

NON-SMALL CELL LUNG CANCER (NSCLC) BIOMARKER TESTING LANDSCAPE

Progress in NSCLC

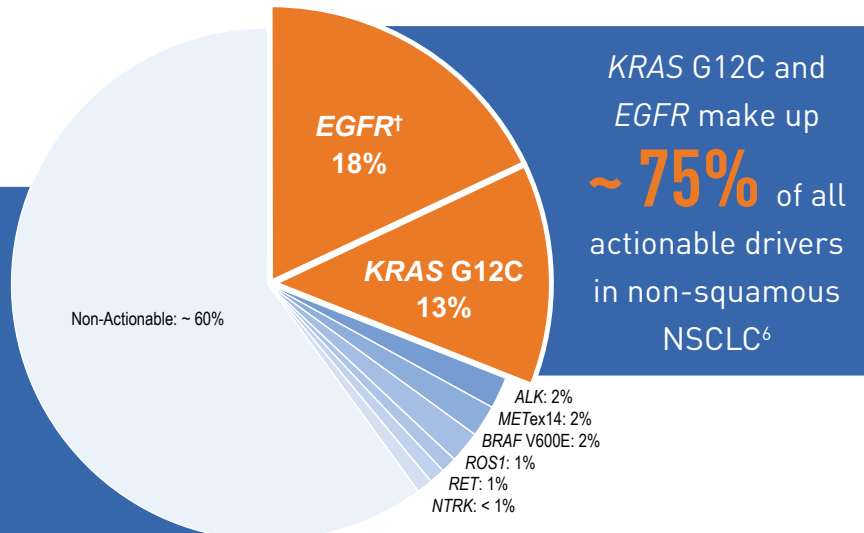
The biomarker landscape has evolved in recent years



- More than 20 targeted therapies have been approved for use in NSCLC¹
- ~ 60% of cancer therapies launched in the US between 2015 and 2020 require or recommend biomarker testing prior to use⁵

Prevalence of Actionable Oncogenic Drivers in NSCLC

~ 2 in every 5
patients with non-squamous NSCLC
have an actionable driver mutation^{6,*}



*From a 2020 analysis of patients with NSCLC in the AACR Genie database (v8.0, N=14,485) and prevalence of KRAS G12C and mutations or alterations with an annotation of "FDA-recognized biomarker predictive of response to an FDA-approved drug in this indication" in non-squamous patients.⁶

[†]EGFR prevalence does not include exon 20 insertions, which can be found in ~ 2% of the overall NSCLC population.⁷

Guidelines Recommend Broad Molecular Testing for Eligible Patients With Advanced NSCLC^{8,9}

NCCN Clinical Practice Guidelines in Oncology (NCCN Guidelines[®]) Recommendations^{8,†,§}

Actionable	Molecular Biomarker								Immune Biomarker	Emerging	Molecular Biomarker	
	EGFR	KRAS G12C	ALK	METex14	BRAF	ROS1	RET	NTRK1/2/3	PD-L1		METamp	HER2
	Testing should be conducted as part of broad molecular profiling								Single-biomarker immunohistochemistry testing recommended		Expanded-panel testing recommended	

[†]The NCCN Clinical Practice Guidelines in Oncology (NCCN Guidelines[®]) for NSCLC provide recommendations for certain individual biomarkers that should be tested and recommend testing techniques but do not endorse any specific commercially available biomarker assays or commercial laboratories.⁸

[§]The NCCN Guidelines[®] for NSCLC recommend broad molecular testing to identify rare driver variants for which targeted therapies may be available to ensure patients receive the most appropriate treatment.⁸

- CAP/AMP/IASLC guidelines recommend testing for actionable and emerging biomarkers utilizing a comprehensive panel or targeted testing⁹

Guideline-Recommended Biomarker Testing May Improve Patient Outcomes^{10,*†}

Adherence to testing for guideline-recommended biomarkers, regardless of therapy

Decreased mortality risk by **11%**

*This was a retrospective study of 28,784 patients diagnosed with advanced NSCLC. Adherence to biomarker testing consisted of patients with evidence of testing for any biomarker, including *EGFR*, *ALK*, *BRAF*, *KRAS*, *ROS1*, or PD-L1 between 14 days prior to and 90 days after diagnosis of advanced NSCLC and the main outcome, overall survival (OS), was agnostic to treatment.¹⁰

†Multivariable analysis was adjusted for age at diagnosis of advanced NSCLC, sex, smoking status, and stage at initial diagnosis of NSCLC.¹⁰

Many Patients With Newly Diagnosed NSCLC Do Not Receive Broad Molecular Testing¹¹



~ 50%
of metastatic patients
received comprehensive
biomarker testing^{11,‡}

Regardless of patient characteristics such as age, race, and smoking status, **biomarker testing** should be conducted in **all eligible patients** with advanced NSCLC¹²

‡A retrospective, observational study assessing real-world biomarker testing patterns in 3,474 patients with metastatic NSCLC from community oncology practices within The US Oncology Network between 2018 and 2020.¹¹

Additional Considerations for Comprehensive Biomarker Testing



Addressing Tissue Insufficiency

- Multigene testing can reduce the number of ordered assays and conserve tissue needed to assess all actionable biomarkers¹²
- Rapid On-Site Evaluation (ROSE) assesses sample adequacy for molecular diagnostic studies to potentially help reduce rebiopsy rates¹³
- Liquid biopsy, which has a high degree of concordance (> 98.2%[§]) and improved turnaround time, can be used when tissue collection is not feasible¹⁴



Shortening Turnaround Time (TAT)

- Broad molecular testing at diagnosis may take less time than consecutive single-gene testing, a process of elimination approach¹⁵
- Reflex testing protocols can reduce average TAT by 37 days¹⁶



Consistent Reporting

- Consider including all actionable biomarkers at the beginning of the report using established nomenclature for genetic alterations¹⁷

§Overall concordance across four genes (*EGFR* exon 19 deletion and L858R, *ALK* fusion, *ROS1* fusion, and *BRAF* V600E).¹⁴

Learn more at [FindKRASG12C.com](https://www.findKRASG12C.com)



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