Title: Molecular Analysis for Targeted Therapy of Non-Small-Cell Lung Cancer

<table>
<thead>
<tr>
<th>Populations</th>
<th>Interventions</th>
<th>Comparators</th>
<th>Outcomes</th>
</tr>
</thead>
</table>
| Individuals:  
- With advanced-stage non-small-cell lung cancer who are being considered for targeted therapy | Interventions of interest are:  
- Testing for *EGFR* variants or *ALK* rearrangements | Comparators of interest are:  
- Management without genetic testing | Relevant outcomes include:  
- Overall survival  
- Disease-specific survival  
- Test accuracy  
- Test validity  
- Quality of life  
- Treatment-related morbidity |

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### Populations

**Individuals:**
- With advanced-stage non-small-cell lung cancer who are being considered for targeted therapy

### Interventions of interest

- Testing for **BRAF** variants or **ROS1** rearrangements

### Comparators of interest

- Management without genetic testing

### Outcomes

Relevant outcomes include:
- Overall survival
- Disease-specific survival
- Test accuracy
- Test validity
- Quality of life
- Treatment-related morbidity

### Individuals

**Individuals:**
- With advanced-stage non-small-cell lung cancer who are being considered for targeted therapy

### Interventions of interest

- Testing for **KRAS** or **HER2** variants, **RET** rearrangements, or **MET** amplifications

### Comparators of interest

- Management without genetic testing

### Relevant outcomes include

- Overall survival
- Disease-specific survival
- Test accuracy
- Test validity
- Quality of life
- Treatment-related morbidity

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**DESCRIPTION**

Over half of patients with non-small-cell lung cancer (NSCLC) present with advanced and therefore incurable disease. Treatment in this setting has been with platinum-based chemotherapy. The identification of specific, targetable oncogenic “driver mutations” in a subset of NSCLCs has resulted in a reclassification of lung tumors to include molecular subtypes that may direct targeted therapy depending on the presence of a specific variant.

**OBJECTIVE**

The objective of this policy is to examine whether testing for **EGFR**, **BRAF**, **KRAS**, and **HER2** variants; **ALK**, **ROS1**, or **RET** rearrangements; or **MET** amplifications improves outcomes in individuals with advanced-stage non-small-cell lung cancer who are being considered for targeted therapy.

**BACKGROUND**

**Non-Small-Cell Lung Cancer**

Treatment options for NSCLC depend on disease stage and include various combinations of surgery, radiation therapy, chemotherapy, and best supportive care. Unfortunately, in up to 85% of cases, the cancer has spread locally beyond the lungs at diagnosis, precluding surgical eradication. In addition, up to 40% of patients with NSCLC present with metastatic disease.\(^1\) When treated with standard platinum-based chemotherapy, patients with advanced NSCLC have a median survival of 8 to 11 months and a 1-year survival of 30% to 45%.\(^2,3\) More recently, the identification of specific, targetable oncogenic “driver mutations” in a subset of NSCLCs has resulted in a reclassification of lung tumors to include molecular subtypes, which are predominantly of adenocarcinoma histology. Testing for **EGFR** variants and **ALK** rearrangements in clinical decision making for the treatment of NSCLC is routine. The use of testing for other variants to direct targeted therapy is not well established and continues to evolve.
**EGFR Gene**

EGFR, a receptor tyrosine kinase (TK), is frequently overexpressed and activated in NSCLC. Drugs that inhibit EGFR signaling either prevent ligand binding to the extracellular domain (monoclonal antibodies) or inhibit intracellular TK activity (small-molecule tyrosine kinase inhibitors [TKIs]). These targeted therapies dampen signal transduction through pathways downstream to the EGFR, such as the RAS/RAF/MAPK cascade. RAS proteins are G proteins that cycle between active and inactive forms in response to stimulation from cell surface receptors, such as EGFR, acting as binary switches between cell surface EGFR and downstream signaling pathways. These pathways are important in cancer cell proliferation, invasion, metastasis, and stimulation of neovascularization.

Variants in 2 regions of the *EGFR* gene (exons 18-24)—small deletions in exon 19 and a point variant in exon 21 (L858R)—appear to predict tumor response to TKIs such as erlotinib. Likewise, tumors with an acquired exon 20 (T790M) substitution variant appear to respond to osimertinib following the failure of TKI therapy.

The prevalence of *EGFR* variants in NSCLC varies by population, with the highest prevalence in nonsmoking Asian women with adenocarcinoma, in whom *EGFR* variants have been reported to be up to 30% to 50%. The reported prevalence in the white population is approximately 10%.

**ALK Gene**

ALK is a TK that, in NSCLC, is aberrantly activated because of a chromosomal rearrangement that leads to a fusion gene and expression of a protein with constitutive TK activity that has been demonstrated to play a role in controlling cell proliferation. The *EML4-ALK* fusion gene results from an inversion within the short arm of chromosome 2.

The *EML4-ALK* rearrangement (“ALK-positive”) is detected in 3% to 6% of NSCLC patients, with the highest prevalence in never-smokers or light ex-smokers who have adenocarcinoma.

**BRAF Gene**

RAF proteins are serine/threonine kinases that are downstream of RAS in the RAS-RAF-ERK-MAPK pathway. In this pathway, the *BRAF* gene is the most frequently mutated in NSCLC, in 1% to 3% of adenocarcinomas. Unlike melanoma, about 50% of the variants in NSCLC are non-V600E variants. Most *BRAF* variants occur more frequently in smokers.

**ROS1 Gene**

*ROS1* codes for a receptor TK of the insulin receptor family, and chromosomal rearrangements result in fusion genes. The prevalence of *ROS1* fusions in NSCLC varies from 0.9% to 3.7%. Patients with *ROS1* fusions are typically never-smokers with adenocarcinoma.
KRAS Gene
The KRAS gene (which encodes RAS proteins) can harbor oncogenic variants that result in a constitutively activated protein, independent of signaling from the EGFR, possibly rendering a tumor resistant to therapies that target the EGFR. Variants in the KRAS gene, mainly codons 12 and 13, have been reported in 20% to 30% of NSCLC, and occur most often in adenocarcinomas in heavy smokers.

EGFR, ALK, ROS1, and KRAS driver mutations are considered to be mutually exclusive.

HER2 Gene
Human epidermal growth factor receptor 2 (HER2) is a member of the HER (EGFR) family of TK receptors and has no specific ligand. When activated, it forms dimers with other EGFR family members. HER2 is expressed in approximately 25% of NSCLC. HER2 variants are detected mainly in exon 20 in 1% to 2% of NSCLC, predominantly in adenocarcinomas in nonsmoking women.

RET Gene
RET (rearranged during transfection) is a proto-oncogene that encodes a receptor TK growth factor. Translocations that result in fusion genes with several partners have been reported. RET fusions occur in 0.6% to 2% of NSCLCs and 1.2% to 2% of adenocarcinomas.

MET Gene
MET amplification is one of the critical events for acquired resistance in EGFR-mutated adenocarcinomas refractory to EGFR TKIs.

PD-1 / PD-L1
PD-1 is a checkpoint protein on T-cells in the immune system and when bound to PD-L1, blocks the recognition of T-cells that the cancer cells are foreign invaders of the body. Some cancer cells will express PD-L1 in greater amounts, which will cause slowing of the immune attack by the T-cells or cause avoidance of the immune attack. Check-point protein inhibitors such as pembrolizumab, block binding of PD-1 with PD-L1 and potentially enhance immune response against cancer cells.

Targeted Therapies
Three orally administered EGFR-selective, small-molecule TKIs have been identified for treating NSCLC: gefitinib (Iressa®; AstraZeneca, Cambridge, England), erlotinib (Tarceva®; OSI Pharmaceuticals, Melville NY) and afatinib (Gilotrif™; Boehringer Ingelheim, Ingelheim, Germany). Gefitinib, erlotinib, and afatinib currently are approved by the U.S. Food and Drug Administration (FDA) for NSCLC.

Crizotinib is an oral small-molecule TKI that is FDA-approved for patients with locally advanced or metastatic NSCLC who are positive for the ALK or ROS1 gene rearrangements. Ceritinib is a potent ALK inhibitor that is approved for ALK-positive
patients who whose cancer has progressed while taking crizotinib or who could not tolerate crizotinib. Alectinib is a selective ALK inhibitor with high central nervous system penetration that is active against several secondary resistance variants to crizotinib. 

Brigatinib is also an ALK inhibitor that may be able to overcome a broad range of the resistance mechanisms in patients who have progressed on or are intolerant to crizotinib. BRAF or MEK inhibition with TKIs (eg, vemurafenib/dabrafenib or trametinib) was originally approved by FDA for treatment of unresectable or metastatic melanoma with \textit{BRAF}V600 variants. The combination of dabrafenib and trametinib was approved for treatment of metastatic NSCLC in 2017.

For the treatment of \textit{KRAS}-mutated NSCLC, EGFR TKIs and anti-EGFR monoclonal antibodies have been investigated as treatment options. Anti-EGFR monoclonal antibodies include cetuximab and panitumumab. Cetuximab may be used in combination with chemotherapy in patients with advanced or recurrent NSCLC as first-line and maintenance therapy. Panitumumab is not used in NSCLC.

Proposed targeted therapies for the remaining genetic alterations in NSCLC addressed are trastuzumab and afatinib for \textit{HER2} variants, crizotinib for \textit{MET} amplification and cabozantinib for \textit{RET} rearrangements.

\textbf{REGULATORY STATUS}

Table 1 summarizes the FDA-approved targeted treatments for patients with NSCLC along with the concurrently approved diagnostic tests.\textsuperscript{5-14}

\textbf{Table 1.} FDA-Approved Treatment for NSCLC and Companion Diagnostic Tests

<table>
<thead>
<tr>
<th>Treatment</th>
<th>Indication</th>
<th>FDA Approval of Companion Diagnostic Test</th>
</tr>
</thead>
<tbody>
<tr>
<td>Afatinib (Gilotrif)</td>
<td>2013: First line for patients with metastatic NSCLC whose tumors have \textit{EGFR} exon 19 deletions or exon 21 (L858R) substitutions as detected by FDA-approved test 2016: Second line for patients with metastatic squamous NSCLC</td>
<td>2013: therascreen® \textit{EGFR} Rotor-Gene Q polymerase chain reaction (RGQ PCR) kit (Qiagen, Netherlands)</td>
</tr>
<tr>
<td>Alectinib (Alecensa)</td>
<td>2015: Second line for patients with \textit{ALK}-positive metastatic NSCLC who have progressed on or are intolerant of crizotinib</td>
<td></td>
</tr>
<tr>
<td>Ceritinib (Zykadia)</td>
<td>2014: Second line for patients with \textit{ALK}-positive metastatic NSCLC who have progressed on or are intolerant of crizotinib</td>
<td></td>
</tr>
<tr>
<td>Crizotinib (Xalkori)</td>
<td>2011: Patients with \textit{ALK}-positive metastatic NSCLC as detected by FDA-approved test</td>
<td>2011: Vysis ALK Break Apart FISH Probe Kit (Abbott Laboratories, Lake Bluff, IL) 2015: Ventana ALK (D5F3) CDx Assay (Ventana Medical Systems, Tucson, AZ)</td>
</tr>
<tr>
<td>Crizotinib (Xalkori)</td>
<td>2016: Patients with \textit{ROS1}-positive metastatic NSCLC</td>
<td>2017: Oncomine™ Dx Target Test (Thermo Fisher Scientific, Waltham, MA)</td>
</tr>
</tbody>
</table>
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<tbody>
<tr>
<td>Erlotinib (Tarceva)</td>
<td>2013: First line for patients with metastatic NSCLC whose tumors have <strong>EGFR</strong> exon 19 deletions or exon 21 (L858R) substitutions as detected by FDA-approved test</td>
<td>2013: cobas® EGFR Mutation Test (tissue test) (Roche Diagnostics, Indianapolis, IN)</td>
</tr>
<tr>
<td></td>
<td>2010: Maintenance for patients with locally advanced or metastatic NSCLC whose disease has not progressed after 4 cycles of platinum-based chemotherapy</td>
<td>2016: cobas® EGFR Mutation Test v2 (tissue or blood test) (Roche, Diagnostics, Indianapolis, IN)</td>
</tr>
<tr>
<td></td>
<td>2004: Second line for patients with locally advanced or metastatic NSCLC</td>
<td></td>
</tr>
<tr>
<td>Gefitinib (Iressa)</td>
<td>2015: First line for patients with metastatic NSCLC whose tumors have <strong>EGFR</strong> exon 19 deletions or exon 21 (L858R) substitutions as detected by FDA-approved test</td>
<td>2015: therascreen® EGFR Rotor-Gene Q polymerase chain reaction (RGQ PCR) kit</td>
</tr>
<tr>
<td></td>
<td>2003: Second line for patients with locally advanced or metastatic NSCLC</td>
<td>2017: Oncomine™ Dx Target Test</td>
</tr>
<tr>
<td>Necitumumab (Portrazza; Eli Lilly, Indianapolis, IN)</td>
<td>2015: In combination with gemcitabine and cisplatin, as first line for patients with squamous NSCLC</td>
<td></td>
</tr>
<tr>
<td>Osimertinib (Tagrisso)</td>
<td>2015: Second line for patients with metastatic NSCLC whose tumors have <strong>EGFR</strong> T790M variants as detected by FDA-approved test, who have not responded to <strong>EGFR</strong>-blocking therapy</td>
<td>2015: cobas® EGFR Mutation Test v2 (blood test)</td>
</tr>
<tr>
<td>Dabrafenib (Tafinlar) and trametinib (Mekinist) combination</td>
<td>2017: Used in combination: for treatment of patients with metastatic NSCLC with <strong>BRAF</strong> V600E variant</td>
<td>2017: Oncomine™ Dx Target Test</td>
</tr>
</tbody>
</table>

ALK: anaplastic lymphoma kinase; 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.

### POLICY

A. Except as noted below, analysis of 2 types of somatic variants within the EGFR gene—small deletions in exon 19 and a single-nucleotide variant in exon 21 (L858R)—may be considered **medically necessary** to predict treatment response to an EGFR tyrosine kinase inhibitor (TKI) therapy (eg, erlotinib [Tarceva®], gefitinib [Iressa®], or afatinib [Gilotrif®]) in patients with stage III or IV disease with histologic subtypes adenocarcinoma, large cell carcinoma, and non-small-cell lung cancer not otherwise specified.

B. Analysis of the T790M variants in the EGFR gene is considered **medically necessary** as a technique to predict treatment response to osimertinib (Tagrisso™) in patients who have progressed on or after EGFR-TKI therapy.

C. Analysis of 2 types of somatic variants within the EGFR gene—small deletions in exon 19 and a point mutation in exon 21 (L858R)—is considered **experimental / investigational** for patients with advanced squamous cell NSCLC.

D. Analysis of other **EGFR** variants within exons 18 to 24, or other applications related to NSCLC, is considered **experimental / investigational**.

*Contains Public Information*
E. Analysis of somatic rearrangement variants of the ALK gene may be considered medically necessary to predict treatment response to ALK inhibitor therapy (e.g., crizotinib [Xalkori®], ceritinib [Zykadia™], alectinib [Alecensa®], or brigatinib [Alunbrig™]) in patients with stage III or IV disease with histologic subtypes adenocarcinoma, large cell carcinoma, and non-small-cell lung cancer not otherwise specified.

F. Analysis of somatic rearrangement variants of the ALK gene is considered experimental / investigational in all other situations.

G. Analysis of the BRAF V600E variant may be considered medically necessary to predict treatment response to BRAF or MEK inhibitor therapy (e.g., dabrafenib [Tafinlar®] and trametinib [Mekinist®]), in patients with stage III or IV disease with histologic subtypes adenocarcinoma, large cell carcinoma, and non-small-cell lung cancer not otherwise specified.

H. Analysis of somatic rearrangement variants of the ROS1 gene may be considered medically necessary to predict treatment response to ALK inhibitor therapy (crizotinib [Xalkori®]) in patients with stage III or IV disease with histologic subtypes adenocarcinoma, large cell carcinoma, and non-small-cell lung cancer not otherwise specified.

I. Analysis of somatic mutations of the KRAS gene is considered experimental / investigational as a technique to predict treatment non-response to anti-EGFR therapy with tyrosine-kinase inhibitors and for the use of the anti-EGFR monoclonal antibody cetuximab in NSCLC.

J. Analysis for genetic alterations in the genes RET, MET, and HER2 for targeted therapy in patients with NSCLC is considered experimental / investigational.

K. Programmed death receptor 1 (PD-1) or its ligand (PD-L1) expression analysis may be considered medically necessary as a technique to predict treatment response to drug therapy.

Policy Guidelines
1. These tests are intended for use in patients with advanced NSCLC. Patients with either small deletions in exon 19 or a point mutation in exon 21 (L858R) of the tyrosine kinase domain of the epidermal growth factor gene are considered good candidates for treatment with erlotinib, gefitinib, or afatinib. Patients with wild-type variants are unlikely to respond to erlotinib or afatinib; for these patients, other treatment options should be considered.
2. The 2017 guidelines from the National Comprehensive Cancer Network recommend that EGFR variants and ALK rearrangement testing (category 1) as well as ROS1 and BRAF testing (category 2A) be performed in the workup of non-small-cell lung
cancer in patients with metastatic disease with histologic subtypes adenocarcinoma, large cell carcinoma, and non-small-cell lung cancer not otherwise specified.

3. The 2013 guidelines issued jointly by the College of American Pathologists, International Association for the Study of Lung Cancer, and Association for Molecular Pathology recommend:
   a) EGFR mutation and ALK rearrangement testing in patients with lung adenocarcinoma regardless of clinical characteristics (eg, smoking history);
   b) In the setting of fully excised lung cancer specimens, EGFR and ALK testing is not recommended in lung cancers when an adenocarcinoma component is lacking (such as pure squamous cell lacking any immunohistochemical evidence of adenocarcinomatous differentiation); and
   c) In the setting of more limited lung cancer specimens (eg, biopsies, cytology) where an adenocarcinoma component cannot be completely excluded, EGFR and ALK testing may be performed in cases showing squamous cell histology. Clinical criteria (eg, young age, lack of smoking history) may be useful to select a subset of these samples for testing.

RATIONALE
The most recent literature search conducted was conducted through August 25, 2017 (see Appendix Table 1 for genetic testing categories).

The evaluation of a prognostic test focuses on 3 main principles: (1) technical reliability (technical accuracy of the test in detecting a variant that is present or in excluding a variant that is absent); (2) clinical validity (prognostic or predictive performance of the test in predicting future risk of clinical disease or response to therapy); and (3) clinical usefulness (ie, a demonstration that the predictive information can be used to improve patient health outcomes).

For any test, it is important to establish its technical reliability, ie, the ability of the test to accurately and reliably measure the characteristic of interest for which the test was designed to identify or measure. Assessment of technical reliability focuses on specific tests and operators and requires review of unpublished and often proprietary information. Review of specific tests, operators, and unpublished data are outside the scope of this evidence review. This review focuses on the clinical validity and clinical usefulness.

Targeted Therapy for Advanced-Stage Non-Small-Cell Lung Cancer
Clinical Context and Test Purpose
The purpose of identifying targetable oncogenic “driver mutations” in patients who have non-small-cell lung cancer (NSCLC) is to inform a decision whether patients should receive a targeted therapy vs another systemic therapy. Patients who present with advanced disease or recurrence following initial definitive treatment typically receive systemic therapy. Traditionally, the systemic therapy was cytotoxic chemotherapy. However, certain patients may be good candidates for treatment with targeted therapies or immunotherapy. The goal of targeted therapies is to preferentially kill malignant cells without significant damage to normal cells so that there is improved therapeutic efficacy along with decreased toxicity.
The question addressed in this evidence review is this: Does testing for epidermal growth factor receptor (EGFR), BRAF, KRAS, or HER2 variants; ALK, ROS, or RET rearrangements; or MET amplifications improve outcomes in individuals with advanced-stage NSCLC who are being considered for targeted therapy?

The following PICOTS were used to select literature to inform this review.

Patients
The relevant population of interest is individuals with advanced NSCLC who are being considered for targeted therapy.

Intervention
The intervention of interest is testing for somatic genome alterations known as "driver mutations," specifically EGFR, BRAF, KRAS, BRAF HER2 variants; ALK, ROS, or RET rearrangements; or MET amplifications.

Comparator
The comparator of interest is standard management without testing for driver mutations. Standard management consists primarily of chemotherapy, although some patients are candidates for immunotherapy.

Outcomes
Beneficial outcomes resulting from a true positive test result are prolonged survival, reduction in toxicity, and improved quality of life associated with receiving a more effective and less toxic targeted therapy instead of chemotherapy in those with driver mutations. Beneficial outcomes from a true negative result are prolonged survival associated with receiving chemotherapy in those without driver mutations. Harmful outcomes resulting from a false negative test result include shorter survival from receiving less effective and more toxic chemotherapy in those with driver mutations; possible harmful outcomes resulting from a false positive test result are shorter survival from receiving potentially ineffective targeted treatment and delay in initiation of chemotherapy in those without driver mutations.

Time
Due to the poor prognosis of advanced NSCLC, the duration of follow-up for the outcomes of interest are 6 months and 1 year.

Setting
Treatment recommendations for patients with advanced NSCLC are usually made in the tertiary care setting, ideally in consultation with a multidisciplinary team of pathologists, thoracic surgeons, and oncologists.

Epidermal Growth Factor Receptor Gene Variants
Somatic variants in the tyrosine kinase domain of the EGFR gene, notably small deletions in exon 19 and a point mutation in exon 21 (L858R, indicating substitution of leucine by arginine at codon position 858) are the most commonly found EGFR variants associated with sensitivity to EGFR tyrosine kinase inhibitors (TKIs; afatinib, erlotinib, gefitinib). These variants are referred to as sensitizing variants. Almost all patients who initially respond to an EGFR TKI eventually develop disease progression. The most common of these secondary variants, called resistance
variants, involves the substitution of methionine for threonine at position 790 (T790M) on exon 20.

**EGFR Variant Frequency**

In 2009, Rosell et al reported EGFR variants in 16.6% of the overall patient sample but noted an increased prevalence in women (69.7%), patients who never smoked (66.6%), and patients with adenocarcinomas (80.9%).

In a comprehensive analysis of 14 studies involving 2880 patients, Mitsudomi et al (2006) reported EGFR variants in 10% of men, 7% of non-Asian patients, 7% of current or former smokers, and 2% of patients with nonadenocarcinoma histologies. Eberhard et al (2005) observed EGFR variants in 6.4% of patients with squamous cell carcinomas (SCCs) and Rosell et al (2009) observed EGFR variants in 11.5% of patients with large cell carcinomas. Both studies had small sample sizes. The acquired EGFR T790M variant has been estimated to be present in 50% to 60% of TKI-resistant cases in approximately 200 patients from 2 studies.

Fang et al (2013) reported EGFR variants (all L858R) in 3 (2%) of 146 consecutively treated Chinese patients with early-stage SCC. In a separate cohort of 63 Chinese patients with SCC who received erlotinib or gefitinib as second- or third-line treatment (63% never-smokers, 21% women), EGFR variant prevalence (all exon 19 deletion or L858R) was 23.8%.

**Clinical Validity**

EGFR-sensitizing and -resistance variants can be detected by direct sequencing, polymerase chain reaction (PCR) technologies, or next-generation sequencing (NGS). Gene sequencing is considered an analytic criterion standard. In 2010, the Canadian Agency for Drugs and Technologies in Health published a rapid response report on EGFR variant analysis. Based on 11 observational studies, the report authors concluded that PCR-based approaches identify EGFR variants with a sensitivity equivalent to that of direct sequencing.

Four tests have been approved as companion diagnostics to detect EGFR-resistance variants (exon 19 deletions or exon 21 L858R substitutions) for at least one of the EGFR TKIs (afatinib, erlotinib, or gefitinib): the therascreen EGFR Rotor-Gene Q polymerase chain reaction (RGQ PCR) kit, cobas EGFR Mutation Test v1 and v2, and Oncomine Dx Target Test. The cobas v2 test also is approved as a companion diagnostic to detect the T790M resistance variant to select patients for treatment with osimertinib.

The clinical validity of the therascreen RGQ PCR kit was demonstrated in a retrospective analysis of patients screened for a phase 3, open-label randomized controlled trial (RCT) comparing afatinib with chemotherapy in treatment-naive patients with stage IIIIB or IV NSCLC, in which the EGFR variants for enrollment were determined using a Clinical Trial Assay (CTA) conducted at central laboratories. The positive percent agreement (PPA) of therascreen vs CTA for detection of EGFR-sensitizing variants was 98% (95% confidence interval [CI], 95% to 99%) and negative percent agreement (NPA) was 97% (95% CI, 94% to 99%). Overall, a statistically significant efficacy benefit for afatinib vs chemotherapy was reported in the EGFR-positive patients as measured by the therascreen EGFR RGQ PCR Kit (hazard ratio [HR], 0.49; 95% CI, 0.35 to 0.69) that was similar to the efficacy in the overall population which was EGFR-positive by the CTA (HR=0.58; 95% CI, 0.43 to 0.78).
The clinical validity of the cobas EGFR Mutation Test v1 was demonstrated in a retrospective analysis of patients screened for a phase 3, open-label RCT comparing erlotinib with chemotherapy in treatment-naive patients with advanced NSCLC. In this RCT, the EGFR variants for enrollment were determined with a CTA at a central laboratory using Sanger sequencing first for determination of EGFR variants status, followed by confirmatory testing for exon 19 deletions and exon 21 L858R variants. The PPA of cobas vs CTA for detection of EGFR-sensitizing variants was 94% (95% CI, 89% to 97%) and NPA was 98% (95% CI, 95% to 99%). Overall, a statistically significant efficacy benefit for erlotinib vs chemotherapy was reported in the EGFR-positive patients as measured by the cobas EGFR Mutation Test v1 (HR=0.34; 95% CI, 0.21 to 0.54) that was similar to the efficacy in the overall population, which was EGFR-positive by the CTA (HR=0.34; 95% CI, 0.23 to 0.49). The cobas EGFR Mutation Test v2 expanded the indication for the use of the cobas EGFR Mutation Test to add the detection of the exon 20 (T790M) substitution variant in NSCLC patients for whom osimertinib (Tagrisso) treatment is indicated. The clinical validity of the cobas EGFR Mutation Test v2 was demonstrated in retrospective analyses of patients enrolled in a phase 2, single-arm study of osimertinib for EGFR-sensitizing variant-positive metastatic NSCLC who had progressed following prior therapy with an approved EGFR TKI. The osimertinib response rate in the patients identified as EGFR T790M-positive by the cobas v2 test was 62% (95% CI, 55% to 69%).

The clinical validity of the Oncomine Dx Target Test was demonstrated in a retrospective analysis of patients screened for a phase 3, open-label RCT, which included newly diagnosed patients with stage IIIIB or IV or recurrent NSCLC, in which the EGFR variant for enrollment was determined using therascreen. The PPA of Oncomine vs therascreen for detection of EGFR-sensitizing variants was 99% (95% CI, 93% to 100%) and NPA was 99% (95% CI, 96% to 100%). No data on the effectiveness of gefitinib in patients identified as EGFR-positive by Oncomine were reported.

EGFR variants may provide prognostic information about disease recurrence and survival as well as predictive information about treatment response. In a 2005 study by Eberhard et al, improved outcomes were observed for EGFR variant–positive patients compared with wild-type patients regardless of treatment (standard chemotherapy or standard chemotherapy plus erlotinib). The objective radiologic response was 38% vs 23% (p=0.01), the median time to progression was 8 months vs 5 months (p<0.001), and the median OS was not reached vs 10 months (p<0.001).

Clinical Utility
A TEC Assessment on this topic was published in 2007. It concluded that there was insufficient evidence to permit conclusions about the clinical validity or utility of EGFR variant testing to predict erlotinib sensitivity or to guide treatment in patients with NSCLC. This Assessment was updated in 2010, with revised conclusions indicating that EGFR variant testing has clinical utility in selecting or deselecting patients for treatment with erlotinib.

A 2013 meta-analysis of 23 trials of erlotinib, gefitinib, and afatinib in patients with advanced NSCLC reported improved progression-free survival (PFS) in EGFR variant–positive patients treated with EGFR TKIs in the first- and second-line settings and for maintenance therapy. Comparators were with chemotherapy, chemotherapy and placebo, and placebo in the first-line, second-line, and maintenance therapy settings, respectively. Among EGFR variant–negative patients, PFS was improved with EGFR TKIs compared with placebo maintenance but not in the first- and second-line settings. Overall survival (OS) did not differ between treatment groups in
either variant-positive or variant-negative patients. Statistical heterogeneity was not reported for any outcome. Similar meta-analyses confirming the PFS and OS results and conclusions for EGFR-positive patients have been published.28-33

Erlotinib
In 2012, Petrelli et al reported on a meta-analysis (13 randomized trials) of 1260 patients with EGFR-mutated NSCLC who received TKIs for first-line, second-line, or maintenance therapy.34 The comparator was standard therapy. Overall, reviewers noted that use of EGFR TKIs increased the chance of obtaining an objective response almost two-fold compared with chemotherapy. Response rates were 70% versus 33% in first-line trials and 47% vs 28.5% in second-line trials. TKIs reduced the hazard of progression by 70% in all trials and by 65% in first-line trials; however, they did not improve OS.

A summary of 3 key RCTs establishing the superiority of erlotinib over chemotherapy in the first-line setting is given in Tables 2 and 3. The 3 RCTs included 555 patients with stage III B or IV NSCLC. All reported clinically and statistically significant improvement in PFS (HR range, 0.16-0.37) but no improvement in OS with erlotinib compared with chemotherapy. Grade 3 or greater adverse events and serious adverse events occurred in fewer patients in the erlotinib groups.

Table 2. Characteristics of RCTs of First-Line Erlotinib vs Chemotherapy in EGFR-Variant NSCLC

<table>
<thead>
<tr>
<th>Author (Year); Study</th>
<th>Countries</th>
<th>Sites</th>
<th>Dates</th>
<th>Participants</th>
<th>Erlotinib</th>
<th>Chemotherapy</th>
</tr>
</thead>
<tbody>
<tr>
<td>Zhou (2011, 2015)35,36; OPTIMAL (NCT00874419)</td>
<td>China</td>
<td>22</td>
<td>NR</td>
<td>165 patients with stage III B/IV NSCLC</td>
<td>83 assigned to erlotinib (150 mg qd)</td>
<td>82 assigned to carboplatin (AUC5) and gemcitabine (1000 mg/m²)</td>
</tr>
<tr>
<td>Rosell (2012)37; EURTAC (NCT00446225)</td>
<td>France, Italy, Spain</td>
<td>42</td>
<td>2007-2011</td>
<td>173 patients with stage III B/IV NSCLC</td>
<td>86 assigned to erlotinib (150 mg qd)</td>
<td>87 assigned to cisplatin (75 mg/m²), docetaxel (75 mg/m²), or gemcitabine (1250 mg/m²)</td>
</tr>
<tr>
<td>Wu (2015)38; ENSURE (NCT01342965)</td>
<td>China, Malaysia, Philippines</td>
<td>30</td>
<td>2011-2012</td>
<td>217 patients with stage III B/IV NSCLC</td>
<td>110 assigned to erlotinib (150 mg qd)</td>
<td>117 assigned to gemcitabine (1250 mg/m²) and cisplatin (75 mg/m²)</td>
</tr>
</tbody>
</table>

AUC5: area under the concentration time curve of 5.0 mg/mL/min; EGFR: epidermal growth factor receptor; NR: not reported; NSCLC: non-small-cell lung cancer; qd: every day; RCT: randomized controlled trial.

Table 3. Results of RCTs of First-Line Erlotinib vs Chemotherapy in EGFR-Variant NSCLC

<table>
<thead>
<tr>
<th>Study (Year)</th>
<th>Median Progression-Free Survival, mo</th>
<th>Median Overall Survival, mo</th>
<th>Serious</th>
<th>Grade 3 or 4</th>
</tr>
</thead>
<tbody>
<tr>
<td>Erlotinib</td>
<td>13.1 (10.6 to 16.5)</td>
<td>22.8</td>
<td>2</td>
<td>Neutropenia 0</td>
</tr>
<tr>
<td>Chemotherapy</td>
<td>4.6 (4.2 to 5.4)</td>
<td>27.2</td>
<td>14</td>
<td>Neutropenia 42</td>
</tr>
<tr>
<td>TE (95% CI)</td>
<td>HR=0.16 (0.10 to 0.26)</td>
<td>HR=1.19 (0.83 to 1.71)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>EURTAC (2012)37</td>
<td>Erlotinib</td>
<td>9.7 (8.4 to 12.3)</td>
<td>19.3</td>
<td>6</td>
</tr>
</tbody>
</table>
### Table 4. Characteristics of RCTs of First-Line Gefitinib vs Chemotherapy in EGFR-Variant NSCLC

<table>
<thead>
<tr>
<th>Author (Year); Study</th>
<th>Countries</th>
<th>Sites</th>
<th>Dates</th>
<th>Participants</th>
<th>Gefitinib</th>
<th>Chemotherapy</th>
</tr>
</thead>
<tbody>
<tr>
<td>Mok (2009)36; IPASS (NCT00322452)</td>
<td>9 East Asian countries</td>
<td>87</td>
<td>2006-2007</td>
<td>1217 patients with stage IIIIB/IV NSCLC (261 EGFR-positive)</td>
<td>609 assigned to gefitinib (250 mg/d)</td>
<td>608 assigned to paclitaxel (200 mg/m²) and</td>
</tr>
</tbody>
</table>
## Description of Interventions

<table>
<thead>
<tr>
<th>Author (Year); Study</th>
<th>Countries</th>
<th>Sites</th>
<th>Dates</th>
<th>Participants</th>
<th>Gefitinib</th>
<th>Chemotherapy</th>
</tr>
</thead>
<tbody>
<tr>
<td>Mitsudomi (2010); West Japan Oncology Group 172 trial; WJ TOG3405</td>
<td>Japan</td>
<td>36</td>
<td>2006-2009</td>
<td>177 patients with stages IIIB/IV or recurrent NSCLC</td>
<td>88 assigned to gefitinib (250 mg/d)</td>
<td>89 assigned to cisplatin (80 mg/m²) and docetaxel (60 mg/m²)</td>
</tr>
<tr>
<td>Maemondo (2010), Inoue (2013); NEJ 002</td>
<td>Japan</td>
<td>43</td>
<td>2006-2009</td>
<td>230 patients with stages IIIB/IV NSCLC or postoperative relapse</td>
<td>115 assigned to gefitinib (250 mg/d)</td>
<td>115 assigned to paclitaxel (200 mg/m²) and carboplatin (AUC6)</td>
</tr>
</tbody>
</table>

AUC5: area under the concentration time curve of 5.0 mg/mL/min; AUC6: area under the concentration time curve of 6.0 mg/mL/min; EGFR: epidermal growth factor receptor; NSCLC: non-small-cell lung cancer; RCT: randomized controlled trial.

## Table 5. Results of RCTs of First-Line Gefitinib vs Chemotherapy in EGFR-Variant NSCLC

<table>
<thead>
<tr>
<th>Study (Year)</th>
<th>Median Progression-Free Survival, mo</th>
<th>Median Overall Survival, mo</th>
<th>Serious</th>
<th>Grade 3 or 4</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>IPASS (2009)</strong>&lt;sup&gt;52&lt;/sup&gt;</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Gefitinib</td>
<td>≈9.6&lt;sup&gt;a&lt;/sup&gt;</td>
<td>NR</td>
<td>16.3%</td>
<td>Rash 3.1; Diarrhea 3.8; Neurotoxic effects 0.3; Neutropenia 3.7; Anemia 2.2; Leukopenia 1.5</td>
</tr>
<tr>
<td>Chemotherapy</td>
<td>≈5.8&lt;sup&gt;b&lt;/sup&gt;</td>
<td>NR</td>
<td>15.6%</td>
<td>Rash 0.8; Diarrhea 1.4; Neurotoxic effects 4.9; Neutropenia 67.1; Anemia 10.6; Leukopenia 35.0</td>
</tr>
<tr>
<td><strong>TE (95% CI)</strong></td>
<td>HR=0.48 (0.36 to 0.64)</td>
<td>HR=0.78 (0.50 to 1.20)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>WJ TOG3405 (2010)&lt;sup&gt;51&lt;/sup&gt;</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Gefitinib</td>
<td>9.2 (8.0 to 13.9)</td>
<td>34.8 (26.0 to 39.5)</td>
<td>ALT/AST elevation 27.5; Rash 2.3; Fatigue 2.3; Neutropenia 0; Leucocytopenia 0; Anemia 0</td>
<td></td>
</tr>
<tr>
<td>Chemotherapy</td>
<td>6.3 (5.8 to 7.8)</td>
<td>37.3 (31.2 to 45.5)</td>
<td>ALT/AST elevation 2.3; Rash 0; Fatigue 2.3; Neutropenia 84; Leucocytopenia 50; Anemia 17</td>
<td></td>
</tr>
<tr>
<td><strong>TE (95% CI)</strong></td>
<td>HR=0.49 (0.34 to 0.71)</td>
<td>HR=1.25 (0.88 to 1.78)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>NEJ 002 (2010, 2013)&lt;sup&gt;52,53&lt;/sup&gt;</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Gefitinib</td>
<td>10.8</td>
<td>27.7</td>
<td>Rash 5.3; Neuropathy 0; Arthralgia 0.9</td>
<td></td>
</tr>
<tr>
<td>Study (Year)</td>
<td>Median Progression-Free Survival, mo</td>
<td>Median Overall Survival, mo</td>
<td>Adverse Events, %</td>
<td></td>
</tr>
<tr>
<td>-------------</td>
<td>-------------------------------------</td>
<td>----------------------------</td>
<td>------------------</td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>Pneumonitis</td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>2.6</td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>Aminotransferase elevation</td>
<td>26.3</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>Neutropenia</td>
<td>0.9</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>Anemia</td>
<td>0.9</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>Thrombocytopenia</td>
<td>0.9</td>
</tr>
<tr>
<td>Chemotherapy</td>
<td>5.4</td>
<td>26.6</td>
<td>Rash</td>
<td>2.7</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>Neuropathy</td>
<td>6.2</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>Arthralgia</td>
<td>7.1</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>Pneumonitis</td>
<td>0.9</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>Aminotransferase elevation</td>
<td>0.9</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>Neutropenia</td>
<td>65.5</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>Anemia</td>
<td>5.3</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>Thrombocytopenia</td>
<td>3.5</td>
</tr>
</tbody>
</table>

TE (95% CI) HR = 0.30 (0.22 to 0.41) HR=0.89 (0.63 to 1.24)

ALT: alanine aminotransferase; AST: aspartate aminotransferase; CI: confidence interval; EGFR: epidermal growth factor receptor; HR: hazard ratio; NR: not reported; NSCLC: non-small-cell lung cancer; RCT: randomized controlled trial; TE: treatment effect.

a Analysis includes EGFR-positive only.
b Analysis includes all patients with safety data.
c Estimated from figure.

**Afatinib**

Unlike erlotinib and gefitinib, which selectively inhibit EGFR, afatinib inhibits not only EGFR but also human epidermal growth factor receptor 2 (HER2) and HER4, and may have activity in patients with acquired resistance to TKIs; such patients often harbor a T790M variant (substitution of threonine by methionine at codon 790) in EGFR exon 20. The efficacy and safety of afatinib were evaluated in the LUX-Lung series of studies.

LUX-Lung 3 was an RCT in 345 patients with stage IIIIB or IV, EGFR variant–positive, lung adenocarcinoma who were previously untreated for advanced disease. Fifty-two percent of patients were Asian, 26% were white, and 90% (308 patients) had common EGFR variants (exon 19 deletion or L858R substitution variant in exon 21). Patients received afatinib or chemotherapy (cisplatin plus pemetrexed). In stratified analysis of patients with common EGFR variants, the median PFS was 13.6 months for the afatinib group and 6.9 months for the chemotherapy group (HR=0.47; 95% CI, 0.34 to 0.65; p=0.001). The median PFS for the 10% of patients who had other EGFR variants was not reported, but the median PFS for the entire patient sample was 11.1 months in the afatinib group and 6.9 months in the chemotherapy group (HR=0.58; 95% CI, 0.43 to 0.78; p=0.001). The incidence of objective response in the entire patient sample was 56% in the afatinib group and 23% in the chemotherapy group (p=0.001). With a median follow-up of 16.4 months, the median OS was not reached in any group; preliminary analysis indicated no difference in OS between the 2 treatment groups in the entire patient sample (HR=1.12; 95% CI, 0.73 to 1.73; p=0.60). Patients in the afatinib group reported greater improvements in dyspnea, cough, and global health status/quality of life than those in the chemotherapy group. Grade 3 or higher diarrhea, rash, and paronychia (nail infection) occurred in 14%, 16%, and 11% of afatinib-treated patients, respectively, and in no patients in the chemotherapy group. Grade 3 or higher mucositis (primarily stomatitis) occurred in 9% of the afatinib group and 1% of the chemotherapy group. Similar results were reported in 2014 phase 3 trial conducted in 364 Asian patients (Lux-Lung 6) comparing afatinib with gemcitabine plus cisplatin. PFS was 11.0 in
the afatinib group and 5.6 months in the chemotherapy group (HR=0.28; 95% CI, 0.20 to 0.39) and the response rate was 67% and 23%.

Three other published LUX-Lung studies evaluated patients with stage IIIIB or IV lung adenocarcinoma who were previously treated for advanced disease, but design features limit interpretation of results.

- LUX-Lung 2 was a single-arm study (2012) of afatinib in 129 patients (87% Asian, 12% white) with EGFR variant–positive disease. Patients had been treated with chemotherapy but not with EGFR-targeted therapy; approximately half of patients (enrolled after a protocol amendment were chemotherapy-naive. Objective responses (primarily partial responses) were observed in 66% of 106 patients with common EGFR variants (exon 19 deletion or L858R) and in 39% of 23 patients with other EGFR variants. The median PFS was 13.7 months in patients with common EGFR variants and 3.7 months in patients with other EGFR variants (p not reported). Results for variant-negative patients were not reported.

- LUX-Lung 1 and LUX-Lung 4 enrolled patients who had progressed on previous treatment with erlotinib, gefitinib, or both for advanced disease. Neither study prospectively genotyped patients. In the LUX-Lung 1 double-blind RCT, 96 (66% Asian, 33% white) of 585 enrolled patients were EGFR variant–positive (76 common EGFR variant–positive). In this group, the median PFS was 3.3 months in the afatinib group and 1.0 month in the placebo group (HR=0.51; 95% CI, 0.31 to 0.85; p=0.009). In 45 variant-negative patients, the median PFS was 2.8 months in the afatinib group and 1.8 months in the placebo group, a statistically nonsignificant difference (p=0.22), possibly due to small group sizes. LUX-Lung 4 was a single-arm study (2013) of afatinib in 62 Japanese patients. Objective responses occurred in 2 (5%) of 36 patients with common EGFR variants and in none of 8 patients with other EGFR variants (p>0.05).

Comparative Effectiveness of EGFR TKIs

As the previous sections have shown, erlotinib, gefitinib, and afatinib all have improved efficacy compared with chemotherapy in patients who have NSCLC and EGFR-sensitizing variants and are well tolerated. RCTs directly comparing the EGFR TKIs have been conducted.

Yang et al (2017) published a meta-analysis of studies comparing erlotinib, gefitinib, and afatinib. Eight randomized trials (n=2225 patients) and 82 cohort studies (n=15,396 patients) were included. However, results were not presented for trials vs cohort studies separately. Of the 8 RCTs identified, three were published in journals not available through PubMed, one was available as an abstract only, one was conducted in squamous cell NSCLC, and one did not require testing to confirm EGFR variant status for enrollment; these RCTs will not be discussed further. The two remaining studies are briefly discussed below.

Both RCTs have compared gefitinib with erlotinib in patients who had EGFR-sensitizing variants. Urata et al (2016) reported on a phase 3 RCT of 401 patients with EGFR variants randomized to gefitinib or erlotinib. The median PFS was 8.3 months (95% CI, 7.2 to 9.7 months) for patients receiving gefitinib and 10.0 months for those receiving erlotinib (95% CI, 8.5 to 11.2 months). Rash was more common with erlotinib (18.1% vs 2.2%) while both alanine aminotransferase elevation and aspartate aminotransferase elevation were more common with gefitinib (6.1% vs 2.2% and 13.0% vs 3.3%, respectively). Similarly, Yang et al (2017) reported a median PFS of
13.0 for erlotinib and 10.4 months for gefitinib (HR=0.81; 95% CI, 0.62 to 1.05) in 256 patients with no differences in rates of grade 3 of 4 adverse events.63

LUX-7 was a phase 2b, head-to-head trial of afatinib vs gefitinib for the treatment of first-line EGFR variant−positive (del19 and L858R) adenocarcinoma of the lung.61 LUX-7 included 319 patients in a 1:1 ratio to afatinib 40 mg/d or gefitinib 250 mg/d, stratified by variant type (del19 and L858R) and brain metastases (present vs absent). In the overall population, PFS was significantly improved with afatinib than with gefitinib (HR=0.73; 95% CI, 0.57 to 0.95; p=0.02). Time-to-treatment failure also showed improvement in favor of afatinib (HR=0.73; 95% CI, 0.58 to 0.92; p=0.01). The objective response rate (ORR) was significantly higher in the afatinib group (70% vs 56%; p=0.01). Several grade 3 or 4 adverse events were more common with afatinib compared with gefitinib including diarrhea (13% vs 1%) and rash (9% vs 3%); liver enzyme elevations were more common with gefitinib (0% vs 9%). Serious events occurred in 11% of patients in the afatinib group and 4% in the gefitinib group.

Osimertinib
In 2015, the U.S. Food and Drug Administration (FDA) granted accelerated approval to osimertinib for treatment of metastatic EGFR T790M variant−positive NSCLC who have progressed on or after EGFR-TKI therapy.64 The therapy was approved along with an FDA-approved companion test, the cobas EGFR Mutation Test v2, which is a blood-based genetic test to detect EGFR variants including the T790M variant. Approval was based on 2 multicenter, single-arm studies.65

The osimertinib label describes the 2 studies.64 Eligible patients had metastatic EGFR T790M variant−positive NSCLC and had progressed on prior systemic therapy, including an EGFR TKI. Patients received osimertinib 80 mg once daily. The first study enrolled 201 patients; the second study enrolled 210 patients. The major efficacy outcome measure of both trials was ORR assessed by a blinded, independent review committee. The median duration of follow-up of 4.2 months in the first study and 4.0 months in the second. The ORR was similar in the 2 studies. The pooled ORR was 59% (95% CI, 54% to 64%); 0.5% achieved complete response and 59% achieved partial response. The most common adverse reactions were diarrhea (42%), rash (41%), dry skin (31%), and nail toxicity (25%). Serious adverse reactions reported in 2% or more patients were pneumonia and pulmonary embolus. Fatal adverse reactions included the following: 4 patients with interstitial lung disease/pneumonitis; 4 patients with pneumonia, and 2 patients with cerebral vascular accident/cerebral hemorrhage.

One RCT has been conducted and is described in Tables 6 and 7. Osimertinib was associated with clinically and statistically significantly prolonged PFS and higher response rates compared with chemotherapy with lower rates of grade 3 and 4 adverse events. However, interstitial lung disease–like adverse events and QT prolongation were more common with osimertinib.

<table>
<thead>
<tr>
<th>Author (Year); Study</th>
<th>Countries</th>
<th>Sites</th>
<th>Dates</th>
<th>Participants</th>
<th>Osimertinib</th>
<th>Chemotherapy</th>
</tr>
</thead>
<tbody>
<tr>
<td>Mok (2017)66; AURA3 (NCT02151981)</td>
<td>18 countries in North America, Europe, Australia, Asia</td>
<td>126</td>
<td>2014-2015</td>
<td>419 T790M-positive advanced NSCLC who had disease progression after first-line EGFR-TKI therapy n=279 to osimertinib (80 mg/d)</td>
<td>n=140 to platinum-pemetrexed (500 mg/m² of body-surface area) plus carboplatin (target</td>
<td></td>
</tr>
</tbody>
</table>
**Table 7. Osimertinib Randomized Controlled Trial Results in EGFR-Variant NSCLC**

<table>
<thead>
<tr>
<th>Study (Year)</th>
<th>PFS, mo</th>
<th>OS, mo</th>
<th>ORR (95% CI%)</th>
<th>Grade ≥3 AEs, %</th>
<th>ILD-Like AEs, %</th>
<th>Prolongation of QT Interval, %</th>
</tr>
</thead>
<tbody>
<tr>
<td>AURA3 (2017)¹⁶⁶</td>
<td>N=419</td>
<td>N=419</td>
<td></td>
<td>N=415</td>
<td>N=415</td>
<td>N=415</td>
</tr>
<tr>
<td>Osimertinib</td>
<td>10.1</td>
<td>NR</td>
<td>71% (65 to 76)</td>
<td>23</td>
<td>4</td>
<td>4</td>
</tr>
<tr>
<td>Platinum-pemetrexed</td>
<td>4.4</td>
<td>NR</td>
<td>31% (24 to 40)</td>
<td>47</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>TE (95% CI)</td>
<td>HR=0.30 (0.23 to 0.41; p&lt;0.001)</td>
<td>OR=5.4 (3.5 to 8.5)</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

AE: adverse event; CI: confidence interval; EGFR: epidermal growth factor receptor; HR: hazard ratio; ILD: interstitial lung disease; NR: not reported; NSCLC: non-small-cell lung cancer; OR: odds ratio; ORR: objective response rate; OS: overall survival; PFS: progression-free survival; TE: treatment effect.

**Section Summary: Epidermal Growth Factor Receptor Gene Variants**

Several RCTs, nonconcurrent prospective studies, and single-arm enrichment studies have demonstrated that patients with EGFR-sensitivity variants benefit from erlotinib, gefitinib, or afatinib therapy and patients with EGFR-resistance variant (T790M) benefit from osimertinib. These observations have been made in populations primarily with adenocarcinomas. Currently, there is little evidence to indicate that EGFR variant testing can guide treatment selection in patients with squamous cell histology. FDA has approved several companion diagnostics for detecting EGFR variants to aid in selecting NSCLC patients for treatment with erlotinib, gefitinib, afatinib and osimertinib.

Patients who are found to have wild-type tumors are unlikely to respond to erlotinib, gefitinib, or afatinib. They patients should be considered candidates for alternative therapies.

**ALK Gene Rearrangements**

ALK gene rearrangements most often consist of an inversion in chromosome 2 which leads to fusion with the echinoderm microtubule-associated protein like 4 (EML4) gene and a novel fusion oncogene EML4-ALK. This inversion causes abnormal expression and activation of ALK tyrosine kinase.⁶⁷ This rearrangement occurs in 3% to 6% of NSCLC.

**Clinical Validity**

Several methods are available to detect ALK gene rearrangements or the resulting fusion proteins in tumor specimens including fluorescence in situ hybridization (FISH), immunohistochemistry, reverse transcription polymerase chain reaction (RT-PCR) of cDNA, and NGS. Two tests have been approved by FDA as companion diagnostics to detect ALK rearrangements for treatment with crizotinib: the Vysis ALK Break Apart FISH Probe Kit and Ventana ALK (D5F3) CDx Assay.

The Vysis kit is a FISH-based assay. The clinical validity of the Vysis ALK Break Apart FISH Probe Kit was demonstrated in a retrospective analysis of patients screened for a phase 2, open-label single-arm study of crizotinib in patients with stage IIIIB or IV NSCLC.⁶⁸ The response rate for crizotinib in 136 ALK-positive patients was 50% (95% CI, 42% to 59%) with a median duration...
of response of 42 weeks (range, 6-42). The response rate for 19 ALK-negative patients was 26% (95% CI, 9% to 51%).

The Ventana assay is an immunohistochemical-based assay. The clinical validity of the Ventana ALK (D5F3) CDx Assay was demonstrated in a retrospective analysis of patients screened for an open-label RCT of crizotinib vs platinum-doublet chemotherapy in patients with stage IIIIB or IV NSCLC. The concordance between the Ventana and Vysis tests were calculated using patient samples analyzed at an independent, central laboratory. The PPA was 86.0% (95% CI, 80.2% to 90.4%) and the NPA was 96.3% (94.7% to 97.4%). Overall, in 343 patients who were ALK-positive by the Vysis assay, crizotinib was associated with longer PFS compared with chemotherapy (HR=0.45; 95% CI, 0.36 to 0.60). In the subset of 141 patients who were also ALK-positive by the Ventana assay, the results were similar (HR=0.40; 95% CI, 0.25 to 0.64). In the 25 patients that were ALK-positive by the Vysis assay and ALK-negative by the Ventana assay, the relative effect of crizotinib was not clear (HR=1.71; 95% CI, 0.43 to 6.79).

Clinical Utility

**Crizotinib**

The accelerated approval of crizotinib by FDA was based on phase 1 and 2 trials in which crizotinib showed marked antitumor activity in patients with ALK-positive advanced NSCLC, with an ORR of 60% and PFS range from 7 to 10 months. These results were confirmed in 2 subsequent phase 3 trials.

A phase 3, open-label trial randomized 347 patients with previously treated, locally advanced, or metastatic ALK-positive lung cancer to oral crizotinib twice daily (n=173) or chemotherapy (n=174) every 3 weeks. All patients had received 1 platinum-based chemotherapy regimen before the trial. The extent of metastatic disease was 95% and 91% in patients in the crizotinib and chemotherapy groups, respectively, and tumor histology was adenocarcinoma in 95% and 94%, respectively. The primary end point was PFS. Patients in the chemotherapy group who experienced progressive disease were allowed to cross over to crizotinib as part of a separate study. The median PFS was 7.7 months in the crizotinib group vs 3.0 months in the chemotherapy group (HR for progression or death with crizotinib, 0.49; 95% CI, 0.37 to 0.64; p<0.001). Partial response rates with crizotinib were 65% (95% CI, 58% to 72%) vs 20% (95% CI, 14% to 26%) with chemotherapy (p<0.001). Interim analysis of OS showed no significant improvement with crizotinib compared with chemotherapy (HR for death in the crizotinib group, 1.02; 95% CI, 0.68 to 1.54; p=0.54). The median follow-up for OS was 12.2 and 12.1 months in the crizotinib and chemotherapy groups, respectively. Patients reported greater reductions in lung cancer symptoms and greater improvement in global quality of life with crizotinib than with chemotherapy.

A phase 3, open-label trial compared crizotinib and chemotherapy in 343 previously untreated patients with ALK-positive advanced nonsquamous NSCLC. Patients were randomized to oral crizotinib twice daily or pemetrexed plus cisplatin or carboplatin every 3 weeks for up to 6 cycles. If there was disease progression for patients receiving chemotherapy, crossover to crizotinib was allowed. PFS was the primary end point. PFS was 10.9 months compared with 7.0 months for the group that received crizotinib vs chemotherapy, respectively (HR for progression or death with crizotinib, 0.45; 95% CI, 0.35 to 0.60; p<0.001); ORRs (complete and partial responses) were 74% and 45%, respectively (p<0.001). The median OS was not reached in either group; the probability of 1-year survival with crizotinib was 84% and 79% with chemotherapy. Crizotinib
was associated with patient-reported greater reduction in lung cancer symptoms and greater improvements in quality of life.

**Other ALK Inhibitors**

Three other ALK TKIs are FDA-approved but without separate companion diagnostics. Ceritinib has demonstrated superior efficacy concerning PFS when compared with chemotherapy in both the first-line and second-line (following crizotinib) settings in the ASCEND-4 and ASCEND-5 RCTs.\(^{71,72}\) Alectinib was associated with response rates of approximately 50% in patients who had progressed on crizotinib in 2 phase 2 studies.\(^{73,74}\) Alectinib has also shown superior efficacy and lower toxicity when compared with crizotinib in the first-line setting in the ALEX and J-ALEX phase 3 RCTs.\(^{75,76}\) Brigatinib has shown promise in early phase 1 and phase 2 studies with PFS of almost 13 months in patients with crizotinib-refractory disease.\(^{77,78}\)

**Section Summary: ALK Gene Rearrangements**

Crizotinib was granted accelerated approval by FDA in 2011 for patients with locally advanced or metastatic NSCLC, based on ORRs observed in 2 single-arm trials. Two subsequent phase 3 trials have shown superior PFS and tumor response rates and improved quality of life in patients with crizotinib vs chemotherapy, in both previously untreated and untreated ALK-positive advanced NSCLC. FDA has approved 2 companion diagnostics or detecting ALK gene rearrangements to aid in selecting NSCLC patients for treatment with crizotinib.

**BRAF Gene Variants**

**Clinical Validity**

BRAF variants are detected by PCR sequencing or NGS methods. The Oncomine Dx Target Test was FDA-approved in 2017 as a companion diagnostic to detect BRAF V600E variants to aid in selecting NSCLC patients for treatment with combination dabrafenib (Tafinlar) and trametinib (Mekinist) therapy. The Oncomine test is an NGS oncology panel that detects, among other variants, BRAF V600E variants from DNA isolated from FFPE tumor tissue samples.\(^{13}\) The clinical validity of the detection of BRAF V600E variants by the test was evaluated by retrospective analyses of a phase 2, multicenter, nonrandomized study including patients with a BRAF V600E variant who had progressed on prior treatment or were treatment naive who were treated with dabrafenib in combination with trametinib in the study. Patients were screened for a BRAF V600E variant based on local lab tests used at each of the enrollment sites. No FDA-approved test was available for detection of BRAF V600E variants in FFPE NSCLC specimens so a validated PCR assay, BRAF V600 PCR Mutation Test, also was used to estimate concordance. The concordance between the Oncomine test and the BRAF V600 PCR Mutation Test was 100% for PPA (95% CI, 95% to 100%) and 100% for NPA (95% CI, 97% to 100%). The response rate in the 57 previously treated patients in the study that were BRAF-positive by local lab test was 67% (95% CI, 53% to 79%) compared with 73% (95% CI, 50% to 89%) for the 22 patients that were also BRAF-positive by Oncomine. The response rate in the 36 treatment-naïve patients who were BRAF-positive by local lab test was 61% (95% CI, 44% to 77%) compared with 61% (95% CI, 39% to 80%) in the 23 patients who were also BRAF-positive by Oncomine.

**Clinical Utility**

In June 2017, FDA approved an additional indication for use of dabrafenib and trametinib combination therapy in patients with NSCLC with BRAF V600E variant as detected by an FDA-approved test. The Oncomine Dx Target Test was approved as a companion diagnostic. The dabrafenib and trametinib product labels describe the results of an open-label, multicenter study.
of patients enrolled 3 cohorts: cohorts A and B had received at least one previous platinum-based chemotherapy regimen with demonstrated disease progression but no more than 3 prior systemic regimens; cohort C could not have received prior systemic therapy for metastatic disease.\textsuperscript{11,12} Trial results for cohorts A and B have also been published in Planchard (2016)\textsuperscript{79} and Planchard (2016)\textsuperscript{80} and are shown in the table below. Cohort A (n=78) received dabrafenib; cohorts B (n=57) and C (n=36) received dabrafenib and trametinib combination therapy. The response rate in cohort C (treatment-naive, received combination therapy) was 61\% (95\% CI, 44\% to 77\%).

The characteristics and results of key nonrandomized trials of BRAF or MEK inhibitors in NSCLC are described in Tables 8 and 9. In summary, the response rate for dabrafenib monotherapy in 78 patients who had progressed on chemotherapy was 33\% at 11 months median follow-up while the response rate for 19 patients (17 of whom had progressed on chemotherapy) treated with vemurafenib monotherapy was 42\% at 8 weeks. Response rates for dabrafenib and trametinib combination therapy were higher than 60\% in patients who had progressed on prior treatment and those that were treatment-naive. Toxicities were similar to those seen in melanoma patients taking BRAF or MEK inhibitors. SCCs and other dermatological side effects were reported.

### Table 8. Study Characteristics of Key Nonrandomized Trials of BRAF or MEK Inhibitors in BRAF-Variant NSCLC

<table>
<thead>
<tr>
<th>Author (Year)</th>
<th>Study Type</th>
<th>Country</th>
<th>Dates</th>
<th>Participants</th>
<th>Treatment</th>
<th>Median FU, mo</th>
</tr>
</thead>
<tbody>
<tr>
<td>Planchard et al (2016)\textsuperscript{79} NCT01336634</td>
<td>Single-arm, open-label phase 2 trial</td>
<td>9 countries in North America, Europe, Asia</td>
<td>2011-2014</td>
<td>Adults, stage IV, BRAF V600E variant, progression after chemotherapy</td>
<td>Dabrafenib (150 mg bid)</td>
<td>10.7</td>
</tr>
<tr>
<td>Planchard et al (2016)\textsuperscript{80}; NCT01336634</td>
<td>Single-arm, open-label phase 2 trial</td>
<td>9 countries in North America, Europe, Asia</td>
<td>2013-2015</td>
<td>Adults, stage IV, BRAF V600E variant, progression after chemotherapy</td>
<td>Dabrafenib (150 mg bid) plus trametinib (2 mg/d)</td>
<td>11.6</td>
</tr>
<tr>
<td>Hyman et al (2015)\textsuperscript{81}; NCT01524978</td>
<td>Single-arm, open-label phase 2 trial</td>
<td>Germany, Spain, U.K., U.S., France</td>
<td>2012-2014</td>
<td>BrafV600 variant-positive nonmelanoma cancers including NSCLC</td>
<td>Vemurafenib (960 mg bid)</td>
<td>6\textsuperscript{a}</td>
</tr>
</tbody>
</table>

bid: twice a day; FU: follow-up; NSCLC: non-small-cell lung cancer.
\textsuperscript{a} Estimated from a figure.

### Table 9. Results of Key Nonrandomized Trials of BRAF or MEK Inhibitors in BRAF-Variant NSCLC

<table>
<thead>
<tr>
<th>Study (Year)</th>
<th>Response (95% CI), %</th>
<th>PFS (95% CI), mo</th>
<th>Overall Survival</th>
<th>Grade 3 or 4</th>
<th>Serious</th>
</tr>
</thead>
<tbody>
<tr>
<td>Planchard et al (2016)\textsuperscript{79}</td>
<td>33 (23 to 45)\textsuperscript{a}</td>
<td>5.5 (3.4 to 7.3) mo</td>
<td>Median, 12.7 mo</td>
<td>• Overall</td>
<td>• 39</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Cutaneous SCC</td>
<td>• 12</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Astenia</td>
<td>• 5</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• BCC</td>
<td>• 5</td>
</tr>
</tbody>
</table>

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Contains Public Information
Case reports have documented a response to vemurafenib in patients with NSCLC and a BRAF variant.\(^8^2-^8^4\)

**Section Summary: BRAF Gene Variants**

FDA has approved a companion diagnostic for detecting BRAF variants to aid in selecting NSCLC patients for treatment with combination BRAF and MEK inhibitors, dabrafenib and trametinib. The clinical validity of the companion diagnostic was established in the Summary of Safety and Effectiveness Data document. FDA expanded the indication for dabrafenib and trametinib to include the treatment of NSCLC patients whose tumors have a BRAF V600E variant based on a multicenter, single-arm study that included a cohort of 57 patients who had progressed on prior therapy and a cohort of 36 treatment-naive patients. Dabrafenib and trametinib combination therapy was very effective in patients with a BRAF V600E variant with a response rate of about 60% in both cohorts. Lower response rates were reported in other nonrandomized studies of BRAF inhibitor monotherapy in patients who had previously progressed on prior treatments.

**ROS1 Gene Rearrangements**

**Clinical Validity**

Several methods are available to detect ROS1 translocations including FISH, immunohistochemistry, quantitative real-time reverse transcription-PCR (qRT-PCR), and some NGS panels. FISH is considered the standard method. The Oncomine Dx Target Test was FDA-approved in 2017 as a companion diagnostic to detect fusions in ROS1 to aid in selecting NSCLC patients for treatment with crizotinib (XALKORI). The Oncomine test is an NGS oncology panel that detects, among other variants, fusions in ROS1 from RNA isolated from formalin-fixed, paraffin-embedded (FFPE) tumor tissue samples.\(^1^3\) The clinical validity of the detection of ROS1 rearrangements by the test was evaluated by retrospective analysis of FFPE NSCLC specimens obtained from patients enrolled in a ROS1 cohort from an ongoing single-arm, phase 1 safety study of crizotinib in patients with advanced cancer. ROS1 fusion status was determined by a validated FISH comparator test for the study. Concordance between the Oncomine Dx Target Test and the FISH test as well as clinical outcomes were reported in the Summary of Safety and Effectiveness Data. A total of 157 specimens were included. The PPA for Oncomine vs FISH was 80% (95% CI, 59 to 93) and NPA was 100% (95% CI, 97 to 100%). For all ROS1-positive patients, as originally detected for enrollment into the ROS1 cohort, the response rate was 72%.

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### Table: Adverse Events, %

<table>
<thead>
<tr>
<th>Study (Year)</th>
<th>Response (95% CI), %</th>
<th>PFS (95% CI), mo</th>
<th>Overall Survival</th>
<th>Adverse Events, %</th>
</tr>
</thead>
<tbody>
<tr>
<td>Hyman et al (2015)(^a)</td>
<td>N=19</td>
<td>N=20</td>
<td>N=20</td>
<td>N=95(^b)</td>
</tr>
<tr>
<td>At 8 wk, 42 (20 to 67)</td>
<td>Median, 7.3 (3.5 to 10.8)</td>
<td>At 12 mo, 66% (36% to 85%)</td>
<td>• Overall</td>
<td>73</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>• Rash</td>
<td>16</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>• Fatigue</td>
<td>12</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>• Arthralgia</td>
<td>4</td>
</tr>
</tbody>
</table>

AE: adverse events; BCC: basal cell carcinoma; CI: confidence interval; NSCLC: non-small-cell lung cancer; PFS: progression-free survival; SCC: squamous cell carcinoma.

\(^a\) The response rate in the Food and Drug Administration product label for this cohort was 27% (18% to 38%).

\(^b\) Only reported for entire cohort including all cancer types.
(95% CI, 58% to 84%). For *ROS1*-positive patients as detected by Oncomine, the response rate was 83% (95% CI, 36% to 99.6%).

Clinical Utility
In 2016, after an expedited review, the FDA expanded the indication for crizotinib to include the treatment of patients whose metastatic NSCLC tumors have a *ROS1* rearrangement. The approval was based on a 2014 multicenter, single-arm study that enrolled 50 patients with advanced NSCLC who tested positive for *ROS1* rearrangement. The study was an expansion cohort of the phase 1 PROFILE 1001 Trial. Patients were given oral crizotinib (250 mg twice daily) in continuous 28-day cycles; the median duration of treatment was 65 weeks. Characteristics and results of this and other nonrandomized studies are shown in Tables 10 and 11. A companion *ROS1* biomarker diagnostic test was not approved at the time of the crizotinib indication expansion. However, the Oncomine Dx Target Test was FDA-approved in 2017 as a companion diagnostic to detect fusions in *ROS1* to aid in selecting NSCLC patients for treatment with crizotinib (Xalkori).

In summary, a nonrandomized trial and an observational study of crizotinib have shown response rates of greater than 70% in patients with *ROS1* rearrangements, the majority of which had progressed on prior therapy. One nonrandomized trial of ceritinib reported response rates of about 60%. Adverse events were similar to those seen in patients with *ALK* rearrangements using ALK TKIs. Common low-grade side effects include gastrointestinal side effects, visual impairment, and pain. Grade 3 or higher adverse events include liver function abnormalities and pneumonia.

Table 10. Study Characteristics of Key Nonrandomized Trials and Observational Studies of Crizotinib or Ceritinib for *ROS1* Rearrangements in NSCLC

<table>
<thead>
<tr>
<th>Author (Year)</th>
<th>Study Type</th>
<th>Country</th>
<th>Dates</th>
<th>Participants</th>
<th>Treatment</th>
<th>Follow-Up, mo</th>
</tr>
</thead>
<tbody>
<tr>
<td>Shaw et al (2014)</td>
<td>Open-label, single-arm, expansion cohort of phase 1 study</td>
<td>Australia, Korea, U.S.</td>
<td>2010-2013</td>
<td>Adults with <em>ROS1</em> rearrangement, 86% had received prior therapy</td>
<td>Crizotinib (250 mg bid)</td>
<td>16.4</td>
</tr>
<tr>
<td>Mazieres et al (2015)</td>
<td>Retrospective study</td>
<td>Six European countries</td>
<td>NR</td>
<td><em>ROS1</em> rearrangement, 97% had received previous chemotherapy</td>
<td>Crizotinib (250 mg bid)</td>
<td>NR</td>
</tr>
<tr>
<td>Lim et al (2017)</td>
<td>Open-label, single-arm, phase 2 study</td>
<td>Korea</td>
<td>2013-2016</td>
<td>Adults with <em>ROS1</em> rearrangement who had progressed on prior therapy, 94% crizotinib-naïve</td>
<td>Ceritinib (750 mg/d)</td>
<td>14.0</td>
</tr>
</tbody>
</table>

bid: twice a day; NR: not reported; NSCLC: non-small-cell lung cancer.

Table 11. Results of Key Nonrandomized Trials and Observational Studies of Crizotinib or Ceritinib for *ROS1* Rearrangements in NSCLC

<table>
<thead>
<tr>
<th>Study (Year)</th>
<th>Response (95% CI), %</th>
<th>Median PFS (95% CI), mo</th>
<th>OS (95% CI), mo</th>
<th>Grade 3 or 4 %</th>
<th>All Grades %</th>
</tr>
</thead>
<tbody>
<tr>
<td>Shaw et al (2014)</td>
<td>72 (58 to 84)</td>
<td>19.2 mo (14.4 to not reached)</td>
<td>At 12 mo: 85% (72% to 93%)</td>
<td>• Hypophosphatemia • Neutropenia</td>
<td>10 • 4</td>
</tr>
<tr>
<td>• Visual impairment • Diarrhea</td>
<td>10 • 4</td>
<td>82 • 40</td>
<td>44</td>
<td>40</td>
<td>44</td>
</tr>
</tbody>
</table>
Kim et al (2013) reported on clinical outcomes in 208 never-smokers with NSCLC adenocarcinoma, according to ROS1-rearrangement status. ALK rearrangements and EGFR variants were concurrently analyzed. The patients had clinical stages ranging from I to IV, but most were stage IV (41.3%). Of the 208 tumors, 3.4% (n=7) were ROS1 rearranged. ROS1 rearrangement was mutually exclusive from ALK rearrangement, but 1 of 7 ROS1-positive patients had a concurrent EGFR variant. Patients with ROS1 rearrangement had a higher ORR and longer median PFS on pemetrexed than those without a rearrangement. In patients with ROS1 rearrangement, PFS with EGFR TKIs was shorter than those patients without the rearrangement. None of the ROS1-positive patients received ALK inhibitors (eg, crizotinib), which is the recommended targeted therapy for patients with NSCLC and this genetic alteration.

**Section Summary: ROS1 Gene Rearrangements**

FDA has approved a companion diagnostic for detecting ROS1 gene rearrangements to aid in selecting NSCLC patients for treatment with crizotinib (Xalkori). The clinical validity of the companion diagnostic was established in the Summary of Safety and Effectiveness Data document. FDA expanded the indication for crizotinib to include the treatment of patients whose tumors have a ROS1 rearrangement based on a multicenter, single-arm study including 50 patients, the majority of whom had progressed on prior therapy. Crizotinib was very effective in patients with ROS1 rearrangements with a response rate of about 70%. Similar response rates were reported in other nonrandomized studies of crizotinib and ceritinib.

**KRAS Gene Variants**

**Clinical Validity**

KRAS variants can be detected by direct sequencing, PCR technologies or NGS. Although KRAS is the most common driver mutation in NSCLC, there are currently no targeted therapies specifically approved for patients with NSCLC with KRAS variants and therefore, no FDA-approved companion diagnostics.
Clinical Utility

**KRAS Variants and EGFR TKIs**

Data on the role of KRAS variants in NSCLC and response to erlotinib are available from post hoc analyses of phase 3 trials of TKIs in patients with wild-type (nonmutated) vs KRAS-mutated lung tumors; phase 2 trials; a large prospective study; retrospective single-arm studies; and meta-analyses.

**Systematic Reviews:** Meta-analyses on the relation between KRAS variants and response to EGFR-TKI therapy are outlined next. The pooled data were insufficient to determine an association between KRAS variant status and treatment effects on PFS or OS.

Linardou et al (2008) performed a meta-analysis of 17 studies with 1008 patients, 165 (16.4%) of whom had a KRAS variant.89 Eligible studies reported response (complete or partial) stratified by KRAS variant status. Primary end points were sensitivity and specificity of KRAS testing, defined as KRAS variant carriers showing no response to erlotinib (stable disease or progressive disease) and KRAS wild-type patients showing a response, respectively. Sensitivity and specificity were assessed overall and in subgroups defined by TKI received (gefitinib and/or erlotinib), response criteria (Response Evaluation Criteria in Solid Tumors [RECIST] or World Health Organization), possible selection bias, and previous chemotherapy, if any. There was no significant difference in sensitivity or specificity across subgroups. The presence of a KRAS variant was associated with a lack of response to TKIs (sensitivity, 21%; 95% CI, 16% to 28%; specificity, 94%; 95% CI, 89% to 97%; positive likelihood ratio, 3.52; negative likelihood ratio, 0.84). (For the analysis, likelihood ratios were calculated using pooled estimates for sensitivity and specificity.) Reviewers concluded that KRAS variants conferred a high level of resistance to anti-EGFR therapies; however, this conclusion is tentative due to study limitations (eg, lack of individual patient data, heterogeneity of response end points, treatment regimens, patient selection criteria, retrospective design of included studies). Furthermore, incomplete reporting of survival data precluded meaningful assessment of the effect of KRAS variant on survival.

Mao et al (2010) performed a meta-analysis of 22 studies in 1470 patients with NSCLC (1335 [91%] evaluable for response), 231 (17%) of whom had KRAS variants.90 Studies were heterogeneous in patient populations (smoking history, tumor histology, stage, ethnicity, treatment received) and response criteria. The primary end point was ORR, defined as the sum of complete and partial response. ORRs for patients with mutated KRAS and wild-type KRAS were 3% and 26%, respectively. Incomplete reporting of survival data precluded meaningful assessment of the effect of KRAS status on survival in NSCLC patients treated with EGFR TKIs. Data for PFS and OS stratified by KRAS status were available in 8 studies. The median PFS in KRAS-mutated and wild-type patients was 3.0 months and 3.9 months, respectively. The median OS in KRAS-mutated and wild-type patients was 4.7 months and 10.7 months, respectively. However, only 2 studies presented HRs with 95% CIs for PFS and OS and, therefore, a pooled analysis to derive an overall HR was not performed.

Pan et al (2016) published a meta-analysis of 41 studies (total N=13,103 patients) of prognostic and predictive values of the KRAS variant in NSCLC.91 The KRAS variant was significantly associated with poorer OS (HR=1.6; 95% CI, 1.4 to 1.8) and DFS (HR=1.57; 95% CI, 1.2 to 2.1) in early-stage resected NSCLC, and with inferior outcomes of EGFR-TKIs treatment (RR=0.21; 95% CI, 0.1 to 0.4) in advanced NSCLC. The KRAS variant was still significantly associated with
poorer OS (HR=1.4; 95 % CI, 1.2 to 1.6) and PFS (HR=1.4; 95 % CI, 1.1 to 1.6) of EGFR TKIs when patients with EGFR variants were excluded.

Retrospective Analyses of Trials: Pao et al (2005) were the first to suggest that patients with KRAS-mutated lung tumors were nonresponsive to treatment with EGFR TKIs.92 Thirty-six patients with bronchioloalveolar carcinoma underwent KRAS variant analysis; 9 (25%) were found to harbor KRAS variants. The response was by a single radiologist, blinded to patient outcome, using RECIST criteria. None of 9 patients with KRAS-mutated tumors responded to erlotinib (p=0.553).

Eberhard et al (2005) performed a post hoc subgroup analysis of KRAS variants in previously untreated patients with advanced NSCLC who had been randomized to chemotherapy with or without erlotinib.17 The original phase 3 trial (TRIBUTE) randomly assigned patients to carboplatin plus paclitaxel either with or without erlotinib.93 Of the original 1079 patients, tumor DNA from 274 (25%) patients was sequenced for KRAS variants. Baseline demographics between patients with available tumor DNA and those without were balanced. KRAS variants were detected in 55 (21%) of 274 patients. The response rate for patients with wild-type KRAS was 26%, regardless of treatment received. In patients with KRAS-mutated tumors, the response rate was 8% for those receiving chemotherapy with erlotinib and 23% for those receiving chemotherapy alone (p=0.16; 95% CI for difference, -5% to 35%); the median OS was 4.4 months (95% CI, 3.4 to 12.9 months) in patients who received erlotinib and 13.5 months (95% CI, 11.1 to 15.9 months) in those who received chemotherapy alone (p=0.019).

In a 2006 phase 2 trial, Giaccone et al studied the response to erlotinib in 53 chemotherapy-naive patients with advanced NSCLC.44 Histologic material was available to assess KRAS variant status from 29 patients, 10 (34%) of whom had variants. All 10 were nonresponders to erlotinib (p=0.125).

In a 2007 phase 2, multicenter, open-label study, Jackman et al evaluated treatment response to erlotinib in chemotherapy-naive patients 70 years of age or older who had advanced NSCLC.45 Of 80 patients eligible for treatment, 41 (51%) had KRAS variant analysis; 6 (15%) patients were variant-positive, none of whom responded to erlotinib. Five (14%) of 35 patients with wild-type KRAS had a partial response.

In a 2008 phase 2 trial, Miller et al assessed response to erlotinib in 101 patients with lung bronchioloalveolar carcinoma (n=12) or adenocarcinoma, bronchioloalveolar subtype (n=89), according to KRAS variant status.42 Eighteen (18%) patients had KRAS-mutated tumors, and none responded to erlotinib (95% CI, 0% to 19%; p<0.01). In patients without a KRAS variant, the response rate was 32%. The median OS in patients with KRAS-mutated tumor was 13 months vs 21 months in patients with KRAS wild-type tumor (p=0.30).

Zhu et al (2008) performed a post hoc subgroup analysis of KRAS variants in patients with advanced NSCLC who had failed standard chemotherapy and had been previously randomized to erlotinib or placebo.46 The original phase 3 trial (National Cancer Institute of Canada Clinical Trials Group Study BR.21; 2005) was the first to demonstrate a significant survival advantage with the use of an EGFR TKI in previously treated NSCLC patients.94 In post hoc analysis, 206 (28%) of the original 731 tumors were tested for KRAS variants, which were identified in 30 (15%) patients. Among the 206 tested patients, 118 (57%) were assessable for response to
erlotinib. Of 98 patients with wild-type KRAS, 10 (10.2%) responded to erlotinib; of 20 patients with mutated KRAS, 1 (5.0%) patient responded (HR [erlotinib vs placebo] in patients with mutated KRAS, 1.67; 95% CI, 0.62 to 4.50; p=0.31; HR in wild-type patients, 0.69; 95% CI, 0.49 to 0.97; p=0.03). In Cox regression, the interaction between KRAS variant status and treatment was not statistically significant (p=0.09).

Rulli et al (2015) reported on results from biomarker analyses in the TAILOR trial. TAILOR enrolled patients from 52 Italian hospitals and genotyped patients for KRAS and EGFR variant status. Wild-type EGFR patients (n=218) received first-line platinum-based chemotherapy and then were randomly allocated at progression to erlotinib or docetaxel. KRAS variants were present in 23% of randomized patients. The presence of a KRAS variant was not associated with PFS (HR=1.01; 95% CI, 0.71 to 1.41; p=0.98) or OS (HR=1.24; 95% CI, 0.87 to 1.77; p=0.23). The treatment effect did not differ by KRAS status (test for interaction: OS p=0.97; PFS p=0.42).

Papadimitrakopoulou et al reported on results of the BATTLE-2 phase 2 study in 2016. The BATTLE-2 program is an umbrella study evaluating effects of targeted therapies focusing on KRAS-mutated cancers. Two hundred patients with advanced NSCLC tumors who did not have EGFR variants or ALK gene fusions whose cancer was refractory to more than 1 prior therapy were assigned to 1 of 4 arms using adaptive randomization: erlotinib (n=22), erlotinib plus MK-2206 (n=42), MK-2206 plus AZD6244 (n=75), or sorafenib (n=61), stratified by KRAS status. AZD6244 and MK2206 are targeted small-molecule drugs that inhibit MEK and AKT, respectively. Sorafenib is a multitargeted signal transduction inhibitor that inhibits raf-kinases, vascular endothelial growth factor receptor 2, platelet-derived growth factor receptor-B, and c-kit. Only 186 evaluable patients were included in analyses. The 8-week disease control rate was 20%, 25%, 62%, and 44% for the 4 treatment groups, respectively, in the KRAS variant− positive patients. For KRAS wild-type patients, disease control rate was 36%, 57%, 49%, and 47%, respectively. The median PFS did not differ by KRAS status.

Observational Studies: Schneider et al (2008) reported on the relation between clinical benefit and putative tumor markers in a subgroup of patients participating in a global open-label, single-arm study of erlotinib in advanced NSCLC, involving 7043 patients in 52 countries (the TRUST study). The subgroup was from German centers and comprised 311 patients with stage IIIB or IV disease who were treated with erlotinib because they had failed or were not medically suitable for standard first-line chemotherapy. Tumor response was assessed using RECIST. Seventeen (15%) patients had KRAS variants, and none responded to erlotinib; 2 patients had stable disease. The impact of KRAS variant status on OS (p=0.06) and PFS (p not reported) was of borderline statistical significance. The authors concluded that current data did not support the selection of patients for treatment with erlotinib on the basis of tumor molecular characteristics and that further studies were needed to determine definitively whether patients with KRAS variants can derive a survival benefit from erlotinib.

In 2009, Boldrini et al reported on the association between KRAS and EGFR variant status and several clinical variables in 411 patients with lung adenocarcinoma, and presented a subgroup analysis of tumor response in patients treated with erlotinib or gefitinib. KRAS variants were observed in 17.9% of all patients. The subset analysis comprised 21 women with stage IV disease who received a TKI as second- or third-line therapy and were assessed for radiographic tumor response using RECIST. The mean age of this subpopulation at the time of diagnosis was 60.8 years (range, 40-86 years). Nineteen (90%) of 21 women were KRAS wild-type, and of
those, 8 (42%) showed partial response, 4 (21%) had stable disease, and 7 (37%) had progressive disease. Two patients with KRAS variants had progressive disease.

Guan et al (2013) reported on 1935 consecutive patients with NSCLC who were treated at a single institution in China. Patients with mutated KRAS were randomly matched on tumor, node, metastasis (TNM) stage, time of first visit within 1 year, and histology, to both EGFR variant–positive and KRAS/EGFR wild-type patients. Seventy (4%) patients received EGFR-TKI therapy. In this group, median PFS was 11.8 months and 2.0 months in patients with EGFR and KRAS variants, respectively, and 1.9 months in wild-type patients; compared with wild-type patients, PFS was statistically longer in patients with EGFR variants (p<0.001) but no different in patients with KRAS variants (p=0.48). The authors observed that “the presence of an EGFR variant, but not a KRAS variant, was predictive of responsiveness to EGFR TKI treatment.”

Fiala et al (2013) reported on a retrospective analysis of patients with squamous cell NSCLC who underwent EGFR, KRAS, and PIK3CA (phosphatidylinositide-3-kinase catalytic subunit-alpha) variant testing. Of 215 patients tested, 16 (7.4%) had mutated KRAS. Of 174 tested patients who were treated with an EGFR-TKI (erlotinib or gefitinib), median PFS in 14 KRAS-mutated patients was 1.3 months vs 2.0 months in KRAS wild-type patients (n=160 [92%]); the difference was not statistically significant (p=0.120). Median OS in this treated group was 5.7 months in KRAS-mutated patients vs 8.2 months in KRAS wild-type patients, a statistically significant difference (p=0.039). The authors concluded that KRAS variant status may have a negative prognostic role but a predictive role was not confirmed. “Patients with squamous cell NSCLC harboring these variants could benefit from targeted treatment and should not be excluded from treatment with EGFR TKIs.”

**KRAS and Anti-EGFR Monoclonal Antibodies**

Two phase 3 trials (BMS099, FLEX) investigated platinum-based chemotherapy with and without cetuximab in the first-line setting for advanced NSCLC. Subsequently, investigations of KRAS variant status and cetuximab treatment were performed for both trials.

In the multicenter, phase 3 BMS099 trial (2010), 676 chemotherapy-naive patients with stage IIIB or IV NSCLC were assigned to taxane and carboplatin with or without cetuximab. The primary end point was PFS; secondary end points were overall response rate, OS, QOL, and safety. The addition of cetuximab did not significantly improve PFS; however, there was a statistically significant improvement in overall response rate in the cetuximab group. There was a trend in OS favoring cetuximab; however, it was not statistically significant. A post hoc correlative analysis was conducted to identify molecular markers for the selection of patients most likely to benefit from cetuximab. Of the original 676 enrolled patients, 202 (29.9%) had tumor samples available for KRAS testing. KRAS variants were present in 35 (17%) patients. Among patients with wild-type KRAS, OS was similar between the cetuximab-containing arm (n=85) and the chemotherapy-alone arm (n=82) (HR=0.93; 95% CI, 0.67 to 1.30; p=0.68; median survival, 9.7 months and 9.9 months, respectively). Among patients with KRAS variants, OS was similar between the cetuximab-containing arm (n=13) and the chemotherapy-alone arm (n=22) (HR=0.91; 95% CI, 0.45 to 2.07; p=0.93; median survival, 16.8 months and 10.8 months, respectively). Overall, the study showed no significant treatment-specific interactions between the presence of KRAS variants and outcomes evaluated; treatment differences favoring the addition of cetuximab in the KRAS-mutated subgroup were consistent with those observed in the wild-type KRAS subgroup and in the overall study population. The authors concluded that the
results did not support an association between KRAS variant and lack of cetuximab benefit similar to that observed in patients with KRAS-mutated metastatic colorectal cancer. However, the results should be interpreted with caution due to small subgroup sample sizes and retrospective nature of the analysis.

In the open-label, randomized, phase 3 FLEX trial (2009), 1125 chemotherapy-naive patients with stage III or IV, NSCLC were randomized to chemotherapy (cisplatin and vinorelbine) plus cetuximab (n=557) or chemotherapy alone (n=568). The primary end point was OS. Patients who received chemotherapy plus cetuximab survived longer than those who received chemotherapy only (median OS, 11.3 months vs 10.1 months, respectively; HR for death, 0.87; 95% CI, 0.76 to 1.00; p=0.04). Subsequently, KRAS variant testing was performed on archival tumor tissue of 395 (35%) of 1125 patients. KRAS variants were detected in 75 (19%) tumors. Among patients with mutated KRAS, the median OS in the cetuximab-containing (n=38) and chemotherapy-alone arms (n=37) was similar (8.9 months vs 11.1 months, respectively; HR=1.00; 95% CI, 0.60 to 1.66; p=1.0). Among patients with wild-type KRAS, the median OS in the cetuximab-containing (n=161) and chemotherapy-alone arms (n=159) was similar (11.4 months vs 10.3 months, respectively; HR=0.96; 95% CI, 0.75 to 1.23; p=0.74). PFS also was similar in the cetuximab-containing and chemotherapy-alone arms in patients with mutated (HR=0.97; 95% CI, 0.76 to 1.24) and wild-type (HR=0.84; 95% CI, 0.50 to 1.40) KRAS. Response rates in the cetuximab-containing arm in patients with KRAS-mutated and wild-type tumors were 36.8% and 37.3%, respectively (p=0.96). Overall, there was no indication that KRAS variant status was predictive of cetuximab effect in NSCLC.

**MEK Inhibitors**

Two RCTs have compared a MEK inhibitor (with or without chemotherapy) with chemotherapy alone in patients with KRAS-variant advanced NSCLC after progression with first-line therapy. The characteristics and results are shown in Tables 12 and 13. MEK inhibitor therapy did not improve PFS compared with docetaxel alone; response rates were similar or marginally improved. Grade 3 or higher adverse events were more frequent with MEK inhibitor therapy compared with docetaxel.

**Table 12. RCT Characteristics of MEK Inhibitors for KRAS-Variant NSCLC**

<table>
<thead>
<tr>
<th>Author (Year); Study</th>
<th>Countries</th>
<th>Sites</th>
<th>Dates</th>
<th>Participants</th>
<th>MEK Inhibitor</th>
<th>Chemotherapy</th>
</tr>
</thead>
<tbody>
<tr>
<td>Blumenschein (2015)\cite{104}; NCT01362296</td>
<td>U.S., Korea, 6 European countries</td>
<td>60</td>
<td>2011-2012</td>
<td>129 patients with stage IV NSCLC and progression following first-line platinum-containing chemotherapy</td>
<td>86 assigned to trametinib (2 mg/d)</td>
<td>43 assigned to docetaxel (75 mg/m²)</td>
</tr>
<tr>
<td>Janne (2017)\cite{105}; SELECT1 (NCT01933932)</td>
<td>25 countries in North and South America, Australia, Europe</td>
<td>202</td>
<td>2013-2016</td>
<td>510 patients with advanced NSCLC and progression following first-line therapy</td>
<td>254 assigned to selumetinib (75 mg bid) plus docetaxel (75 mg/m²)</td>
<td>256 assigned to docetaxel (75 mg/m²)</td>
</tr>
</tbody>
</table>

bid: twice a day; NSCLC: non-small-cell lung cancer; RCT: randomized controlled trial.
Table 13. RCT Results for MEK Inhibitors for *KRAS*-Variant NSCLC

<table>
<thead>
<tr>
<th>Study (Year)</th>
<th>Progression-Free Survival</th>
<th>Overall Survival</th>
<th>ORR (95% CI), %</th>
<th>Grade ≥3 %</th>
<th>Adverse Events, %</th>
</tr>
</thead>
<tbody>
<tr>
<td>Trametinib</td>
<td>12 wk</td>
<td>8 mo</td>
<td>12</td>
<td>• Overall</td>
<td>41</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Rash</td>
<td>6</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Diarrhea</td>
<td>5</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Asthenia</td>
<td>5</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Hypertension</td>
<td>9</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Neutropenia</td>
<td>0</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Decreased neutrophils</td>
<td>0</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Rash</td>
<td>6</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Diarrhea</td>
<td>5</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Asthenia</td>
<td>5</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Hypertension</td>
<td>9</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Neutropenia</td>
<td>0</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Decreased neutrophils</td>
<td>0</td>
</tr>
<tr>
<td>Docetaxel</td>
<td>11 wk</td>
<td>Not yet reached</td>
<td>12</td>
<td>• Overall</td>
<td>37</td>
</tr>
<tr>
<td></td>
<td></td>
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<td></td>
<td>• Rash</td>
<td>0</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Diarrhea</td>
<td>2</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Asthenia</td>
<td>0</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Hypertension</td>
<td>0</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Neutropenia</td>
<td>14</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Decreased neutrophils</td>
<td>7</td>
</tr>
<tr>
<td>TE (95% CI)</td>
<td>HR=1.14 (0.75 to 1.75)</td>
<td>HR=0.97 (0.52 to 1.83)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Selumetinib plus docetaxel</td>
<td>3.9 mo</td>
<td>8.7 mo</td>
<td>20.1</td>
<td>• Overall</td>
<td>67</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Rash</td>
<td>7</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Diarrhea</td>
<td>9</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Asthenia</td>
<td>8</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Dyspnea</td>
<td>5</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Anemia</td>
<td>7</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Neutropenia</td>
<td>7</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Rash</td>
<td>3</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Diarrhea</td>
<td>3</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Asthenia</td>
<td>2</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Dyspnea</td>
<td>4</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Anemia</td>
<td>4</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>• Neutropenia</td>
<td>4</td>
</tr>
<tr>
<td>TE (95% CI)</td>
<td>HR=0.93 (0.77 to 1.12)</td>
<td>HR=1.05 (0.85 to 1.30)</td>
<td>OR=1.61 (1.00 to 2.62)</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

CI: confidence interval; HR: hazard ratio; NSCLC: non-small-cell lung cancer; OR: odds ratio; ORR: objective response rate; RCT: randomized controlled trial; TE: treatment effect.

Section Summary: *KRAS* Gene Variants

Data on the role of *KRAS* variants in NSCLC and response to erlotinib are available from post hoc analysis of trials, observational studies, and meta-analyses. Although studies have shown that *KRAS*-variants in patients with NSCLC confer a high level of resistance to TKIs, data are insufficient to assess any additional benefit to *KRAS* testing beyond *EGFR* testing.

A lack of response to EGFR monoclonal antibodies has been established in metastatic colorectal cancer, and use of these drugs is largely restricted to patients with wild-type *KRAS*. The expectation that *KRAS* variant status also would be an important predictive marker for cetuximab response in NSCLC has not been shown. In 2 randomized trials with post hoc analyses of *KRAS* variant status and use of cetuximab with chemotherapy, *KRAS* variants did not identify patients...
who would benefit from anti-EGFR antibodies, because outcomes with cetuximab were similar regardless of KRAS variant status.

Two RCTs have compared a MEK inhibitor with docetaxel in patients with KRAS-variant advanced NSCLC who had progression following first-line therapy. The MEK inhibitor did not improve PFS compared with docetaxel; the response rate was marginally improved. Grade 3 or higher adverse events were more frequent with the MEK inhibitors.

**HER2 Gene Variants**

Mazières et al (2013) reported on a retrospective review of a consecutive series of patients with NSCLC tested for a HER2 variant, and they assessed clinicopathologic characteristics and patient outcomes by variant status. A HER2 variant was identified in 65 (1.7%) of 3800 patients, and was mutually exclusive of other driver variant (EGFR, ALK, BRAF), with the exception of 1 case in which both an HER2 and KRAS variant were identified. The patient population in which a HER2 variant was found had a median age of 60 years (range, 31-86 years), 69% were women, and 52% were never-smokers. All tumors were adenocarcinomas, and 50% were stage IV (n=33). Patients with stage IV disease received conventional chemotherapy and, of these, 16 patients also received HER2-targeted therapy as additional lines of therapy (for a total of 22 individual anti-HER2 treatments that were evaluable). Four patients had progressive disease, 7 had disease stabilization, and 11 with partial response. PFS for patients with HER2 therapies was 5.1 months.

Mok et al reported on the biomarker subgroup analyses from the FASTACT-2 study in 2016. FASTACT-2 is a multicenter, randomized, placebo-controlled, double-blind, phase 3 study of intercalated first-line erlotinib or placebo with gemcitabine and platinum, followed by maintenance therapy with erlotinib or placebo, for Asian patients with stage IIIIB or IV NSCLC. In addition to analyzing for EGFR, HER2 and HER3 biomarkers were analyzed by immunohistochemistry. Only EGFR variants (p<0.001) were predictive of outcomes; HER2 and HER3 biomarkers were not significant in a treatment-by-biomarker interaction test.

Shen et al (2015) retrospectively reviewed 111 patients from a Uygur population who received gefitinib 250 mg once daily and were evaluated for HER2 expression. HER2 overexpression was detected in 24 patients. The ORR in patients with and without HER2 overexpression was 29% and 14%, respectively (p=0.12). The median PFS and OS in patients with and without HER2 overexpression did not differ statistically significantly (PFS, 4.7 months vs 3.9 months, p=0.09; OS, 21 months vs 19 months, p=0.09).

**RET Gene Rearrangements**

In a phase 2, prospective trial (2013) for patients with RET-fusion–positive tumors, preliminary data on 3 patients treated with cabozantinib showed a partial response in 2 patients and stable disease in the third, approaching 8 months.

**MET Gene Amplifications**

A phase 2 trial (2013) of MET-positive NSCLC, in which patients were treated with an anti-MET antibody plus erlotinib, showed improved PFS and OS.

**PD-1 / PD-L1**

As a primary immunosuppressive driver, PD-L1 overexpression may be an important facilitator for tumor growth and metastasis. PD-L1 has been detected in up to 50% of human cancers, making
the PD-L1 pathway a focus of cancer research. NCCN recommends IHC testing for PD-L1 expression before first-line treatment in patients with metastatic NSCLC with negative or unknown test results for EGFR mutations, ALK rearrangements, and ROS1 rearrangements. Although it is not an optimal biomarker, PD-L1 expression is currently the best available biomarker to assess whether patients are candidates for pembrolizumab. PD-L1 expression is continuously variable and dynamic; thus, a cutoff value for a positive result is artificial. Patients with PD-L1 expression levels just below and just above 50% will probably have similar responses. The definition of a positive PD-L1 test result varies depending on which biomarker assay is used.

SUMMARY OF EVIDENCE

For individuals who have advanced-stage NSCLC who are being considered for targeted therapy who receive testing for EGFR variants and ALK rearrangements, the evidence includes phase 3 studies comparing TKIs with chemotherapy. Relevant outcomes are overall survival, disease-specific survival, test accuracy and validity, quality of life, and treatment-related morbidity. Studies have shown that TKIs are superior to chemotherapy regarding tumor response rate and progression-free survival, with a reduction in toxicity and improvement in quality of life. The evidence is sufficient to determine that the technology results in a meaningful improvement in the net health outcome.

For individuals who have advanced-stage NSCLC who are being considered for targeted therapy who receive testing for BRAF variants and ROS1 rearrangements, the evidence includes nonrandomized trials and observational studies of BRAF and MEK inhibitors and crizotinib or ceritinib, respectively. Relevant outcomes are overall survival, disease-specific survival, test accuracy and validity, quality of life, and treatment-related morbidity. Studies have shown that combination therapy with dabrafenib and trametinib for BRAF V600E variant NSCLC and crizotinib for NSCLC with ROS1 rearrangements result in response rates of 60% and 70%, respectively, with acceptable toxicity profiles. The evidence is sufficient to determine that the technology results in a meaningful improvement in the net health outcome.

For individuals who have advanced-stage NSCLC who are being considered for targeted therapy who receive testing for KRAS or HER2 variants, RET rearrangements, or MET amplifications, the evidence includes for KRAS post hoc analyses trials, observational studies, and meta-analyses; for the other variants, the evidence includes a phase 2 trial with preliminary data, and retrospective analyses of very small case series and case reports. Relevant outcomes are overall survival, disease-specific survival, test accuracy and validity, quality of life, and treatment-related morbidity. Studies have shown that KRAS variants in patients with NSCLC confer a high level of resistance to TKIs; data are insufficient to assess any additional benefit to testing for KRAS variants to select for EGFR TKIs beyond EGFR testing. In 2 randomized trials with post hoc analyses of KRAS variant status and use of the anti-EGFR monoclonal antibody cetuximab with chemotherapy, KRAS variants did not identify patients who would benefit from anti-EGFR antibodies, because outcomes with cetuximab were similar regardless of KRAS variant status. In two randomized controlled trials of advanced KRAS-variant positive disease, MEK inhibitors did not improve progression-free survival compared with docetaxel. Studies for HER2, RET, and MET variant testing have reported response rates and progression-free survival in numbers of patients too small from which to draw conclusions. The evidence is insufficient to determine the effects of the technology on health outcomes.
PRACTICE GUIDELINES AND POSITION STATEMENTS

National Comprehensive Cancer Network Guidelines

EGFR Testing

National Comprehensive Cancer Network (NCCN) guidelines (v.8.2017) for the treatment of metastatic non-small-cell lung cancer (NSCLC) recommend the following:

- **EGFR [epidermal growth factor receptor] mutation testing** is recommended (category 1) in patients with non-squamous NSCLC (ie, adenocarcinoma, large cell carcinoma) or in NSCLC not otherwise specified, because erlotinib or afatinib (category 1 for both) is recommended for patients who are positive for **EGFR** mutations.
- When an **EGFR** mutation is discovered prior to first-line chemotherapy, erlotinib (category 1), afatinib (category 1), or gefitinib (category 1) are recommended.
- When an **EGFR** mutation is discovered during first-line chemotherapy, interrupt or continue chemotherapy, then follow with erlotinib, afatinib, or gefitinib.
- If progression occurs following first-line treatment, **EGFR T790M** testing is recommended (category 2A). If T790M+, osimertinib (category 1), local therapy, or continuing with erlotinib, afatinib, or gefitinib are recommended (depending on symptoms, location of metastases, and number of lesions).
- Erlotinib is not recommended as first-line therapy or subsequent therapy following progression to for patients negative for **EGFR** mutations or with unknown **EGFR** status.
- In patients with squamous cell carcinoma, **EGFR** mutation testing should be considered in never-smokers; when histology is assessed using small biopsy specimens (rather than surgically resected samples); or when histology is mixed adenosquamous (category 2A).

ALK Gene

NCCN guidelines (v.8.2017) state the following on anaplastic lymphoma kinase (**ALK**) rearrangement testing:

- **ALK**-rearrangement testing is recommended (category 1) in patients with nonsquamous NSCLC (ie, adenocarcinoma, large cell carcinoma) or in NSCLC not otherwise specified.
- If **ALK**-positive status is discovered before first-line chemotherapy, alectinib (category 1; preferred), crizotinib (category 1), or ceritinib (category 1) is recommended.
- If **ALK** rearrangement is discovered during first-line chemotherapy, interrupt or complete planned chemotherapy and start alectinib, crizotinib or ceritinib.
- If there is progression on first-line therapy, continue alectinib, crizotinib or ceritinib, switch to ceritinib, alectinib or brigatinib, or consider local therapies are recommended (depending on symptoms, location of metastases, and number of lesions).
- In patients with squamous cell carcinoma, **ALK**-rearrangement testing should be considered in never-smokers; when histology is assessed using small biopsy specimens (rather than surgically resected samples); or when histology is mixed adenosquamous (category 2A).

ROS1 Testing

NCCN guidelines (v.8.2017) state the following on **ROS1**-rearrangement testing:

- **ROS1**-rearrangement testing is recommended (category 2A) in patients with nonsquamous NSCLC (ie, adenocarcinoma, large cell carcinoma) or in NSCLC not otherwise specified.
- If **ROS1**-positive status is discovered, crizotinib is recommended.
BRAF Testing
NCCN guidelines (v.8.2017) state the following on BRAF testing:
- BRAF testing is recommended (category 2A) in patients with nonsquamous NSCLC (i.e., adenocarcinoma, large cell carcinoma) or in NSCLC not otherwise specified.
- If BRAF V600E mutation-positive status is discovered, combination dabrafenib and trametinib or other first-line options are recommended.

KRAS Gene
NCCN guidelines (v.8.2017) state that “KRAS mutations are associated with intrinsic TKI [tyrosine kinase inhibitor] resistance, and KRAS gene sequencing could be useful for the selection of patients as candidates for TKI therapy.” Targeted therapy for patients with the KRAS variants is currently unavailable.

Other Genes
NCCN guidelines (v.8.2017) do not give specific recommendations for testing for genetic alterations in the genes HER2, RET, or MET in NSCLC. However, it states that the following emerging targeted agents are available for patients with one of these specific genetic alterations:
- High-level MET amplification or MET exon 14 skipping mutation: crizotinib (category 2A)
- HER2 mutations: trastuzumab and afatinib (category 2B)
- RET rearrangements: cabozantinib or vandetanib (category 2A)

College of American Pathologists et al
In 2013, the College of American Pathologists, the International Association for the Study of Lung Cancer, and the Association for Molecular Pathology published evidence-based guidelines for molecular testing to select patients with lung cancer for treatment with EGFR TKI therapy. Based on excellent quality evidence (category A), the guidelines recommend EGFR mutation testing in patients with lung adenocarcinoma regardless of clinical characteristics, such as smoking history.

American Society of Clinical Oncology
In 2014, the American Society of Clinical Oncology reviewed and endorsed the 2013 College of American Pathologists, International Association for the Study of Lung Cancer, and Association for Molecular Pathology guidelines, and highlighted 3 evolving areas: advances in ALK testing methodology, considerations for selecting appropriate populations for molecular testing, and the emergence of other targeted molecular alterations. The American Society of Clinical Oncology recommendations stated that testing for EGFR should be prioritized over other molecular markers in lung adenocarcinoma, and that, after EGFR testing, testing for ALK should be prioritized over other proposed molecular markers in lung adenocarcinomas, for which published evidence is insufficient to support testing guideline development at the present time.

The American Society of Clinical Oncology also updated its evidence-based recommendations on systemic therapy for patients with stage IV NSCLC in 2017. The following are recommendations and associated quality and strength of evidence (see Table 14).

<table>
<thead>
<tr>
<th>Recommendation</th>
<th>QOE</th>
<th>SOR</th>
</tr>
</thead>
<tbody>
<tr>
<td>First-line therapy</td>
<td>High</td>
<td>Strong</td>
</tr>
</tbody>
</table>

Sensitizing EGFR variants: afatinib, erlotinib, or gefitinib
**Recommendation**

| ALK rearrangements: crizotinib | Intermediate | Moderate |
| ROS1 rearrangement: crizotinib | Low | Weak |
| **Second-line therapy** | | |
| Sensitizing EGFR variants and T790M resistance variant: osimertinib | High | Strong |
| ROS1 rearrangement who have not received prior crizotinib: crizotinib | Low | Moderate |
| BRAF variants who have received prior immune checkpoint therapy: dabrafenib alone or in combination with trametinib | Insufficient | Moderate |

QOE: quality of evidence; SOR: strength of recommendation.

**American College of Chest Physicians Guidelines**

American College of Chest Physicians (ACCP) updated its evidence-based clinical practice guidelines on the treatment of stage IV NSCLC in 2013. Based on review of the literature, guideline authors reported improved response rates, progression-free survival, and toxicity profiles with first-line erlotinib or gefitinib compared with first-line platinum-based therapy in patients with EGFR mutations, especially exon 19 deletion and L858R. ACCP recommended “testing patients with NSCLC for EGFR mutations at the time of diagnosis whenever feasible, and treating with first-line EGFR TKIs if mutation-positive.”

**U.S. PREVENTIVE SERVICES TASK FORCE RECOMMENDATIONS**

Not applicable.

**ONGOING AND UNPUBLISHED CLINICAL TRIALS**

Currently unpublished trials that might influence this review are listed in Table 15.

**Table 15. Summary of Key Trials**

<table>
<thead>
<tr>
<th>NCT No.</th>
<th>Trial Name</th>
<th>Planned Enrollment</th>
<th>Completion Date</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Ongoing</strong></td>
<td></td>
<td></td>
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</tr>
<tr>
<td>NCT01306045</td>
<td>Pilot Trial of Molecular Profiling and Targeted Therapy for Advanced Non-Small Cell Lung Cancer, Small Cell Lung Cancer, and Thymic Malignancies</td>
<td>600</td>
<td>Dec 2019</td>
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<tr>
<td>NCT01248247</td>
<td>BATTLE-2 Program: A Biomarker-Integrated Targeted Therapy Study in Previously Treated Patients With Advanced Non-Small Cell Lung Cancer</td>
<td>334</td>
<td>Jun 2019</td>
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<tr>
<td>NCT03225664</td>
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<td>Sep 2020</td>
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<tr>
<td>NCT02117167</td>
<td>Intergroup Trial UNICANCER UC 0105-1305/ IFCT 1301: SAFIR02_Lung - Evaluation of the Efficacy of High Throughput Genome Analysis as a Therapeutic Decision Tool for Patients With Metastatic Non-small Cell Lung Cancer</td>
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<tr>
<td>NCT02465060</td>
<td>Molecular Analysis for Therapy Choice (MATCH)</td>
<td>6452</td>
<td>Jun 2022</td>
</tr>
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</table>

NCT: national clinical trial.

**CODING**

The following codes for treatment and procedures applicable to this policy are included below for informational purposes. Inclusion or exclusion of a procedure, diagnosis or device code(s) does not constitute or imply member coverage or provider reimbursement. Please refer to the member’s contract benefits in effect at the time of service to determine coverage or non-coverage of these services as it applies to an individual member.
CPT/HCPCS
81235  EGFR (epidermal growth factor receptor) (eg, non-small cell lung cancer) gene analysis, common variants (eg, exon 19 LREA deletion L858R, T790M, G719A, G719S, L861Q)
81275  KRAS (Kirsten rat sarcoma viral oncogene homolog) (eg, carcinoma) gene analysis, variants in exon 2 (eg, codons 12 and 13)
81276  KRAS (Kirsten rat sarcoma viral oncogene homolog) (eg, carcinoma) gene analysis; additional variant(s) (eg, codon 61, codon 146)
81404  Molecular pathology procedure, Level 5 (eg, analysis of 2-5 exons by DNA sequence analysis, mutation scanning or duplication/deletion variants of 6-10 exons, or characterization of a dynamic mutation disorder/triplet repeat by Southern blot analysis)
81405  Molecular pathology procedure, Level 6 (eg, analysis of 6-10 exons by DNA sequence analysis, mutation scanning or duplication/deletion variants of 11-25 exons, regionally targeted cytogenomic array analysis)
81406  Molecular pathology procedure, Level 7 (eg, analysis of 11-25 exons by DNA sequence analysis, mutation scanning or duplication/deletion variants of 26-50 exons, cytogenomic array analysis for neoplasia)
81479  Unlisted molecular pathology procedure
88342  Immunohistochemistry or immunocytochemistry, per specimen; initial single antibody stain procedure
88365  In situ hybridization (eg, FISH), per specimen; initial single probe stain procedure
0022U  Targeted genomic sequence analysis panel, non-small cell lung neoplasia, DNA and RNA analysis, 23 genes, interrogation for sequence variants and rearrangements, reported as presence/absence of variants and associated therapy(ies) to consider

- There is a specific CPT code for testing for common variants of EGFR: 81235.
- If testing is done by immunohistochemical assay, CPT code 88342 would likely be reported. If testing is done by fluorescence in situ hybridization (FISH), CPT code 88365 would likely be reported.
- There are specific CPT codes for testing for KRAS: 81275, 81276.
- CPT code 81404 has a listing for RET testing.
- CPT code 81405 has listings for both KRAS and RET testing.
- CPT code 81406 has a listing for BRAF testing.
- Testing for variants in the other genes listed above would be reported with the unlisted molecular pathology code 81479.

ICD-10 Diagnoses
C34.01  Malignant neoplasm of right main bronchus
C34.02  Malignant neoplasm of left main bronchus
C34.11  Malignant neoplasm of upper lobe, right bronchus or lung
C34.12  Malignant neoplasm of upper lobe, left bronchus or lung
C34.2   Malignant neoplasm of middle lobe, bronchus or lung
C34.31  Malignant neoplasm of lower lobe, right bronchus or lung
C34.32  Malignant neoplasm of lower lobe, left bronchus or lung
C34.81  Malignant neoplasm of overlapping sites of right bronchus and lung
C34.82  Malignant neoplasm of overlapping sites of left bronchus and lung
- ICD-10-CM does not have specific coding for non-small-cell lung cancer. The following malignant neoplasm of lung codes would be used.

## REVISIONS

<table>
<thead>
<tr>
<th>Date</th>
<th>Description</th>
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<tbody>
<tr>
<td>09-28-2014</td>
<td>Policy added to the bcbsks.com web site on 08-29-2014. Effective on 09-28-2014, 30 days after posting.</td>
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| 02-08-2015 | Title of policy changed from "Epidermal Growth Factor Receptor Mutation Analysis for Patients with Non-Small Cell Lung Cancer". Updated Description section. In Policy section:  
  - Added "D. Analysis of somatic mutations of the KRAS gene is considered experimental / investigational as a technique to predict treatment non-response to anti-EGFR therapy with tyrosine-kinase inhibitors and for the use of the anti-EGFR monoclonal antibody cetuximab in NSCLC."
  - Added "E. Testing for genetic alterations in the genes ROS, RET, MET, BRAF, and HER2, for targeted therapy in patients with NSCLC, is considered experimental / investigational." |
|           | Updated Rationale section. |
|           | In Coding section:  
  - The following CPT codes were added: 81275, 81404, 81405, 81406, 81479, 88342, 88365. |
|           | Updated References section. |
| 05-14-2015 | Updated Description section. In Policy section:  
  - Added Item D, "Analysis of somatic rearrangement mutations of the ALK gene may be considered medically necessary to predict treatment response to crizotinib in patients with advanced lung adenocarcinoma or in whom an adenocarcinoma component cannot be excluded (see Policy Guidelines)."
  - Added Item E, "Analysis of somatic rearrangement mutations of the ALK gene is considered experimental / investigational in all other clinical situations."
  - In Item G, added "Analysis" and removed "Testing", to read "Analysis for genetic alterations in the genes ROS, RET, MET, BRAF, and HER2, for targeted therapy in patients with NSCLC, is considered experimental / investigational."
  - In Policy Guidelines, Item 2, added "The 2015", "as a category 1 recommendation that", and "and ALK rearrangement testing be performed in the workup of NSCLC in patients with histologic subtypes adenocarcinoma, large-cell carcinoma, and NSCLC not otherwise specified," and removed "a) for patients with advanced lung cancer, nonsquamous cell type, or b) when biopsy specimens are small and histology is mixed," to read, "2. The 2015 guidelines from the National Comprehensive Cancer Network recommend as a category 1 recommendation that EGFR mutation testing and ALK rearrangement testing be performed in the workup of NSCLC in patients with histologic subtypes adenocarcinoma, large-cell carcinoma, and NSCLC not otherwise specified."
  - In Policy Guidelines, Item 3, added "The", "and ALK rearrangement" and "and ALK", and removed "Current", to read, "The 2014 guidelines issued jointly by the College of American Pathologists, International Association for the Study of Lung Cancer, and Association for Molecular Pathology recommend: a) EGFR mutation and ALK rearrangement testing in patients with lung adenocarcinoma regardless of clinical characteristics (eg, smoking history); b) In the setting of fully excised lung cancer specimens, EGFR and ALK mutation testing is not recommended in lung cancers when an adenocarcinoma component is lacking (such as pure squamous cell lacking any
immunohistochemical evidence of adenocarcinomatous differentiation); and c) In the setting of more limited lung cancer specimens (e.g., biopsies, cytology) where an adenocarcinoma component cannot be completely excluded, EGFR and ALK testing may be performed in cases showing squamous cell histology. Clinical criteria (e.g., young age, lack of smoking history) may be useful to select a subset of these samples for testing."

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<tr>
<th>Date</th>
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<th>Rationale</th>
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<th>References</th>
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<td>10-01-2017</td>
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|            | ▪ In Item A, added "an EGFR tyrosine kinase inhibitor (TKI) therapy (e.g., "[Tarceva®], gefitinib [Iressa®],", and "[Gilotrif®])" to read, "Except as noted below, analysis of 2 types of somatic mutation within the EGFR gene—small deletions in exon 19 and a point mutation in exon 21 (L858R)—may be considered medically necessary to predict treatment response to an EGFR tyrosine kinase inhibitor (TKI) therapy (e.g., erlotinib [Tarceva®], gefitinib [Iressa®], or afatinib [Gilotrif®]) in patients with advanced lung adenocarcinoma or in whom an adenocarcinoma component cannot be excluded (see Policy Guidelines)."
|            | ▪ Added new Item B, "Analysis for the T790M mutation in the gene for the EGFR is considered medically necessary as a technique to predict treatment response to osimertinib (Tagrisso™) in patients who have progressed on or after EGFR-TKI therapy." |
|            | ▪ In Policy Guidelines, revised guideline dates for Items 2 and 3 and added "Genetic Counseling." |
|            | Updated     |           |        |            |        |
| 03-14-2018 | Updated     |           |        |            | Updated|
|            | Description | Rationale | Coding | References | Policy |
|            |             |           |        |            |        |
patients with metastatic disease with histologic subtypes adenocarcinoma, large cell carcinoma, and non-small-cell lung cancer not otherwise specified."

• In Item B, removed "mutation" and added "variants" to read, "Analysis of the T790M variants in the EGFR gene is considered medically necessary as a technique to predict treatment response to osimertinib (Tagrisso™) in patients who have progressed on or after EGFR-TKI therapy."

• In Item C, removed "mutation" and added "variants" to read, "Analysis of 2 types of somatic variants within the EGFR gene—small deletions in exon 19 and a point mutation in exon 21 (L858R)—is considered experimental / investigational for patients with advanced squamous cell NSCLC."

• In Item D, removed "mutations" and added "variants" to read, "Analysis of other EGFR variants within exons 18 to 24, or other applications related to NSCLC, is considered experimental / investigational."

• In Item E, removed "mutations" and "with advanced lung adenocarcinoma or in whom an adenocarcinoma component cannot be excluded (see Policy Guidelines)" and added "variants", "ALK inhibitor therapy (eg, [Xalkori®], ceritinib [Zykadia™], alectinib [Alecensa®], or brigatinib [Alunbrig™])", and "with metastatic disease with histologic subtypes adenocarcinoma, large cell carcinoma, and non-small-cell lung cancer not otherwise specified" to read, "Analysis of somatic rearrangement variants of the ALK gene may be considered medically necessary to predict treatment response to ALK inhibitor therapy (eg, crizotinib [Xalkori®], ceritinib [Zykadia™], alectinib [Alecensa®], or brigatinib [Alunbrig™]) in patients with metastatic disease with histologic subtypes adenocarcinoma, large cell carcinoma, and non-small-cell lung cancer not otherwise specified."

• In Item F, removed "mutations" and "clinical" and added "variants" to read, "Analysis of somatic rearrangement variants of the ALK gene is considered experimental / investigational in all other situations."

• Added new Item G, "Analysis of the BRAF V600E variant may be considered medically necessary to predict treatment response to BRAF or MEK inhibitor therapy (eg, dabrafenib [Tafinlar®] and trametinib [Mekinist®]), in patients with metastatic disease with histologic subtypes adenocarcinoma, large cell carcinoma, and non-small-cell lung cancer not otherwise specified."

• Added new Item H, "Analysis of somatic rearrangement variants of the ROS1 gene may be considered medically necessary to predict treatment response to ALK inhibitor therapy (crizotinib [Xalkori®]) in patients with metastatic disease with histologic subtypes adenocarcinoma, large cell carcinoma, and non-small-cell lung cancer not otherwise specified."

• In previous Item G (now Item I), removed "mutations" and added "variants" to read, "Analysis of somatic variants of the KRAS gene is considered experimental / investigational as a technique to predict treatment non-response to anti-EGFR therapy with tyrosine-kinase inhibitors and for the use of the anti-EGFR monoclonal antibody cetuximab in NSCLC."

• In Previous Item H (now Item J), removed "ROS" and "BRAF" to read, "Analysis for genetic alterations in the genes RET, MET, and HER2 for targeted therapy in patients with NSCLC is considered experimental / investigational."

• Added new Item K, "Programmed death receptor 1 (PD-1) or its ligand (PD-L1) expression analysis may be considered medically necessary as a technique to predict treatment response to drug therapy."

• Updated Policy Guidelines.

Updated Rationale section.

In Coding section:

• Updated nomenclature for CPT codes: 88342, 88365.
• Updated coding bullets.
<table>
<thead>
<tr>
<th>09-12-2018</th>
<th>In Policy section:</th>
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<tbody>
<tr>
<td>• In Item A, removed &quot;metastatic&quot; and added &quot;stage III or IV&quot; to read, &quot;Except as noted below, analysis of 2 types of somatic variants within the EGFR gene—small deletions in exon 19 and a single-nucleotide variant in exon 21 (L858R)—may be considered medically necessary to predict treatment response to an EGFR tyrosine kinase inhibitor (TKI) therapy (eg, erlotinib [Tarceva®], gefitinib [Iressa®], or afatinib [Gilotrif®]) in patients with stage III or IV disease with histologic subtypes adenocarcinoma, large cell carcinoma, and non-small-cell lung cancer not otherwise specified.&quot;</td>
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<tr>
<td>• In Item E, removed &quot;metastatic&quot; and added &quot;stage III or IV&quot; to read, &quot;Analysis of somatic rearrangement variants of the ALK gene may be considered medically necessary to predict treatment response to ALK inhibitor therapy (eg, crizotinib [Xalkori®], ceritinib [ZykadiaTM], alectinib [Alecensa®], or brigatinib [Alunbrig™]) in patients with stage III or IV disease with histologic subtypes adenocarcinoma, large cell carcinoma, and non-small-cell lung cancer not otherwise specified.&quot;</td>
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<tr>
<td>• In Item G, removed &quot;metastatic&quot; and added &quot;stage III or IV&quot; to read, &quot;Analysis of the BRAF V600E variant may be considered medically necessary to predict treatment response to BRAF or MEK inhibitor therapy (eg, dabrafenib [Tafinlar®] and trametinib [Mekinist®]), in patients with stage III or IV metastatic disease with histologic subtypes adenocarcinoma, large cell carcinoma, and non-small-cell lung cancer not otherwise specified.&quot;</td>
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<tr>
<td>• In Item H, removed &quot;metastatic&quot; and added &quot;stage III or IV&quot; to read, &quot;Analysis of somatic rearrangement variants of the ROS1 gene may be considered medically necessary to predict treatment response to ALK inhibitor therapy (crizotinib [Xalkori®]) in patients with stage III or IV metastatic disease with histologic subtypes adenocarcinoma, large cell carcinoma, and non-small-cell lung cancer not otherwise specified.&quot;</td>
<td></td>
</tr>
</tbody>
</table>

**Updated References section.**

**REFERENCES**


57. Miller VA, Hirsh V, Cadranel J, et al. Afatinib versus placebo for patients with advanced, metastatic non-small-cell lung cancer after failure of erlotinib, gefitinib, or both, and one or two lines of


Other References
1. Blue Cross and Blue Shield of Kansas Oncology Liaison Committee, February 2015; August 2017.
2. Blue Cross and Blue Shield of Kansas Pathology Liaison Committee, July 2016; May 2018.

APPENDIX

Appendix Table 1. Categories of Genetic Testing Addressed

<table>
<thead>
<tr>
<th>Category</th>
<th>Addressed</th>
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<tbody>
<tr>
<td>1. Testing of an affected individual’s germline to benefit the individual</td>
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<tr>
<td>1a. Diagnostic</td>
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<tr>
<td>1b. Prognostic</td>
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<tr>
<td>1c. Therapeutic</td>
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<tr>
<td>2. Testing cancer cells from an affected individual to benefit the individual</td>
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<tr>
<td>2a. Diagnostic</td>
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</tr>
<tr>
<td>2b. Prognostic</td>
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<tr>
<td>2c. Therapeutic</td>
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<tr>
<td>3. Testing an asymptomatic individual to determine future risk of disease</td>
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<tr>
<td>4. Testing of an affected individual’s germline to benefit family members</td>
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<tr>
<td>5. Reproductive testing</td>
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<tr>
<td>5a. Carrier testing: preconception</td>
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<tr>
<td>5b. Carrier testing: prenatal</td>
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<tr>
<td>5c. In utero testing: aneuploidy</td>
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<tr>
<td>5d. In utero testing: mutations</td>
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<tr>
<td>5e. In utero testing: other</td>
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<tr>
<td>5f. Preimplantation testing with in vitro fertilization</td>
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