li> <li>Rapid TAT</li> <li>For ALK rearrangement, IHC is valid with any quantity and percentage of positive tumor cells (useful in biopsy) or cytology samples with a small amount of tumor cells<sup>190,191</sup></li> </ul>                                    | <ul> <li>ROS1-positive IHC requires confirmation by FISH<br/>(≥15% tumor cells)</li> <li>LDT IHC tests require careful validation</li> </ul>   |
| NGS        | <ul> <li>Same advantages as detailed for mutation studies</li> <li>There are some commercial lung cancer-specific fusion panels<sup>153,192</sup></li> </ul>   | <ul><li>Same disadvantages as detailed for mutation studies</li><li>Good quality RNA is required</li></ul>   |

Note: Data from Tsao et al.<sup>105</sup>

Abbreviations: ALK, anaplastic lymphoma kinase; cDNA, circulating DNA; CDx, companion diagnostic test; FFPE, formalin-fixed paraffin-embedded; FISH, fluorescence in situ hybridization; IHC, immunohistochemistry; LDT, laboratory-developed test; mRNA, messenger RNA; NGS, next-generation sequencing; NSCLC, non-small-cell lung cancer; RT PCR, reverse transcriptase polymerase chain reaction; TAT, turnaround time.

| Table 3 US FDA-Approved Drugs and | Companion | Diagnostics |
|-----------------------------------|-----------|-------------|
| for Advanced NSCLC                |           |             |

| Targeted Agent                                     | Companion Diagnostic Test  |
|--|--|
| EGFR   |  |
| Erlotinib<br>Gefitinib<br>Afatinib<br>Osimertinib  | Cobas <sup>®</sup> EGFR Mutation Test v2<br>Therascreen EGFR RGQ PCR Kit<br>Therascreen EGFR RGQ PCR Kit<br>Cobas <sup>®</sup> EGFR Mutation Test v2 |
| ALK  |  |
| Crizotinib<br>Ceritinib<br>Alectinib<br>Lorlatinib | Vysis ALK Break Apart FISH Probe Kit<br>Ventana ALK (D5F3) CDx Assay<br>Ventana ALK (D5F3) CDx Assay<br>Ventana ALK (D5F3) CDx Assay                 |

**Note:** Adapted with permission from Schwartzberg L, Kim ES, Liu D, Schrag D. Precision oncology: who, how, what, when, and when not? Am Soc Clin Oncol Educ Book. 2017;37:160–169<sup>88</sup> with data from these studies.<sup>106,193</sup>

Abbreviations: ALK, anaplastic lymphoma kinase; CDx, companion diagnostic test; EGFR, epidermal growth factor receptor; FDA, Food and Drug Administration; FISH, fluorescence in situ hybridization; NSCLC, non-small-cell lung cancer; PCR, polymerase chain reaction; RGQ, Rotor-Gene Q.

coverage of detection for *EGFR* mutations.<sup>95</sup> For example, one Japanese study demonstrated the Amoy 9-in-1 qPCR test had high concordance with the Oncomine comprehensive assay 3.0 NGS test, with 99% overall agreement when detecting ex20ins.<sup>89</sup> Pathologists should consider the available approaches and the advantages and disadvantages of each method, including analytical sensitivity and TAT. One testing platform is usually sufficient for *EGFR* testing. However, when the sample contains a small amount of DNA, testing for most frequent *EGFR* mutations, such as exon 19 deletions and L858R mutations, should be prioritized.<sup>10,15,97</sup>

In the setting of active disease progression while exposed to targeted therapy, retesting a tumor sample can shed light on the next appropriate therapeutic steps. Liquid biopsy analysis of ctDNA has become the recommended approach for detecting *EGFR* T790M mutations.<sup>17</sup> This technique assesses the release of DNA fragments from tumor cells into the peripheral circulation due to apoptosis and necrosis.<sup>17,98–100</sup> Recent real-world data suggest droplet-digital PCR has high sensitivity and specificity for low-abundance T790M mutation.<sup>18</sup> The NCCN and ESMO guidelines now recommend repeat genomic testing at progression to identify *EGFR*- and *ALK*-resistance targets.<sup>10,14</sup> Newer applications for ctDNA, such as real-time therapy response monitoring, minimal residual disease (MRD) testing, and their use as a predictive biomarkers for immunotherapy, are currently under investigation.<sup>101,102</sup> When there is no

evidence of *EGFR* T790M, testing for alternate mechanisms of resistance (eg *MET* amplification, *ERBB2* amplification) may be used to direct patients to additional therapies.<sup>14</sup>

# ALK Testing

Testing for *ALK* rearrangement should be systematically carried out in all patients with advanced non-SCC NSCLC.<sup>10,15</sup> There are currently three detection methods widely available in clinical practice, including fluorescence in situ hybridization (FISH), IHC and NGS technology. FISH is considered the 'gold standard' and the most widely used assay for *ALK* rearrangement detection.<sup>10,15</sup> The FISH assay (Vysis LSI *ALK* Break Apart Rearrangement Probe Kit; Abbott Molecular) is a US FDA-approved CDx for *ALK* rearrangement detection for crizotinib, which received approval in 2011.<sup>103</sup> The newer ALK inhibitors, alectinib, brigatinib, ceritinib, and lorlatinib are also paired with this diagnostic test. However, in recent years NGS testing has gained increased use for *ALK* rearrangement detection, overtaking FISH testing (46.0% compared with 37.7%) in 2019.<sup>103</sup>

FISH has several limitations, including the need for expensive equipment, combined with technical and interpretative expertise. Lung cancer cells may overlap with normal lung tissue, and confidently distinguishing between the two in a dark field can be a challenge. Furthermore, appropriate long-term storage conditions that control for light, heat and humidity, are necessary to preserve tissue samples and reduce signal decay.<sup>104,105</sup> The type of ALK rearrangement can also affect the outcome of FISH testing, producing either falsenegative or false-positive results. Although the FISH assay has been firmly validated, the potential for false-negatives highlights the importance of combining more than one testing modality to achieve 100% accuracy with different ALK rearrangements in NSCLC.<sup>104</sup> Other limitations of FISH include the requirement of  $\geq$ 50 tumor cells for the determination of ALK status.<sup>104</sup>

IHC testing is routinely used in diagnostic pathology labs for a variety of reasons, including determination of tumor lineage and subtype. Several antibodies are currently available that detect *ALK* protein expression in NSCLC, using formalin-fixed paraffin-embedded (FFPE) tissue, including the mouse monoclonal antibody 5A4 (Abcam) and the rabbit monoclonal antibody D5F3 (Ventana Medical Systems Inc, Tucson, AZ, Cell Signaling Technology). In 2012, the Ventana *ALK* (D5F3) assay was approved in Europe as a CDx test to aid in the identification of *ALK* rearrangement in patients with NSCLC. The assay was approved by the National Medical Products Administration (NMPA, formerly known as the Chinese Food and Drug Administration) in 2013<sup>104</sup> and by the US FDA in 2015.<sup>106</sup>

There are technical challenges in detecting *ALK* rearrangement effectively by IHC in a spectrum of tissue types. Therefore, it is important for positive/negative controls to ensure accurate interpretation of staining intensity without subjectivity.<sup>104</sup> Until recently, there was considerable debate over testing algorithms for *ALK* detection, and previous guidelines recommended that all *ALK*-IHC-positive results should be confirmed by *ALK* FISH.<sup>107–109</sup> However, given the cost-effectiveness and rapid TAT of IHC, combined with the accuracy of the technique when combined with high-performance antibodies, IHC is now considered to be an equivalent alternative to FISH for *ALK* testing in patients with NSCLC.<sup>10,14,15,80,81</sup>

#### Next-Generation Sequencing

In response to the ever-growing list of predictive biomarkers, NGS technology has emerged as a preferred method for comprehensive testing in NSCLC. NGS enables the simultaneous assessment of many DNA or RNA alterations beyond *EGFR* and *ALK* that are rapidly becoming clinically relevant. Targeted testing provides higher coverage of genomic regions of interest to improve detection of relevant alterations and to allow critical molecular information to be available for therapeutic decisions in an adequate time frame. Performed in an optimized way, NGS can result in improved tissue use and efficiency.<sup>110,111</sup> The US FDA recently approved two DNA-based NGS platforms for molecular testing in advanced malignancies: the Memorial Sloan Kettering-Integrated Mutation Profiling of Actionable Cancer Targets and FoundationOne CDx.<sup>112</sup>

DNA is more stable than RNA, and for many clinical purposes, DNA-based sequencing using panels of specific disease-related genes is sufficient. However, at the DNA level, it is not always possible to detect gene-fusion expression, particularly if breakpoints involve long intronic regions that cannot be covered by hybridization-capture probes.<sup>113</sup> To address this issue, targeted RNA-based NGS assays have been developed for gene-fusion detection, and emerging evidence suggests that these platforms can be more sensitive, efficient and functionally definitive, considering that many DNA variants (eg multiple intronic breakpoints) give rise to the same oncogenic transcript.<sup>114–116</sup> One potential drawback to this approach is that it requires good-quality RNA, which is sometimes difficult to obtain

from FFPE specimens.<sup>99</sup> Although RNA-based assays can be based on hybrid-capture or amplicon-based methods, most use the latter.<sup>99</sup>

# International Patterns of Biomarker Testing

To better understand patterns of uptake for biomarker testing in advanced NSCLC, we searched the PubMed and Embase databases for articles published in English with abstracts between 01 January 2000 and 24 May 2021. Articles were selected by combining search results generated with exploded medical subject headings (MeSH) or keywords from the following categories: (1) NSCLC (ie non-small-cell lung cancer, carcinoma, non-small-cell lung, etc), (2) *ALK* (ie anaplastic lymphoma kinase), (3) *EGFR* (ie epidermal growth factor receptor) and (4) testing (ie molecular testing, molecular diagnosis, testing patterns, testing frequency, diagnostic testing, diagnostic patterns, etc). We hand-searched the reference lists of retrieved articles for additional studies. Abstracts only and/or conference proceedings were excluded.

We identified 24 articles incorporating data from 45 countries. Articles are presented in Table 4 according to region (to account for ethnic differences in molecular profiles) and in chronological order (eg oldest to most recent). Among the articles retrieved, 14 were retrospective reviews,<sup>42,117–129</sup> eight were based on data from surveys and/or interviews with healthcare professionals,<sup>130–137</sup> and two were observational studies.<sup>138,139</sup>

Despite general consensus that biomarker testing is necessary for patients with NSCLC, our review demonstrates this is not being translated into clinical practice, given that testing is highly variable. Although *EGFR* testing has been recognized as standard practice since 2011, the implementation of this assessment can still be inconsistent. However, biomarker testing for *EGFR* mutations and *ALK* rearrangements appear to have increased in most regions over time. In the largest real-world assessment of *ALK*-testing patterns in patients with NSCLC from the US, *ALK* testing rates increased over time from 32% in 2011 to 62% in 2016.<sup>119</sup> Similarly, in a study from the Netherlands by Sluga et al the rate of performing molecular diagnostic testing for NSCLC increased from 11% in 2008 to 75% in 2014.<sup>127</sup>

Several international studies have also identified variable rates of testing between different countries. In the PiVOTAL study, a multinational retrospective study

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|--------------------------------|--|--|--------|----------------------------------|---------------------|--|---------------------------------------|--|--|---|
|                                |  |  |        |                                  | Testing F           | Testing Frequency                            |                                       |  |  |   |
| Author                         | Period   | Location                                   | Method | No. Centers/HCPs                 | c                   | Overall, n (%)                               | EGFR, n (%)                           | ALK, n (%)   | Dx Tests   |   |
| North America                  | ca   |  |        |                                  |                     |  |                                       |  |  |   |
| Pan <sup>125</sup>             | 2007–2011  | VSN  | RR     | iKnowMedTMdatabase <sup>a</sup>  | 1168                |  | 128 (11%) 2%<br><2010; 32% in<br>2011 | 28 (2%)  |  |   |
| Enewold <sup>117</sup>         | 2010   | NSA  | RR     | SEER Database <sup>b</sup>       | 1358                |  | 228 (17%) 157<br>(21%) in AdC         |  |  |   |
| MacLean <sup>123</sup>         | 2010–2012  | ASU  | RR     | Humana Research Database         | 2623                | 1579 (61%)                                   |                                       |  |  |   |
| Lim <sup>122</sup>             | 2010-2013  | CAN  | RR     | l center                         | 258°                |  | 150 (58%)                             | 50 (19%)   | IHC/FISH/NGS   |   |
| Shen <sup>126</sup>            | 2013-2014  | ASU  | RR     | Marketscan Database <sup>d</sup> | 5842                | 1039 (18%)                                   | 1039 (18%)                            |  |  |   |
| Gutierrez <sup>I 18</sup>      | 2013-2015  | ASU  | RR     | 15 centers/89 oncol.             | 814                 | 479 (59%)                                    |                                       |  | PCR/FISH/NGS   |   |
| Schink <sup>135</sup>          | 2014   | ASU  | S      | 57 centers                       | NA                  | 57 (100%)                                    | 57 (100%)                             | 57 (100%)  | EGFR – PCR 44%; ALK – FISH 64%   |   |
| lllei <sup>119</sup>           | 2011–2017  | NSA  | RR     | Flatiron Health Database         | 31,483              |  |                                       | 16,726 (53%) 32%<br>2011; 62% 2016                 |  |   |
| South America                  | a  |  |        |                                  |                     |  |                                       |  |  |   |
| Palacio <sup>124</sup>         | 2011–2016  | BRA  | RR     | All public and private settings  | 11,684              |  | 4440 (38%)  3%<br>2011; 42% 2016      |  | SS   |   |
| Europe                         |  |  |        |                                  |                     |  |                                       |  |  | 1 |
| Sluga <sup>127</sup>           | 2008–2014  | NL   | RR     | 4 centers                        | 2206                | 879 (40%) <sup>e</sup> 11%<br>2008; 75% 2014 |                                       |  |  |   |
| Ess <sup>42</sup>              | 2008–2014  | SUI  | RR     | l center                         | 718                 |  | 447 (62%)                             | 265 (37%)  |  |   |
| Don-<br>Carolis <sup>130</sup> | 2014   | UK   | S      | 15 centers                       | 23,131 <sup>f</sup> | 12,140 (53%)                                 | 12,140 (53%) include<br>testing       | 12,140 (53%) includes both ALK and EGFR<br>testing |  |   |
| Gobbini <sup>138</sup>         | 2014–2015  | ш  | S      | 38 centers                       | 1787                | 1388 (78%)                                   | 1353 (76%)                            | 942 (53%)  | EGFR-direct sequencing ALK-IHC and/or<br>break apart FISH; NGS in some centers | 1 |
| Ryska <sup>136</sup>           | 2014–2015  | Central,<br>Eastern<br>Europe <sup>g</sup> | S      | 42 oncol.                        | AN                  |  | 75-100%                               | ALK testing – all<br>except BG                     | EGFR: RT-PCR; ALK-IHC + FISH and/or FISH<br>alone. ISR, NGS                    | 1 |
|                                |  |  |        |                                  |                     |  |                                       |  |  |   |

Table 4 Testing Frequency for EGFR Mutations and/or ALK Rearrangements in Patients with Advanced NSCLC

| Middle East  |  |  |  |   |   |  |   |  |   |
|--|--|--|--|---|---|--|---|--|---|
| Bar <sup>137</sup>   | 2013   | ISR  | S  | 24 oncol.   |   |  | 19 (79.2%)  |  |   |
| Asia-Pacific   |  |  |  |   |   |  |   |  |   |
| McKeage <sup>139</sup>   | 2010–2016  | ZN   | so   | All non-SCC NSCLC from Auckland,<br>Manukau, Wáitemata and Northland  | 3130  |  |   | 407 (13%)  | Vysis break apart FISH  |
| Yatabe <sup>128</sup>  | 2011–2012  | Asia-Pac   | RR   | 40 centers/I I countries <sup>h</sup>   | 22,193  |  | 18% (CHN) – 65%<br>(JPN) <sup>†</sup>   |  | DNA sequencing. 75% – IHC (poorly<br>differentiated samples)  |
| Hotta <sup>131</sup>   | 2012   | JPN  | s  | 871 oncol.  | NA  |  | 775 (89%)   | 525 (60%)  |   |
| lsobe <sup>120</sup>   | 2011–2013  | Ndĺ  | RR   | 5 centers   | 129 <sup>c</sup>  | 105 (81%)  | 105 (81%)   | 25 (19%)   |   |
| Zhou <sup>129</sup>  | 2015-2016  | CHN  | RR   | 12 centers  | 932   |  | 665 (71%)   | 416 (45%)  |   |
| Prabhash <sup>134</sup>  | 2015–2016  | IND  | INTV   | III oncol.  | NA  | 105 (95%)  | 105 (95%)   |  |   |
| International  |  |  |  |   |   |  |   |  |   |
| Lee <sup>121</sup>   | 2011–2013  | INT  | RR   | 78 centers/8 countries  | 1440  | 43% (BRA) – 85%<br>(TVV)   | 41% (DE) – 97%<br>(TW) <sup>i</sup>   | 3% (TW) – 27%<br>(ITL)   |   |
| Peters <sup>133</sup>  | 2014–2016<br>2016–2016   | INT  | 2<br>surveys <sup>k</sup>  | S1: 562 oncol. <sup>1</sup><br>S2: 707 oncol. <sup>m</sup>  | AN  |  | 450 (80%)   |  |   |
| Jankovic <sup>132</sup>  | 2017–2018  | INT  | S  | 36 oncol/18 countries <sup>n</sup>  | AN  | 90% guided by<br>test results  | 85%   | 58%  | PCR 60%; SS 26%; NGS 26%; FISH 26%; IHC<br>16%  |
| Notes: <sup>a</sup> Captu<br>SEER Database<br>NOS or AdC,<br>Singapore, Taiw<br>Brazil (81%). <sup>k</sup> T | res demographic, "Data are repor<br>26 had SCC. "Pai<br>an, Thailand and<br>wo surveys, 18 r | , clinical, laborato<br>rted for patients<br>tients eligible for<br>Vietnam. <sup>1</sup> Data 1<br>months apart. <sup>1</sup> C | ory and treat<br>with non-SC<br>an EGFR dia<br>not provided<br>anada, France | <b>Notes:</b> <sup>a</sup> Captures demographic, clinical, laboratory and treatment data for patients receiving care within US<br>SEER Database. <sup>c</sup> Data are reported for patients with non-SCC only. <sup>4</sup> Marketscan Database contains health<br>NOS or AdC, 26 had SCC. <sup>1</sup> Patients eligible for an <i>EGF</i> diagnostic test. <sup>8</sup> Bulgaria, Croatia, Czech Republi<br>Singapore, Taiwan, Thailand and Vientam. <sup>1</sup> Data not provided by Indonesia, Malysia, Singapore, the Philippin<br>Brazil (81%). <sup>k</sup> Two surveys, 18 months apart. <sup>1</sup> Canada, France, Germany, Italy, Japan, South Korea, Spain, Ta<br>Brazil Korea, Scin, Auroris Erenco, Socia, Auroris Erenco, Spain, Ta | S Oncology<br>insurance c<br>ic, Israel, Slı<br>1es or Vietr<br>1iwan, UK a | 's network of approx<br>claims for 50 million i<br>ovakia, Slovenia, Pola<br>nam. <sup>1</sup> Italy (69%), Spa<br>nad. <sup>1</sup> Survey 2 i,<br>nd USA. <sup>m</sup> Survey 2 i, | imately 1200 commun<br>ndividuals from >100<br>nd, Hungary, Turkey. <sup>†</sup><br>in (56%), Germany (4)<br>noluded physicians fro | ity-based oncologists.<br>large- or medium-size(<br>'China, Hong Kong, In<br>1%), Australia (64%), J<br>m China. "Portugal, Se | Notes: <sup>a</sup> Captures demographic, clinical, laboratory and treatment data for patients receiving care within US Oncology's network of approximately 1200 community-based oncologists. <sup>b</sup> Random selection of patients with NSCLC in the SEER Database. <sup>C</sup> Data are reported for patients with non-SCC only. <sup>d</sup> Marketscan Database contains health insurance claims for 50 million individuals from >100 large- or medium-sized US-based employers. <sup>e</sup> 853 patients had NSCLC-NOS or AdC, 26 had SCC. <sup>*</sup> Patients eligible for an <i>EGFR</i> diagnostic test. <sup>®</sup> Bulgaria, Croatia, Czech Republic, Israel, Slovenia, Poland, Hungary, Turkey. <sup>In</sup> China, Hong Kong, Indonesia, Japan, South Korea, Malaysia, Philippines, Singapore, Taiwan, Thailand and Vietnam. <sup>U</sup> Data not provided by Indonesia, Singapore, the Philippines or Vietnam. <sup>1</sup> Italy (69%), Spain (56%), Germany (41%), Australia (64%), Japan (81%), South Korea (44%), Taiwan (77%) and Brazil (81%). <sup>k</sup> Twon surves, I8 Montenam, Italy, Japan, South Korea (44%), Japan, Uzbekistan, Peru, <sup>1</sup> China, Hong South South Korea (44%), Japan, 10 <sup>3</sup> , South Korea (44%), Paterbaijan, Uzbekistan, Peru, <sup>1</sup> Kuosu surves, I8 Montenam, <sup>1</sup> Italy, Japan, Cuena, Mantan, South Korea (44%), Japan, South Korea, Spain, Juvau, Teorea (66%), Spain, Justo (66%), Germany (41%), Australia (67%), Australia (75, and 176, and 176, and 176, and 176, and 176, and 178, and 176, and 178, and 188, and 178, and 178, and 188, and |

Abbreviations: AdC, adenocarcinoma: ALK, anaplastic lymphoma kinase; BRA, Brazil; CAN, Canada; CHN, China; DE, Germany; Dx, diagnostic; EGFR, epidermal growth factor receptor; FISH, fluorescence in situ hybridization; HCR healthcare professional; IHC, immunohistochemistry; IND, India; INT, interview; ITL, Italy; ISR, Israel; JPN, Japan; NA, not available; NGS, next-generation sequencing; NL, Netherlands; NOS, not otherwise specified; NR, nor responses; NS, Nos and and and the second of the second strates of second strates and the second strates and st Brazil, Kuwait, France, Spain, Austria, Estonia, Belgium, Montenegro, Nigeria, USA and Poland.

conducted in Italy, Spain, Germany, Australia, Japan, South Korea, Taiwan and Brazil, a total of 1440 patients with newly diagnosed advanced NSCLC were enrolled from 78 centers in eight countries.<sup>121</sup> The EGFR testing rate ranged between 43% in Brazil to 85% in Taiwan. Molecular testing rates (non-specific to NGS/PCR) were similar in Spain, South Korea and Japan (76%, 76% and 74%, respectively).<sup>121</sup> In the same study, the proportion of patients with ALK testing ranged from 41% in Germany to 97% in Taiwan.<sup>121</sup> In a more recent survey of oncologists from multiple developed countries, the proportion of physicians who requested EGFR testing before first-line therapy was 80%.<sup>133</sup> In Asia, EGFR mutation testing was ordered before first-line therapy in 92% and 84% of patients in 2015 and 2016, respectively. In Europe, rates of EGFR testing for the same periods were 77% and 81%, whereas in North America, 76% of patients with advanced NSCLC were assessed for EGFR mutations in 2015 compared with 77% in 2016.<sup>133</sup> A global survey by the IASLC of their members and allied healthcare professionals (n=2537) across 102 countries suggested that fewer than 50% of patients received molecular testing.<sup>140</sup> Many respondents expressed dissatisfaction with the current state of molecular testing in their country, including 41% of those performing and interpreting assays, with identified issues including difficulties in understanding results (37%) and the quality of the samples (23% reported a >10% rejection rate). Despite concerns regarding the quality of testing, 47% of respondents involved in performing and interpreting assays stated there was no policy or strategy to improve quality in their country.<sup>140</sup>

Several studies reported survival differences between cohorts who were tested for molecular biomarkers and, if appropriate, received targeted therapy, versus those who never received a molecular test. In a study from Switzerland involving 718 patients, OS was 10 months (interquartile range [IQR] 4-23 months) for tested patients compared with 3 months (IQR 1-8 months) in patients who were not tested.<sup>42</sup> Similarly, in the PiVOTAL study, patients who were tested had longer OS than patients who were not tested: the median OS from start of first-line therapy ranged from 10.0 (Japan) to 26.7 (Taiwan) months for patients who were tested, and 7.6 (Australia/Brazil) to 19.3 (Taiwan) months for those who were not.<sup>121</sup> Additional studies have found that the probability of molecular testing varies according to practice (eg private versus public),<sup>124</sup> de novo versus subsequent-line therapy,<sup>141</sup> the degree of life expectancy (<1 month versus  $\geq 1$  month),<sup>42</sup> patient age,<sup>126</sup> rural versus urban practice,<sup>142</sup> and the presence of  $\ge 2$  comorbidities.<sup>42</sup>

Direct comparisons of testing uptake between regions should be treated with caution. The rate of biomarker testing uptake depends on many factors, not least of which is the availability of targeted therapies. For example, in the PiVOTAL study, Taiwan had the highest rate of biomarker testing (85%), driven largely by the reimbursement of gefitinib. However, Brazil had the lowest testing rate (43%), mainly because the necessary assays were not covered by the public healthcare system during the study period.<sup>121</sup> Other possible explanations for the different rates of uptake may relate to higher detection rates for EGFR mutations in certain populations (eg Asia). Because many clinical trials demonstrating the superiority of EGFR TKIs over cytotoxic chemotherapy were conducted in Asia, it is reasonable to expect that the experience gained by clinicians and pathologists in those countries has led to more efficient diagnostic services and an increase in the likelihood of testing and treating patients with NSCLC in accordance with guidelines.<sup>133</sup>

# **Barriers to Biomarker Testing**

Barriers to the uptake of biomarker testing are multifactorial and often country-specific. However, access to targeted therapies and absence of reimbursement for testing and/or availability of testing facilities are the main barriers cited in the literature (Table 5);<sup>120,121,132,136,138,139,143,144</sup> assav quality and standards, awareness, and TAT have also been highlighted as additional hurdles to testing.<sup>140</sup> As expected, testing for a specific driver mutation tends to be associated with access to the associated drug,<sup>121,143</sup> and in general, targeted treatments are more accessible for clinicians in the US and Europe than for their colleagues Africa, non-European countries and in South America.<sup>132,144</sup> On a similar note, limited and/or absent

Table 5 Potential Barriers to Biomarker Testing

| Access to targeted therapies <sup>118,135,137–139,143</sup>           |
|---|
| Reimbursement for testing <sup>133,138</sup>                          |
| Tissue sample quality <sup>98,143</sup>                               |
| TAT for test results <sup>42,120,131</sup>                            |
| Coordination among multiple specialist groups <sup>126</sup>          |
| Accurate interpretation of results/physician education <sup>118</sup> |
| Abbreviation: TAT, turnaround time.                                   |

reimbursement is a significant barrier to biomarker testing and a disincentive to reflex testing.<sup>132,136,143</sup> In an international survey of medical oncologists, 38% of respondents noted that biomarker tests were paid for by patients, whereas in 29% and 23% of cases, the expenses were covered by insurance or hospital funds, respectively.<sup>132</sup> Another study of biomarker testing practices in Eastern European countries found that pharmaceutical sponsorship was necessary in order to subsidize the cost of testing in Hungary, Poland and Slovenia. Furthermore, the study reported that in Bulgaria and Croatia, pharmaceutical sponsorship was the only source of financial support for testing.<sup>136</sup> In these situations, creative discussions with all interested parties are critical to finding new solutions. One potential strategy is the development of policies to promote access to testing by establishing links between health systems, and science and technology offices with active participation from the pharmaceutical industry.<sup>144</sup>

Use of validated models for estimating *EGFR* mutation status has been proposed for patients with non-SCC.<sup>145</sup> Based on retrospective patient cohort data in New Zealand, logistical regression modeling of recognized risk factors (including sex, ethnicity, and smoking status) has enabled predictive mutation modeling that may assist clinical decision-making in patients where tissue-based mutation testing is difficult, or as a supplement to mutation testing.<sup>145</sup>

Nonetheless, obtaining adequate tissue for diagnosis, tissue subtyping and molecular profiling are imperative for treatment planning in patients with advanced NSCLC. However, tissue samples from biopsies are often insufficient or inadequate for biomarker testing,<sup>86,122,133,136,146</sup> necessitating repeat biopsies and delays in treatment.<sup>100</sup> Optimizing tissue handling after biopsies are obtained to maximize available material for molecular studies is essential, and standardized algorithms for diagnostic procedures should be defined in routine practice. These should involve reflex testing for biomarker testing, which will shorten TAT and preserve tissue.<sup>135,147</sup> Quality control is also essential to ensure internal reproducibility and the validity of biomarker results. All laboratories should be certified and participate regularly in established qualitycontrol programs.147

As mentioned, reflex testing can reduce the time to initiating targeted treatment.<sup>82,85</sup> Indeed, the reliance solely on clinician judgment decreases the likelihood of full adherence to testing guidelines.<sup>118</sup> For example, in one study, only 21% of patients had biomarker test results

available at the time of the oncology consultation, leading to significant delays in treatment initiation.<sup>122</sup> Moreover, 19% of patients (eligible for targeted therapy) started firstline chemotherapy before biomarker results became available. Avoiding the need for clinicians to specify each biomarker test, through a testing policy that groups tests as a set of common requests, could help ensure testing occurs consistently and rapidly.<sup>122</sup>

Poor cooperation between pathology and molecular testing laboratories has been cited as a potential barrier to biomarker testing.<sup>128</sup> With rapid progress in the molecular profiling of NSCLC and its increasing complexity, collaboration and frequent communication between clinicians obtaining tumor samples, oncologists and pathologists (ie a multidisciplinary team) are essential to successful and timely biomarker testing. Although the ideal approach will vary by country and region, the multidisciplinary team should endeavor to create a center-specific diagnostic and therapeutic plan.<sup>148</sup> The pathologist's input is fundamental to ensure that the appropriate collection method will be used and that sufficient tissue will be collected to allow for morphological, IHC and molecular studies to be conducted.<sup>6</sup> Collectively, the multidisciplinary team should provide individual centers with "stewardship" regarding tissue collection and biomarker testing.<sup>149</sup>

Although many barriers to successful biomarker testing exist at institutional and regional levels, initiatives are often necessary at the individual level. Knowledge translation in this area has demonstrated significant improvements in specialist understanding about tissue sampling, molecular testing and treatment in lung cancer.<sup>150</sup> Furthermore, education of clinicians involved in obtaining diagnostic tumor specimens and reporting pathologists can increase the likelihood of reporting biomarker results by the time of initial oncology assessment.<sup>122</sup> Wherever possible, approaching and educating individuals involved in lung cancer diagnosis and molecular testing, including interventional radiologists, respiratory physicians, pathology technicians and pathologists, should be considered in the context of increasing biomarker testing rates.<sup>110</sup>

# Testing for EGFR Mutations and ALK Rearrangements: Experiences from South Korea, Russia, Turkey, Brazil, Mexico, Argentina and China

Lack of medical and clinical infrastructure complicates the ability to collect country-specific statistics on incidence

and prevalence of NSCLC, including standard practices and outcomes. However, the paucity of available data emphasizes the clear need to develop evidence and correct the "unbalance" of country-level information.

# South Korea

Biomarker testing for EGFR mutations is considered standard practice in South Korea, provided tissue samples are adequately collected. Both EGFR testing and EGFR TKIs are subsidized under the national reimbursement policy. Two real-time PCR-based assays, Cobas® EGFR Mutation Test v2 and PANAMutyper<sup>™</sup> EGFR, have been approved by the Ministry of Food and Drug Safety (formerly known as the Korea FDA). The assays are also approved for ctDNA-based EGFR testing, and their use is evenly distributed between laboratories. Recently, a droplet digital PCR-based assay, the GenesWell<sup>™</sup> ddEGFR Mutation Test, has been approved by the Ministry of Food and Drug Safety. All three methods are reimbursed in South Korea. Regarding the detection of ALK rearrangements in patients with NSCLC, the Ventana ALK (D5F3) assay was approved as a CDx by the Ministry of Food and Drug Safety in 2018. Most laboratories in South Korea have subsequently changed their method for ALK testing from FISH to Ventana ALK CDx D5F3. Both ALK testing and ALK inhibitors are subsidized under the national reimbursement policy.

## Russia

Improving the diagnosis of malignant neoplasms, particularly in patients with NSCLC, is a priority area for the modernization of cancer care in the Russian Federation. Clinical guidelines issued by the Russian Federation Ministry of Health advocate testing for mutations in the EGFR gene and ALK and ROS1 translocations, enabling the administration of effective targeted therapy. However, in practice, molecular testing faces several challenges. First, testing requires highly qualified and experienced technicians. Four large laboratories based at national medical research centers are currently responsible for molecular testing, together with 30 reference laboratories supported by the Russian Society of Clinical Oncology.<sup>151</sup> A second limitation is the transportation of samples from remote regions of the Russian Federation to suitably qualified laboratories. A federal project, titled "Fight against oncological diseases," is currently working to increase the number of reference laboratories, replace obsolete equipment and establish new laboratories in regional centers.

Biomarker testing for *EGFR* mutations is predominantly conducted using the Cobas<sup>®</sup> EGFR Mutation Test v2 or Sanger sequencing.<sup>152</sup> NGS is not used for routine testing due to its high cost. In most centers, IHC is used for the detection of *ALK* rearrangements. However, the presence of artifacts, and 5–10% of doubtful cases, requires the use of alternative methods (eg FISH) for confirmation.<sup>146</sup> At the time of writing, PCR-based methods for the identification of *ALK* translocations have not been registered in the Russian Federation. The methods described above for the detection of *EGFR* mutations and *ALK* rearrangements are funded by the Obligatory Medical Insurance Fund.

# Turkey

In Turkey, routine qPCR assessment of *EGFR* mutation status is conducted by most centers and is reimbursed by government-held healthcare insurance. Despite this, an insufficient amount of tumor cells in the biopsy specimen and poor tissue quality remain significant issues in approximately 5–25% of cases.<sup>136</sup> FISH assessment of *ALK* rearrangements is reimbursed for all patients with NSCLC.<sup>136</sup> IHC is commonly used as a pre-screening method by most centers, with subsequent confirmation by FISH analysis.<sup>136</sup> RT-PCR or NGS methods are used in cases where no signal can be obtained with FISH, or where there is discordance between IHC and FISH results.<sup>153</sup>

In *ALK*+ NSCLC patients treated with lorlatinib, oneyear OS rate was estimated at 65%. In response-evaluable patients (n=55), the ORR and disease control rate were 68.6% and 87.0%, respectively; *ALK*+ patient responses were 69.6% and 87.0%, respectively.<sup>154</sup> Patients receiving erlotinib therapy had significantly improved OS rates compared with patients who received non-TKI treatments (288 versus 119 weeks; p=0.004), whereas PFS rates were not significantly different in patients who did and did not receive erlotinib (32±5 versus 33±3 weeks; p=0.755). Patients expressing both *EGFR* and *KRAS* mutations reported the lowest OS rate. Erlotinib therapy was associated with increased survival in these patients.<sup>155</sup>

The overall frequency of all *EGFR* mutations in Turkish patients with NSCLC is estimated at over 16%, <sup>156,157</sup> with 32% and 20% of cases expressing exon 19 and 21 mutations, respectively.<sup>158</sup> Frequency estimates were significantly higher in female patients and non-smokers.<sup>159</sup> Following lung surgery, *EGFR* driver mutations have been reported in

up to 40% of patients.<sup>160</sup> Data from Turkish patients with Stage III NSCLC suggest neoadjuvant chemotherapy with three agents prior to surgery is associated with outcome benefits.<sup>161</sup>

## Brazil

Although no formal surveys exist, an estimated 70% of all AdC NSCLC cases in Brazil undergo molecular testing.<sup>124,162,163</sup> There are no national outcome or survival statistics, as local data collection is variable or simply not performed across several states. In addition, testing practices vary between centers: several laboratories have adopted reflex testing policies, whereas others approach molecular testing as an on-demand service. Similarly, some laboratories prefer NGS technologies, 162-164 sequencing 14-20 genes at one time, 124,162,165 whereas others have adopted strategies to improve TAT by using RT-PCR assays for EGFR and BRAF mutations, 162,164-166 for ALK, IHC/FISH ROS1 and and PD-L1 rearrangements.<sup>163,164,167</sup>

In the absence of government reimbursement for molecular testing, several pharmaceutical companies have recently established a national "Lung Mapping" consortium. The consortium provides oncologists with access to a subsidized NGS-based test using the FoundationOne<sup>®</sup> platform and has considerably improved access to testing in several Brazilian centers. This ongoing project has yet to report national detection rates of important driver genes.

## Mexico

In Mexico, lung cancer accounts for 10% of all cancerrelated mortality,<sup>168</sup> and only 56.5% of NSCLC cases have a history of tobacco smoking, which suggests that other environmental factors such as hydrocarbons, metals, air pollution, and wood-smoke exposure could have a greater impact in the development of lung cancer.<sup>169,170</sup> The prevalence of *EGFR* mutations and *ALK* rearrangements among patients with NSCLC has been estimated at 34.3% and 7.6%, respectively.<sup>47,171</sup>

Approximately 90% of *EGFR* mutations are exon 19 deletions and L858R mutations in exon 21, although 20.5% of patients express rare mutations in exons 18-21.<sup>172</sup> The most frequent *EGFR* mutations in patients from Mexico are Q787 (15.6%), exon 19 deletions (11.1%), L858R mutation in exon 21 (7.8%), and T790M mutation in exon 20 (1.1%). Other mutations have been identified, such as in *KRAS, MET* and *PDGFRA* (20%), *HNF1A* (14.4%), *APC* (12.2%), *HER2* (11.1%), and *MSH6* 

(10%), as well as alterations of lower frequency in *PIK3CA, GUSB, ALK* rearrangements, *KSR1, KIT, STK11, FLT3, ERBB4, VHL, NOTCH1, GNAS, FGFR3, CDH1, BRAF, ABL1* and *RB1.*<sup>173</sup> In addition, the prevalence of *ALK* rearranged NSCLC is estimated at 7.6% in Mexico. Although there is a country and continental variability of *ALK* rearrangement frequency, the overall incidence of *ALK* rearranged NSCLC in Latin America does not differ from the rest of the word.<sup>171</sup>

However, the uptake of molecular testing as part of the standard diagnostic pathway in NSCLC remains variable. Access to EGFR TKIs is restricted due to cost, and in turn, molecular testing for patients with NSCLC is not reimbursed by the healthcare system. Apart from several large diagnostic centers, such as the Instituto Nacional de Cancerología, very few laboratories have the financial means or adequate infrastructure to undertake testing on a regular basis. Current testing methods available in Mexico include FISH, RT-PCR and IHC. Unfortunately, the FISH test can be expensive, especially for low-income countries, requires experience for accurate interpretation, and does not identify specific fusion transcript variants. In some cases, pharmaceutical companies with novel molecular therapies will pay for the cost of genomic testing as a means of recruiting patients to clinical trials. Although these settings may help a subset of eligible patients in the short term, they benefit only a restricted number of patients and come with substantial burdens.

## Argentina

Access to, and reimbursement of, molecular testing for patients with NSCLC remains a major issue in Argentina. Approximately 90% of all testing is concentrated in a handful of private laboratories and large academic centers in major cities. In general, molecular tests are ordered by the treating oncologists; testing following surgical resection is very infrequent (approximately 1% of all samples). However, reflex testing protocols are possible in large academic centers with dedicated pathology services. There are no comprehensive studies on patient access to testing. In larger centers, an estimated 60-80% of all patients with AdC NSCLC undergo molecular testing, whereas the average TAT for EGFR/BRAF and ALK/ ROS1 tests is 5-7 days and 1-2 days, respectively. Depending on the type of test and the platform used, approximately 5-10% of cases are rejected due to inadequate samples.

Molecular testing for NSCLC follows the IASLC and NCCN guidelines, whereby EGFR, BRAF, ALK and ROS1 are routinely assessed. Of all EGFR mutations, uncommon EGFR mutations (excluding L858R and ex19del) are detected in 10-20% of cases.<sup>174</sup> Cobas® EGFR Mutation Test v2 or Therascreen<sup>®</sup> gPCR is commonly used to assess EGFR and BRAF mutations. IHC with the D5F3 Ventana IHC kit is used for the assessment of ALK, whereas ROSI status is determined using the D4D6 signaling antibody. ROS1 IHC-positive cases are confirmed by FISH. NGS, such as the Oncomine Focus Assay for solid tumors, is available in a few laboratories. In the absence of reimbursement for clinical application, occasionally patients pay privately for the assay; however, the technique is more frequently used for research-based activities. Liquid biopsy is also available and frequently used in patients with resistance to EGFR TKIs. In some instances, it is ordered for firstline testing when the tissue sample is insufficient.

Analysis of clinical and pathologic data from two academic centers in Buenos Aires suggests that *KRAS* mutations occur in approximately 23% of patients with NSCLC, a higher frequency than that reported across Latin America.<sup>175</sup> There was a higher proportion of male patients (65%) and smoking history (94%); mean patient age was 66 years (IQR: 61–72.5 years). In total, 94% of tumor histology samples were identified as adenocarcinoma. Of patients with metastatic disease, 90% received treatment. First-line treatment included chemotherapy (94%) or immunotherapy (6%). Subsequently, half of all patients received second-line treatment, of which 75% received immunotherapy. With a follow-up of 38 months, median OS of patients with metastatic disease was 14.2 months (95% confidence interval [CI], 7.7–30.3).<sup>175</sup>

When considering uncommon EGFR mutations (other than L858R and exon 19 deletion), distribution of tumors comprised exon 18 G719X (46.7%), exon 21 L861Q (24.4%), exon 20 T790M (20%), exon 20 S768I (11.1%), exon 20 insertion (4.4%),and exon 19 pLys745 Ala750del (2.2%).<sup>174</sup> Among patients included in the database review, the most frequent EGFR TKI regimen received was afatinib, followed by erlotinib/gefitinib, and osimertinib. Overall, 22% of tumors were considered complex, defined as  $\geq 2$  coexisting and distinct EGFR mutations, and were associated with a significantly better response to first-line EGFR TKI (ORR 90%) than those with single, non-resistant uncommon EGFR mutations (ORR 52%) (p=0.06). Similarly, patients with complex mutations showed a better OS (median not reached versus 20.3 months [95% CI, 8.7–31.9], respectively; p=0.04).<sup>174</sup>

Private and academic laboratories have different reimbursement systems. Approximately 90% of all molecular tests conducted by private laboratories are supported by the pharmaceutical industry. In academic hospitals, pharmaceutical support is more variable and often confined to the largest centers in Buenos Aires. In those institutions, approximately 50% of molecular tests receive pharmaceutical support, with the remaining tests supported by health insurance and/or privately paying patients. To date, establishing laboratories with the expertise to undertake molecular testing in regional healthcare facilities has been hampered by the absence of a clear and dedicated reimbursement system.

#### China

The National Medical Products Administration of China has approved genetic testing platforms for *EGFR* mutations, including PCR-based methods, Sanger sequencing, Luminex liquid chip and NGS. In China, not all hospitals have the ability or equipment to perform genetic testing and analysis, and so the frequency of *EGFR* testing for NSCLC varies between regions. In Northern China, Cheng et al reported a screening rate of 42.5%.<sup>176</sup> PCR-based methods, such as ARMS (72.1%), Sanger sequencing (5.36%) and Luminex liquid chip (5.10%), were the most frequently used platforms.<sup>176</sup> The median time from tumor diagnosis to *EGFR* or *ALK* status confirmation was 7 and 5 days, respectively.<sup>177</sup>

The efficacy of targeted therapies in NSCLC has been analyzed using a network meta-analysis of 128 clinical trials with 39,501 participants across 14 therapeutic groups. Compared with chemotherapy, ORR was significantly improved for afatinib, alectinib and crizotinib. Cabozantinib and alectinib showed the highest probability for first-line treatment ranking in ORR (62.5%).<sup>178</sup> Targeted therapies have varying effects on OS in NSCLC patients with different gene mutations.<sup>179–181</sup> For example, Yang et al found that OS was increased in patients with EGFR exon 19 deletion compared with the exon 21 L858R point mutation (92 versus 65 months; p<0.001).<sup>180</sup> Compared with chemotherapy, ALK-targeted treatment was associated with a significantly higher PFS (hazard ratio [HR]=0.48; 95% CI, 0.42-0.55), but not significantly higher OS (HR=0.88; 95% CI, 0.72-1.07) in five eligible studies (n=1404).<sup>181</sup>

Several centers in China have adopted reflex testing policies to help streamline the time between pathological diagnosis of NSCLC and identification of molecular markers. Once a sample is diagnosed as NSCLC, a second sample is prepared, and the process of DNA extraction/ molecular identification initiated. Using this process, the molecular report can be sent out at the same time as the routine pathology report. The typical driver genes detected include receptor tyrosine kinases (*RTKs*), *EGFR*, *HER2*, *DDR2* (mutation), *ALK*, *ROS1*, *RET* (gene rearrangement), *MET*, *FGFR1*, *PDGFRA* (gene amplification), *KRAS*, *BRAE*, *PIK3CA*, and *PTEN*.<sup>182</sup>

In the case of lung biopsy specimens, unstained slices or wax rolls are pre-cut at the same time as the pathology sample. When staining or IHC results are indicative of NSCLC, the unstained slices or wax rolls are used for DNA extraction/molecular detection, and the molecular pathology report is available 1–2 days after routine diagnosis. Recent analyses of neoadjuvant targeted therapy suggest significant improvements in radiographic response rates compared with conventional chemotherapy regimens, resulting in longer disease-free survival in real-world settings.<sup>183,184</sup>

All countries indicated above are based on the experiences and perspectives of the authors in their respective countries.

# Conclusion

In summary, the selection of patients with advanced NSCLC based on their EGFR and ALK status is vital, on account of the high response rates observed with the EGFR- and ALK-targeted agents. Despite this, the uptake of biomarker testing varies substantially between countries, and the translation of guideline recommendations into clinical practice remains challenging. Inequitable access to targeted therapies and the absence of reimbursement for biomarker testing are commonly cited as barriers to uptake. Tissue sample quality, delayed TAT and the accurate interpretation of test results create additional barriers. Strategies to address these issues will necessarily be context- and country-dependent but could include the development of multidisciplinary tumor boards to ensure tissue and testing stewardship, standard operating procedures (SOPs) for routine biomarker screening in newly diagnosed patients and further education of specialists who obtain diagnostic cancer specimens.

# **Clinical Practice Points**

Clinical Governance

- Close working relationships between physicians obtaining tumor samples, oncologists and pathologists are essential, and where possible, multidisciplinary tumor boards should be considered to optimize the diagnosis and treatment of lung cancer.
- Center-specific SOPs should be established as part of routine clinical practice to streamline the NSCLC diagnostic pathway.

Infrastructure and Quality Control

- From a laboratory perspective, adequate infrastructure and staffing are necessary to facilitate rapid TAT for biomarker testing.
- Quality control is essential to ensure consistent and reliable diagnostic results. All laboratories that undertake biomarker testing should participate in external quality assessment programs.

#### Education

• Guidance and education of physicians involved in diagnostic tumor specimens and reporting can increase the likelihood of reporting biomarker test results by the time of the initial oncology consultation.

#### Biomarker Testing

- Biomarker testing is necessary for determining the optimal treatment of patients newly diagnosed with NSCLC. Biomarker testing for *EGFR* mutation, *ALK* and *ROS1* rearrangements, *BRAF* mutation and *PD-L1* should be initiated as soon as a pathological diagnosis on non-SCC NSCLC is confirmed (or SCC NSCLC, in selected cases).
- If NGS is available, alterations in genes such as *RET*, *MET*, *HER2* and *KRAS* should also be assessed.
- Reflex biomarker testing at the level of the pathologist should be part of SOPs for confirmed non-SCC NSCLC. This bypasses the time delay for oncology consultation and allows prioritization of sample processing for biomarker testing.
- Timely feedback from pathologists to clinicians acquiring lung cancer diagnostic samples regarding sample adequacy is important.

- Sanger sequencing, pyrosequencing, real-time PCR, and NGS are recommended for *EGFR* testing, and validated tests, including FISH and IHC, may be used for *ALK* testing. NGS can result in a dramatic reduction in cost and should be considered as a means of addressing the financial burden associated with biomarker testing.
- Repeat biomarker testing at the time of disease progression is recommended to identify *EGFR*- and *ALK*resistance targets.

#### Acknowledgments

Medical writing support and editorial assistance were provided by Brad Dalton, MSc, on behalf of Synergy Vision Ltd, London, according to Good Publication Practice guidelines, and were funded by Takeda Pharmaceuticals International AG – Singapore Branch.

#### Disclosure

MD reports personal fees, non-financial support for speaker honoraria and/or congress travel assistance from Pfizer, Takeda, and Roche, outside the submitted work. AA and HW are employees of Takeda Pharmaceuticals International AG – Singapore Branch. The authors report no other conflicts of interest in this work.

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