

<b>POLICY TITLE</b>	<b>MOLECULAR ANALYSIS FOR TARGETED THERAPY FOR NON-SMALL CELL LUNG CANCER</b>
<b>POLICY NUMBER</b>	<b>MP- 2.241</b>

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**I. POLICY**

***EGFR Testing***

Analysis of somatic variants in exons 18 through 21 (e.g., G719X, L858R, T790M, S6781, L861Q) within the epidermal growth factor receptor (*EGFR*), may be considered **medically necessary** to predict treatment response to an EGFR tyrosine kinase inhibitor therapy (e.g., erlotinib [Tarceva], gefitinib [Iressa], afatinib [Gilotrif], or osimertinib [Tagrisso]) in patients with advanced lung adenocarcinoma, large cell carcinoma, advanced squamous cell non-small-cell lung cancer, and non-small-cell lung cancer not otherwise specified.

Analysis of other *EGFR* variants within exons 22 to 24, or other applications related to NSCLC, is considered **investigational**.

***ALK Testing***

Analysis of somatic rearrangement variants of the anaplastic lymphoma kinase (*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 advanced lung adenocarcinoma or in whom an adenocarcinoma component cannot be excluded (see Policy Guidelines section).

Analysis of somatic rearrangement variants of the *ALK* gene is considered **investigational** in all other situations. There is insufficient evidence to support a conclusion concerning the health outcomes or benefits associated with these procedures.

***BRAF V600E TESTING***

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 advanced lung adenocarcinoma or in whom an adenocarcinoma component cannot be excluded (see Policy Guidelines section).

***ROS1 TESTING***

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

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patients with advanced lung adenocarcinoma or in whom an adenocarcinoma component cannot be excluded (see Policy Guidelines section).

**KRAS Testing**

Analysis of somatic variants of the *KRAS* gene is considered **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. There is insufficient evidence to support a conclusion concerning the health outcomes or benefits associated with these procedures.

**NTRK Testing**

Analysis of gene fusions may be considered **medically necessary** to predict treatment response to larotrectinib in patients with advanced lung adenocarcinoma or in whom an adenocarcinoma component cannot be excluded (see Policy Guidelines section.)

**Other Genes**

Testing for genetic alterations in the genes *HER2*, *RET*, and *MET* for targeted therapy in patients with NSCLC is considered **investigational**. There is insufficient evidence to support a conclusion concerning the health outcomes or benefits associated with these procedures.

**Tumor Mutational Burden**

Tumor mutational burden testing is a biomarker of outcomes with immunotherapy in multiple tumor types, including lung cancer. This procedure is considered **investigational**. There is insufficient evidence to support a conclusion concerning the health outcomes or benefits associated with these procedures.

**Policy Guidelines**

These gene tests are intended for use in patients with advanced non-small-cell lung cancer. 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 receptor (*EGFR*) 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.

The 2019 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. The guidelines add that testing should be conducted as part of broad molecular profiling and should include the NTRK gene fusion.

The 2018 guidelines issued jointly by the College of American Pathologists, International Association for the Study of Lung Cancer, and Association for Molecular Pathology have recommended the following:

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“One set of genes must be offered by all laboratories that test lung cancers, as an absolute minimum: EGFR, ALK, and ROS1. A second group of genes should be included in any expanded panel that is offered for lung cancer patients: BRAF, MET, RET, ERBB2 (HER2), and KRAS, if adequate material is available. KRAS testing may also be offered as a single-gene test to exclude patients from expanded panel testing. All other genes are considered investigational at the time of publication.”

**II. PRODUCT VARIATIONS**

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This policy is only applicable to certain programs and products administered by Capital BlueCross please see additional information below, and subject to benefit variations as discussed in Section VI below.

**FEP PPO** - Refer to FEP Medical Policy Manual 2.04.45 Molecular Analysis for Targeted Therapy of Non-Small-Cell Lung Cancer. The FEP Medical Policy manual can be found at: <https://www.fepblue.org/benefit-plans/medical-policies-and-utilization-management-guidelines/medical-policies>.

**III. DESCRIPTION/BACKGROUND**

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**NON-SMALL-CELL LUNG CANCER**

Treatment options for non-small-cell lung cancer (NSCLC) depend on disease stage and include various combinations of surgery, radiotherapy, systemic therapy, and best supportive care. Unfortunately, in up to 85% of cases, the cancer has spread locally beyond the lungs at diagnosis, precluding surgical eradication. Also, up to 40% of patients with NSCLC present with metastatic disease. 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%. 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 epidermal growth factor receptor (EGFR) variants and anaplastic lymphoma kinase (ALK) rearrangements is routine in clinical decision making for the treatment of NSCLC. The use of testing for other variants to direct targeted therapy 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.

## MEDICAL POLICY

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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.

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**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.

**NTRK Gene Fusions**

NTRK gene fusions encode tropomyosin receptor kinase fusion proteins that act as oncogenic drivers for solid tumors including lung, salivary gland, thyroid, and sarcoma. It is estimated that NTRK gene fusions occur in 0.2% of patients with NSCLC and do not typically overlap with other oncogenic drivers.

**Tumor Mutational Burden**

Tumor mutational burden is an emerging biomarker of outcomes with immunotherapy in multiple tumor types, including lung cancer.

**TARGETED THERAPIES**

Four orally administered EGFR-selective, small-molecule TKIs have been identified for treating NSCLC: gefitinib (Iressa; AstraZeneca), erlotinib (Tarceva; OSI Pharmaceuticals), afatinib (Gilotrif; Boehringer Ingelheim), and osimertinib (Tagrisso; AstraZeneca). Gefitinib, erlotinib, afatinib, and osimertinib currently are approved by the U.S. Food and Drug Administration (FDA) for NSCLC when *EGFR* status is confirmed through a companion diagnostic test.

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 confirmed through a companion diagnostic test. 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 *BRAF* V600 variants confirmed through a companion diagnostic test. The combination of dabrafenib and trametinib was approved for treatment of metastatic NSCLC in 2017 for patients with confirmed *BRAF* V600 variants.

For the treatment of *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

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patients with advanced or recurrent NSCLC as first-line and maintenance therapy. Panitumumab is not used in NSCLC.

Targeted therapies currently under investigation and not FDA-approved for the remaining genetic alterations in NSCLC are trastuzumab and afatinib for *HER2* variants, crizotinib for *MET* amplification, and cabozantinib for *RET* rearrangements.

Larotrectinib was approved in 2018 for the treatment of patients with solid tumors harboring an NTRK gene fusion. There is currently no FDA approved companion diagnostic test for larotrectinib. The clinical review states, "The clinical review team and CDRH agreed that it is in the best interest of U.S. patients to approve larotrectinib before one or more companion diagnostic assays are ready for a PMA submission. Loxo Oncology has agreed to a postmarketing commitment to work with diagnostic developers to develop an analytically and clinically validated companion diagnostic test for the selection of patients with NTRK fusion-positive solid tumors for whom larotrectinib is safe and effective."

Nivolumab in combination with ipilimumab has been investigated as a treatment option for patients with NSCLC with tumor mutational burden >10 mutations per megabase. There is no FDA companion diagnostic test for tumor mutational burden.

Targeted therapies currently under investigation and not FDA-approved for the remaining genetic alterations in NSCLC are trastuzumab and afatinib for *HER2* variants, crizotinib for *MET* amplification, and cabozantinib for *RET* rearrangements.

**REGULATORY STATUS**

Table 1 summarizes the FDA-approved targeted treatments for patients with NSCLC along with the concurrently approved diagnostic tests.

**Table 1. FDA-Approved Targeted Treatment for NSCLC and Companion Diagnostic Tests**

<b>Treatment</b>	<b>Indication</b>	<b>FDA Approval of Companion Diagnostic Test</b>
<b>Afatinib (Gilotrif)</b>	<ul style="list-style-type: none"> <li>• 2013: First line for patients with metastatic NSCLC whose tumors have <i>EGFR</i> exon 19 deletions or exon 21 (L858R) substitutions</li> <li>• 2016: Second line for patients with metastatic squamous NSCLC</li> <li>• 2018: First line for patients with nonresistant <i>EGFR</i> variants other than exon 19 or exon 21 NSCLC</li> </ul>	<ul style="list-style-type: none"> <li>• 2013: theascreen® <i>EGFR</i> Rotor-Gene Q polymerase chain reaction (RGQ PCR) kit (Qiagen)</li> <li>• 2017: FoundationOne CDx™ (Foundation Medicine)</li> </ul>
<b>Alectinib (Alecensa)</b>	<ul style="list-style-type: none"> <li>• 2015: Second line for patients with <i>ALK</i>-positive metastatic NSCLC who have progressed on or are intolerant of crizotinib</li> </ul>	2017: FoundationOne CDx™ (Foundation Medicine)

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	<ul style="list-style-type: none"> <li>• 2017: First line for patients with <i>ALK</i>-positive NSCLC who have not received prior systemic therapy for metastatic disease</li> </ul>	
<b>Brigatinib (Alunbrig)</b>	<ul style="list-style-type: none"> <li>• 2017: Second line for patients with metastatic <i>ALK</i>-positive NSCLC who have progressed on or are intolerant of crizotinib</li> </ul>	Test not specified in FDA approval
<b>Ceritinib (Zykadia)</b>	<ul style="list-style-type: none"> <li>• 2014: Second line for patients with <i>ALK</i>-positive metastatic NSCLC who have progressed on or are intolerant of crizotinib</li> <li>• 2017: First line for patients with <i>ALK</i>-positive metastatic NSCLC</li> </ul>	<ul style="list-style-type: none"> <li>• 2015: Ventana <i>ALK</i> (D5F3) CDx Assay (Ventana Medical Systems)</li> <li>• 2017: FoundationOne CDx™ (Foundation Medicine)</li> </ul>
<b>Crizotinib (Xalkori)</b>	<ul style="list-style-type: none"> <li>• 2011: First line for patients with <i>ALK</i>-positive metastatic NSCLC</li> </ul>	<ul style="list-style-type: none"> <li>• 2011: Vysis <i>ALK</i> Break Apart FISH Probe Kit (Abbott Laboratories)</li> <li>• 2015: Ventana <i>ALK</i> (D5F3) CDx Assay (Ventana Medical Systems)</li> <li>• 2017: FoundationOne CDx™ (Foundation Medicine)</li> </ul>
<b>Crizotinib (Xalkori)</b>	<ul style="list-style-type: none"> <li>• 2016: Patients with <i>ROS1</i>-positive metastatic NSCLC</li> </ul>	<ul style="list-style-type: none"> <li>• 2017: Oncomine™ Dx Target Test (Thermo Fisher Scientific)</li> </ul>
<b>Dacomitinib (Vizimpro)</b>	<ul style="list-style-type: none"> <li>• 2018: First line for patients with metastatic NSCLC with <i>EGFR</i> exon 19 deletion or exon 21 (L858R) substitutions</li> </ul>	Test not specified in FDA approval
<b>Dabrafenib (Tafinlar) plus trametinib (Mekinist)</b>	<ul style="list-style-type: none"> <li>• 2017: Used in combination for treatment of patients with metastatic NSCLC with <i>BRAF</i> V600E variant</li> </ul>	<ul style="list-style-type: none"> <li>• 2017: Oncomine™ Dx Target Test</li> <li>• 2017: FoundationOne CDx™ (Foundation Medicine)</li> </ul>
<b>Erlotinib (Tarceva)</b>	<ul style="list-style-type: none"> <li>• 2013: First line for patients with metastatic NSCLC whose tumors have <i>EGFR</i> exon 19 deletions or exon 21 (L858R) substitutions</li> <li>• 2010: Maintenance for patients with locally advanced or metastatic NSCLC whose disease has not progressed after 4 cycles of platinum-based chemotherapy</li> </ul>	<ul style="list-style-type: none"> <li>• 2013: cobas® <i>EGFR</i> Mutation Test (tissue test) (Roche Diagnostics)</li> <li>• 2016: cobas® <i>EGFR</i> Mutation Test v2 (tissue</li> </ul>

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	<ul style="list-style-type: none"> <li>• 2004: Second line for patients with locally advanced or metastatic NSCLC</li> </ul>	<ul style="list-style-type: none"> <li>or blood test) (Roche Diagnostics)</li> <li>• 2017: FoundationOne CDx™ (Foundation Medicine)</li> </ul>
<b>Gefitinib (Iressa)</b>	<ul style="list-style-type: none"> <li>• 2015: First line for patients with metastatic NSCLC whose tumors have <i>EGFR</i> exon 19 deletions or exon 21 (L858R) substitutions</li> <li>2003: Second line for patients with locally advanced or metastatic NSCLC</li> </ul>	<ul style="list-style-type: none"> <li>• 2015: theascreen® EGFR Rotor-Gene Q polymerase chain reaction (RGQ PCR) kit</li> <li>• 2017: Oncomine™ Dx Target Test</li> <li>• 2017: FoundationOne CDx™ (Foundation Medicine)</li> <li>• 2017: cobas® EGFR Mutation Test (tissue test) (Roche Diagnostics)</li> </ul>
<b>Osimertinib (Tagrisso)</b>	<ul style="list-style-type: none"> <li>• 2015: Second line for patients with metastatic NSCLC whose tumors have <i>EGFR</i> T790M variants as detected by FDA-approved test, who have not responded to <i>EGFR</i>-blocking therapy</li> <li>• 2018: First line for patients with metastatic NSCLC whose tumors have <i>EGFR</i> exon 19 deletions or exon 21 L858R variants</li> </ul>	<ul style="list-style-type: none"> <li>• 2015: cobas® EGFR Mutation Test v2 (blood test)</li> <li>• 2017: FoundationOne CDx™ (Foundation Medicine)</li> </ul>
<b>Larotrectinib (Vitrakvi)</b>	<ul style="list-style-type: none"> <li>• 2018: Adult and pediatric patients with solid tumors that:                             <ul style="list-style-type: none"> <li>○ have a neurotrophic receptor tyrosine kinase (NTRK) gene fusion without a known acquired resistance mutation,</li> <li>○ are metastatic or where surgical resection is likely to result in severe morbidity, and</li> <li>○ have no satisfactory alternative treatments or that have progressed following treatment.</li> </ul> </li> </ul>	Test not specified in FDA approval

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.



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**IV. RATIONALE**

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**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 (eg, afatinib, erlotinib, gefitinib, osimertinib) with chemotherapy. The relevant outcomes are OS, disease-specific survival, test validity, QOL, and treatment-related morbidity. Studies have shown that TKIs are superior to chemotherapy regarding tumor response rate and PFS, with a reduction in toxicity and improvement in the QOL. 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. The relevant outcomes are OS, disease-specific survival, test validity, QOL, 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 amplification, 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. The relevant outcomes are OS, disease-specific survival, test validity, QOL, 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 two 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 RCTs of advanced KRAS-variant positive disease, MEK inhibitors did not improve PFS compared with docetaxel. Studies for HER2, RET, and MET variant testing have reported response rates and PFS 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.

For individuals who have advanced-stage NSCLC who are being considered for targeted therapy who receive NTRK gene fusion testing, the evidence includes prospective observational studies. The relevant outcomes are OS, disease-specific survival, test validity, QOL, and treatment-related morbidity. In 55 patients with consecutively and prospectively identified TRK fusion-positive solid tumors, including 4 patients with lung tumors, the overall response rate was 80%

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(95% CI, 67 to 90). The median PFS had not been reached after a median follow-up duration of 9.9 months (range, 0.7 to 25.9). Responses were observed regardless of tumor type or age of the patient. 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 TMB testing, the evidence includes an RCT and retrospective observational studies. In a subgroup analysis of an ongoing RCT, PFS was significantly longer with nivolumab plus ipilimumab than with chemotherapy among patients with NSCLC and a high TMB (>10 mutations per megabase). In exploratory analyses, retrospective observational studies have reported an association between higher TMB and longer PFS and OS in patients receiving immunotherapy. These results need to be confirmed in additional, well-designed prospective studies. The evidence is insufficient to determine the effects of the technology on health outcomes.

**V. DEFINITIONS**

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**VI. BENEFIT VARIATIONS**

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The existence of this medical policy does not mean that this service is a covered benefit under the member's health benefit plan. Benefit determinations should be based in all cases on the applicable health benefit plan language. Medical policies do not constitute a description of benefits. A member's health benefit plan governs which services are covered, which are excluded, which are subject to benefit limits and which require preauthorization. There are different benefit plan designs in each product administered by Capital BlueCross. Members and providers should consult the member's health benefit plan for information or contact Capital BlueCross for benefit information.

**VII. DISCLAIMER**

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*Capital BlueCross's medical policies are developed to assist in administering a member's benefits, do not constitute medical advice and are subject to change. Treating providers are solely responsible for medical advice and treatment of members. Members should discuss any medical policy related to their coverage or condition with their provider and consult their benefit information to determine if the service is covered. If there is a discrepancy between this medical policy and a member's benefit information, the benefit information will govern. If a provider or a member has a question concerning the application of this medical policy to a specific member's plan of benefits, please contact Capital BlueCross' Provider Services or Member Services. Capital BlueCross considers the information contained in this medical policy to be proprietary and it may only be disseminated as permitted by law.*

**VIII. CODING INFORMATION**

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**Note:** This list of codes may not be all-inclusive, and codes are subject to change at any time. The identification of a code in this section does not denote coverage as coverage is determined by the terms of member benefit information. In addition, not all covered services are eligible for separate reimbursement.

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**Investigational; therefore, not covered for KRAS testing for NSCLC when used to predict treatment non-response to anti-EGFR therapy:**

CPT Codes®								
81275	81276	88363						

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**Investigational; therefore, not covered when used to test for genetic alterations in the genes RET, MET, HER2, and TMB for targeted therapy in patients with NSCLC:**

CPT Codes®								
81404	81405	81479	0022U	0037U				

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**Covered when medically necessary:**

CPT Codes®								
81191	81192	81193	81194	81210	81235	81401	88342	88365

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ICD-10-CM Diagnosis Codes	Description
C34.00	Malignant neoplasm of unspecified main bronchus
C34.01	Malignant neoplasm of right main bronchus
C34.02	Malignant neoplasm of left main bronchus
C34.10	Malignant neoplasm of upper lobe, unspecified bronchus or lung
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
C34.30	Malignant neoplasm of lower lobe, unspecified 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
C34.80	Malignant neoplasm of overlapping sites of unspecified bronchus and 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
C34.90	Malignant neoplasm of unspecified part of unspecified bronchus or lung
C34.91	Malignant neoplasm of unspecified part of right bronchus or lung
C34.92	Malignant neoplasm of unspecified part of left bronchus or lung
D38.1	Neoplasm of uncertain behavior of trachea, bronchus and lung

**IX. REFERENCES**

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**MEDICAL POLICY**

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<b>MP 2.241</b>	<b>CAC 4/26/11 New Policy.</b> Adopted BCBSA (medically necessary).
	<b>CAC 6/26/12 Consensus review.</b> No changes, references updated. FEP variation added.
	<b>12/20/2013 Administrative update.</b> 2013 Codes added
	<b>05/20/13 Administrative update.</b> Code review completed.
	<b>8/1/13 Administrative update.</b> Add Medicare variation referencing LCD L33142.
	<b>CAC 11/26/13 Consensus review.</b> No change to policy statements. Added rationale section. References updated.
	<b>7/24/14 Administrative update.</b> For the Medicare variation - For Novitas MAC jurisdictions, the LCD has been assigned a new number. Biomarkers for Oncology LCD changed from L33124 to L34796
	<b>CAC 1/27/15 Minor revision.</b> <ul style="list-style-type: none"> <li>▪ Policy title revised to “Molecular Analysis for Targeted Therapy for Non-Small Cell Lung Cancer” for this review.</li> <li>▪ First policy statement revised to state analysis of two types of somatic mutation within the epidermal growth factor receptor (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 erlotinib or afatinib in patients with advanced lung adenocarcinoma or in whom an adenocarcinoma component cannot be excluded. Previously, afatinib was not included and the medically necessary indication as for non-small cell lung cancer.</li> </ul>

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	<ul style="list-style-type: none"> <li>▪ A new statement was added that testing for ROS, RET, MET, BRAF and HER2 mutations is considered investigational.</li> <li>▪ Policy MP-2.240 KRAS Mutation Analysis in Non-Small Cell Lung Cancer (NSCLC) merged with this policy.</li> </ul> <p>Background, rationale, references, and guidelines updated. FEP variation updated to reflect combining two policies. (MP 2.240 Info added into this policy). Cellular and histologic classification of NSCLC added to policy guidelines. No coding changes.</p>
	<p><b>11/2/15 Administrative update.</b> LCD number changed from L34796 to L35396 due to Novitas update to ICD-10.</p>
	<p><b>CAC 3/29/16 Minor revision.</b> Policy updated only to address <i>ALK</i> rearrangement testing. Policy statement added that analysis of somatic rearrangement mutations of the <i>ALK</i> gene is considered medically necessary to predict treatment response to crizotinib in patients with advanced lung adenocarcinoma or in whom an adenocarcinoma component cannot be excluded. Policy statement added that testing for <i>ALK</i> gene rearrangements is investigational for all other clinical situations. Reference and rationale update. Coding reviewed.</p>
	<p><b>1/1/17 Administrative update.</b> Product variation section reformatted.</p>
	<p><b>CAC 3/28/17 Minor revision.</b> Added testing for T790M mutation in patients who have progressed on or after EGFR-TKI therapy to the EGFR Gene medically necessary statement. The medically necessary statement language for the EGFR and ALK Genes were revised to reference the gene inhibitor therapy with examples of therapy types/drugs. FEP variation updated. Policy Guidelines, Description/Background, Rationale and Reference sections updated. Appendix added. Coding Reviewed</p>
	<p><b>1/1/18 Administrative update.</b> Medicare variations removed from Commercial Policies.</p>
	<p><b>1/25/18 Minor revision.</b> Policy section updates:</p> <ul style="list-style-type: none"> <li>• Genetics nomenclature update ‘mutation’ to ‘variant’;</li> <li>• 'Point mutation' replaced with 'single-nucleotide variant' within the EGFR Gene policy statement;</li> <li>• <i>ROS1</i> and <i>BRAF</i> testing added as medically necessary when criteria are met; and</li> <li>• Genes within the ‘Other Genes’ statement updated.</li> </ul> <p>FEP variation removed as policy was archived December 2015. Policy Guidelines, Description/Background, Rationale and Reference sections updated. Coding revised to mirror policy statements.</p>
	<p><b>12/14/18 Minor review.</b> The policy section on <i>EGFR</i> Testing was changed. Testing for additional variants in the <i>EGFR</i> gene was added as medically necessary. Policy statements on other variants unchanged.</p>

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	Description/Background and Reference sections updated. Rationale condensed. Coding reviewed.
	<b>11/7/19 Minor review.</b> New indications for NTRK testing and tumor mutational burden (TMB) testing added. Medically necessary statement for NTRK testing and investigational statement for TMB testing. Literature reviewed and references and coding updated. Effective 5/1/2020.
	<b>9/10/2020 Consensus review.</b> No changes made to the policy statement. Policy guidelines, product variation, description, rationale, benefit variation, disclaimer, and references updated. Coding reviewed.
	<b>9/22/2020 Administrative update.</b> New codes 81191, 81192, 81193, 81194 added. Effective 1/1/2021.

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