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Best Practices When Choosing Tissue or Liquid Biopsy Specimens For Biomarker Testing in Non-Small Cell Lung Cancer (NSCLC)

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Biomarker testing for NSCLC: Background

Molecular Diagnostics in Clinical Oncology

Genetic Testing for Hereditary Cancer Risk

Assays

- Single gene analysis
- Multigene Panels
- WES
- WGS

Applications

- Identification of patients at risk
- Familial risk
- Treatment selection

Screening tests

Assays

- ctDNA assays (i.e. methylation, fragmentomics)
- CTCs
- Immunoassays

Applications

- Early cancer detection
- Screening

Tests for Early-Stage Cancer

Assays

- CTC assays
- Gene expression panels
- PCR/RT-PCR
- NGS hotspot & multigene panels
- ctDNA MRD & monitoring assays
- Immunoassays

Applications

- Prognosis/Risk stratification
- Minimal residual disease
- Tumor response monitoring
- Therapy Selection

Tumor Molecular Profiling for Advanced/Metastatic Cancer

Assays

- PCR/RT-PCR
- NGS hotspot & multigene panels
- WES
- WGS
- Immunoassays
- ctDNA monitoring assays
- CTC assays

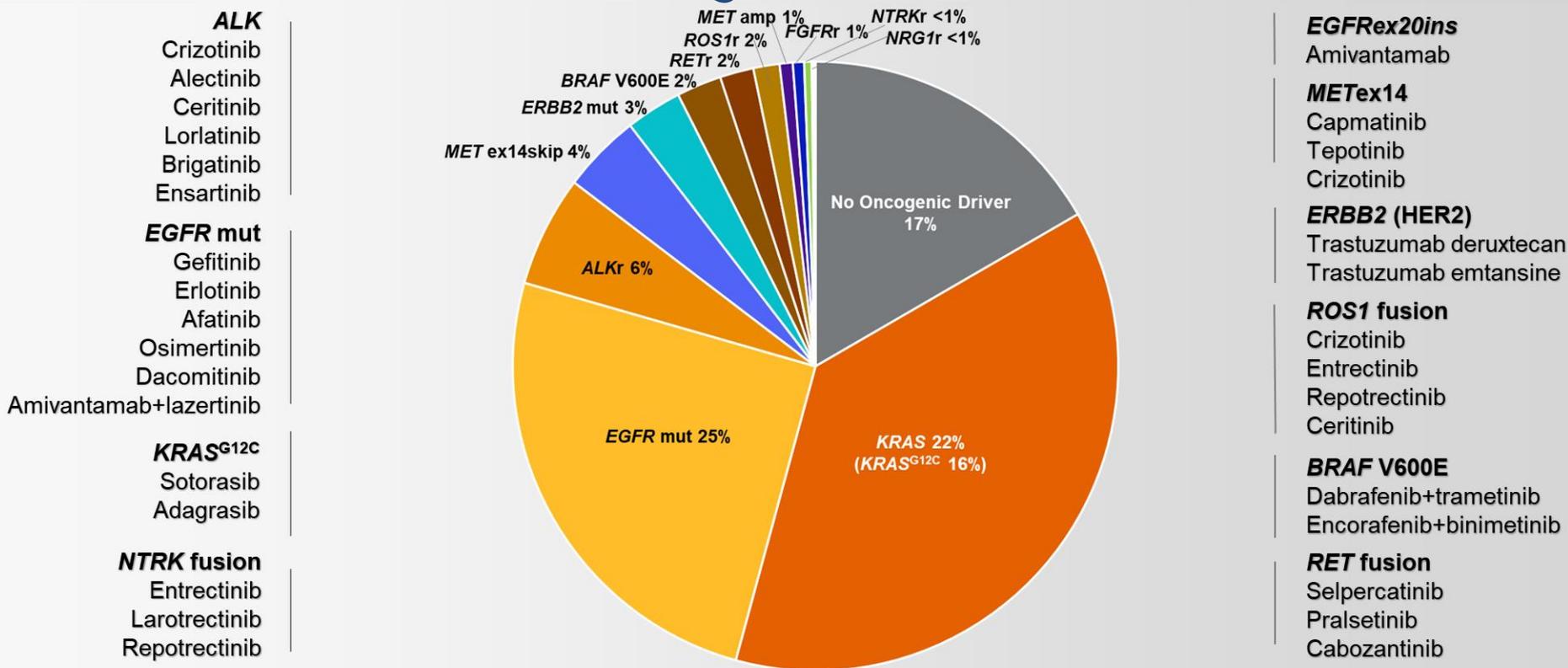
Applications

- Chemotherapy toxicity
- Toxicity and efficacy of supportive care medication
- Targeted therapy selection
- Immunotherapy selection

WES=whole-exome sequencing; WGS=whole-genome sequencing; CTCs= circulating tumor cells; ctDNA=circulating tumor DNA; PCR=polymerase chain reaction; RT-PCR=reverse transcription polymerase chain reaction; NGS=next generation sequencing; MRD=minimal residual disease
Adapted from Sokolenko AP, Imyanitov EN. Front Mol Biosci. 2018 Aug 27;5:76



>50% of Advanced NSCLC Patients Harbor an Actionable Oncogenic Driver Alteration



NSCLC=non-small cell lung cancer; amp=amplification; mut=mutation; r=rearrangement

Adapted from Odintsov I, Sholl LM. Pathology. 2024;56(2):192-204.

National Cancer Institute. (n.d.). Targeted therapy drug list. National Institutes of Health.

<https://www.cancer.gov/about-cancer/treatment/types/targeted-therapies/approved-drug-list#targeted-therapy-approved-for-lung-cancer>



Real-World Biomarker Testing Rates and Turnaround Times Within a Large Community-Based Oncology Network

Retrospective observational chart review of patients with metastatic NSCLC (n=3474)



90% pts had testing for at least one biomarker, and **<50% received all 5 biomarker tests***



TAT from test order to results was ~2 weeks. Median time from diagnosis to 1L treatment was ~5 weeks



Only 35% had test results for **all 5 biomarkers** before 1L treatment



NGS testing rates increased with time, but **remained below 50%** over the study period

*EGFR, ALK, ROS1, BRAF, and PD-L1 were assessed
NSCLC=non-small cell lung cancer; TAT=turnaround time; 1L=first-line; pts=patients; NGS=next-generation sequencing
Robert NJ, Espirito JL, Chen L, et al. Lung Cancer. 2022 Apr;166:197-204



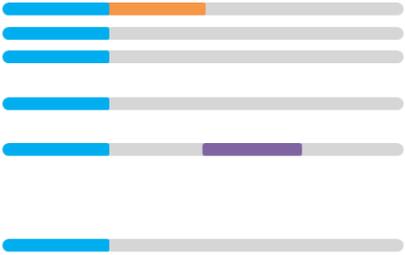
Comprehensive Genomic Profiling (CGP) can Identify More Clinically Relevant Variants than Conventional Testing Approaches

Single Marker Testing



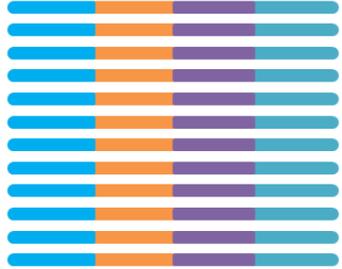
- Identifies mutations in a single gene using conventional approaches like FISH, PCR, or IHC¹
- Would miss clinically relevant mutations in other genes^{2,3}

Hotspot Panel

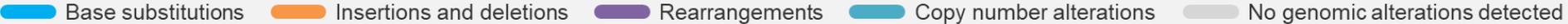


- Interrogates specific mutations in select genes^{4,5}
- Would miss other clinically relevant classes of alterations and mutations in other genes^{2,4-7}

Comprehensive Genomic Profiling

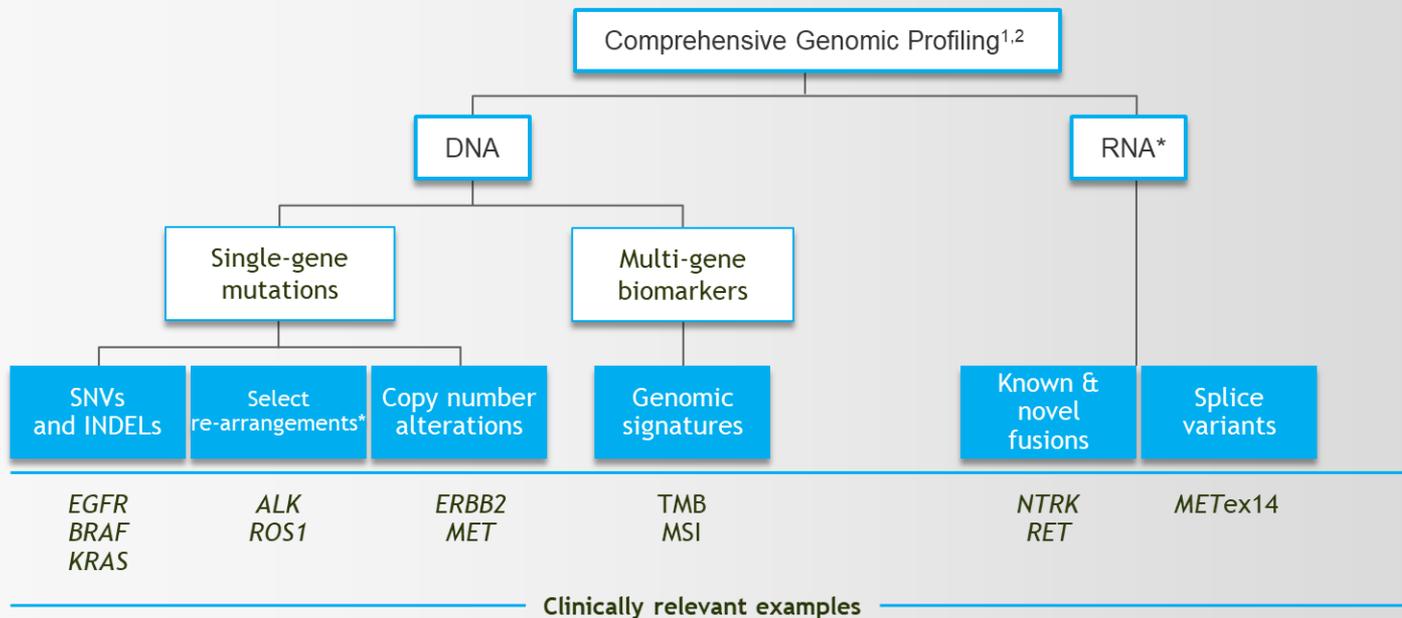


Analyzes all four types of genomic mutations across a large panel of cancer-related genes and genomic signatures (ie, TMB, MSI)^{8,9}



CGP=comprehensive genomic profiling; FISH=fluorescence in situ hybridization; IHC=immunohistochemistry; MSI=microsatellite instability; PCR=polymerase chain reaction; TMB=tumor mutational burden
1. Malone ER, Oliva M, Sabatini PJB, Stockley TL, Siu LL. *Genome Med.* 2020;12:8; 2. Reitsma M, Fox J, Borre PV, et al. *J Manag Care Spec Pharm.* 2019;25(5):601-611; 3. Ali SM, Hensing T, Schrock AB, et al. *Oncologist.* 2016;21:762-770; 4. Singh RR, Patel KP, Routbort MJ, et al. *J Mol Diagn.* 2013;15(5):607-622; 5. Kopetz S, Mills Shaw KR, Lee JJ, et al. *JCO Precis Oncol.* 2019;3:PO.18.00213; 6. Zehir A, Benayed R, Shah RH, et al. *Nat Med.* 2017;23(6):703-713; 7. Drilon A, Wang L, Arcila ME, et al. *Clin Cancer Res.* 2015;21(16):3631-3639; 8. Kroeze LI, de Voer RM, Kamping EJ, et al. *J Mol Diagn.* 2020;22(6):757-769; 9. Pestinger V, Smith M, Sillo T, et al. *Mol Diagn Ther.* 2020;24(3):339-349

Comprehensive Genomic Profiling (CGP) Facilitates Biomarker Testing



Advantages of CGP:

- ✓ Optimizes the yield of actionable findings^{1,2}
- ✓ Enables tissue stewardship³
- ✓ Detects rare or infrequent variants, including fusions²
- ✓ Broad genomic coverage for accurate calling of genomic signatures⁴

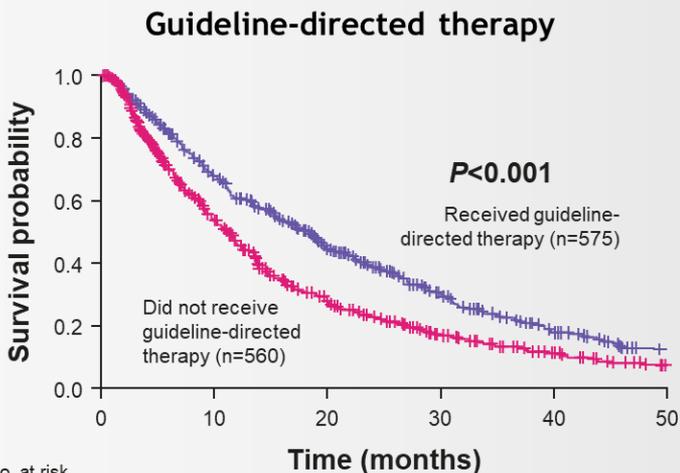
*Not all CGP assays bait for rearrangements in DNA and RNA

CGP=comprehensive genomic profiling; NGS=next-generation sequencing; INDELs=insertions and deletions; SNVs=single nucleotide variants; HRD=homologous recombination deficiency; TMB= tumor mutational burden; MSI=microsatellite instability

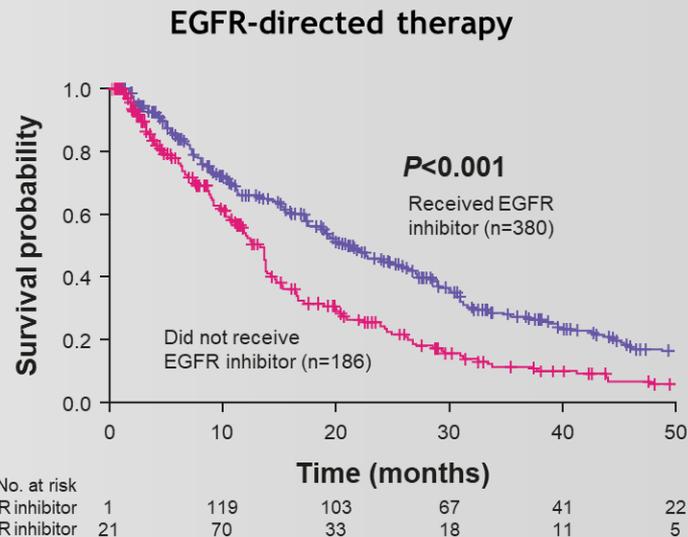
1. Kopetz S, Mills Shaw KR, Lee JJ, et al. JCO Precis Oncol. 2019;3:PO.18.00213; 2. Dilon A, Wang L, Arcila ME, et al. Clin Cancer Res. 2015;21(16):3631-3639; 3. Yu TM, Morrison C, Gold EJ, et al. Clin Lung Cancer. 2019 Jan;20(1):20-29.e8; 4. Chalmers ZR, Connelly CF, Fabrizio D, et al. Genome Med. 2017 Apr 19;9(1):34



Improved Survival in NSCLC Patients Receiving CGP with Genomically-Matched Targeted Therapies



	No. at risk					
	0	10	20	30	40	50
Received guideline-directed therapy	2	168	142	91	59	32
Did not receive guideline-directed therapy	55	196	93	54	28	12



	No. at risk					
	0	10	20	30	40	50
Received EGFR inhibitor	1	119	103	67	41	22
Did not receive EGFR inhibitor	21	70	33	18	11	5



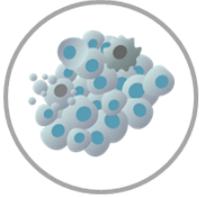


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Choosing tissue and/or liquid biopsy specimens

Biomarker Testing for NSCLC

Benefits & Limitations When Using Tissue or Liquid Biopsies for Biomarker Testing

	Advantages	Limitations
<p>Tumor tissue</p> 	<ul style="list-style-type: none">• Pathology information• PD-L1 assessment• High sensitivity• RNA for fusion detection	<ul style="list-style-type: none">• Invasive• Represents snapshot of tumor genomics• Longer TAT• Rebiopsy not always feasible
<p>Liquid biopsy</p> 	<ul style="list-style-type: none">• Less invasive, easy serial testing• Captures intra- & intertumoral heterogeneity• Rapid TAT	<ul style="list-style-type: none">• ctDNA fraction depends on tumor shed rate• Lower sensitivity for detecting all alteration types when ctDNA fraction is low• Result interference by CHIP

ctDNA=circulating tumor DNA; CHIP=clonal hematopoiesis of indeterminate potential; PD-L1=programmed cell death ligand 1; TAT=turnaround time.

Rolfo C, Mack P, Scagliotti GV, et al. J Thorac. Oncol. 2021;16(10):1647-1662.



Opportunities for Precision Medicine are Missed Up to 30% of the Time Due To Tissue Insufficiency

Frequency of tissue insufficiency



Tissue insufficiency due to *quantity or quality* is a barrier to NGS

Factors influencing QNS:

- Biopsy procedure³
- Tumor percentage⁴
- Tissue handling and fixation⁵
- Prior testing⁶

NGS=next-generation sequencing; NSCLC=non-small cell lung cancer; QNS=quantity not sufficient

1. Hagemann IS, Devarakonda S, Lockwood CM, et al. *Cancer*. 2015;121(4):631-639; 2. Aggarwal C, Thompson JC, Black TA, et al. *JAMA Oncol*. 2019;5(2):173-180; 3. Mata DA, Harries L, Williams EA, et al. *Arch Pathol Lab Med*. 2023;147(3):338-347; 4. Goswami RS, Luthra R, Singh RR, et al. *Am J Clin Pathol*. 2016;145(2):222-237; 5. Hussain M, Corcoran C, Sibilla C, et al. *Clin Cancer Res*. 2022;28(8):1518-1530; 6. Drilon A, Wang L, Arcila ME, et al. *Clin Cancer Res*. 2015;21(16):3631-3639

Factors Influencing ctDNA Levels and Detection

Disease characteristics

Tumor type

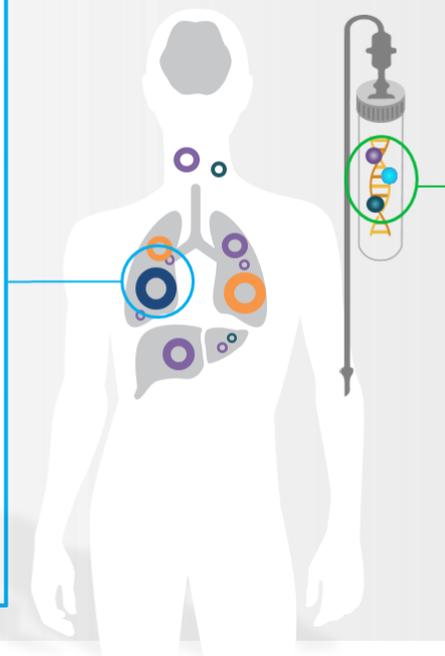
Tumor burden and staging

Disease status

(stable vs progressive disease)

Tumor microenvironment

(stroma and vascularization)



Other factors

Sample timing related to treatment or procedures

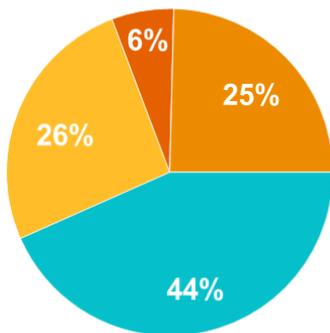
Sample acquisition, transport, and processing procedures

Type of variant

Assay sensitivity

Not all patients may be candidates for liquid biopsy at a given time

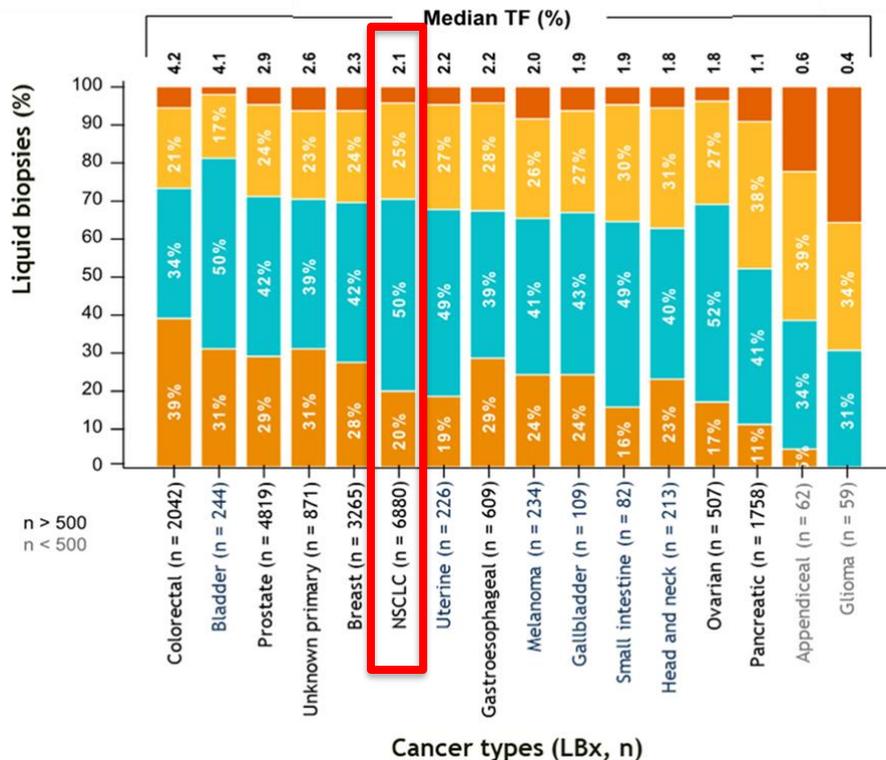
ctDNA Detection Rate Varies Across Tumor Types



Entire cohort (23,482)

ctDNA fraction

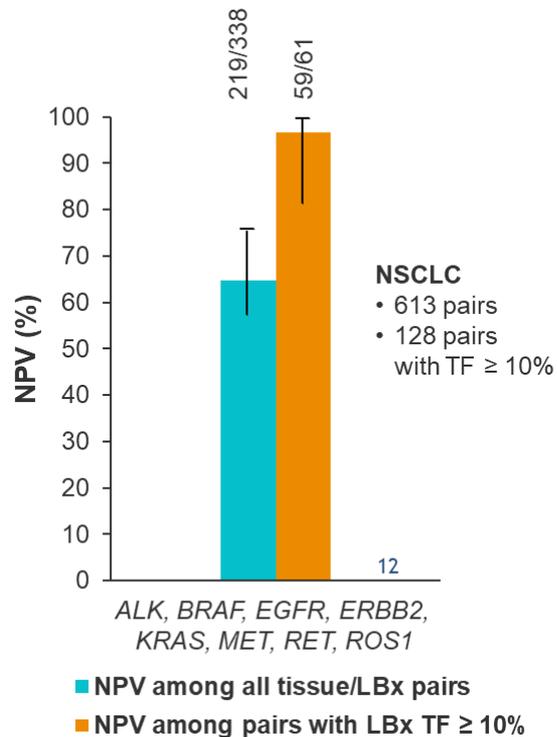
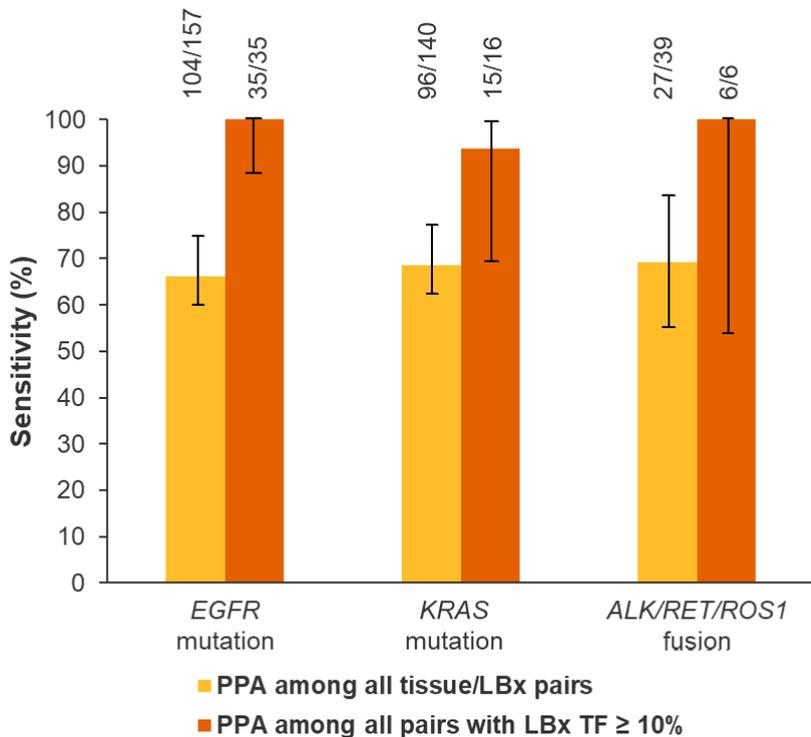
- ≥ 10%
- (1%-10%)
- 0%-1%
- Not detected



Cancer types (LBx, n)

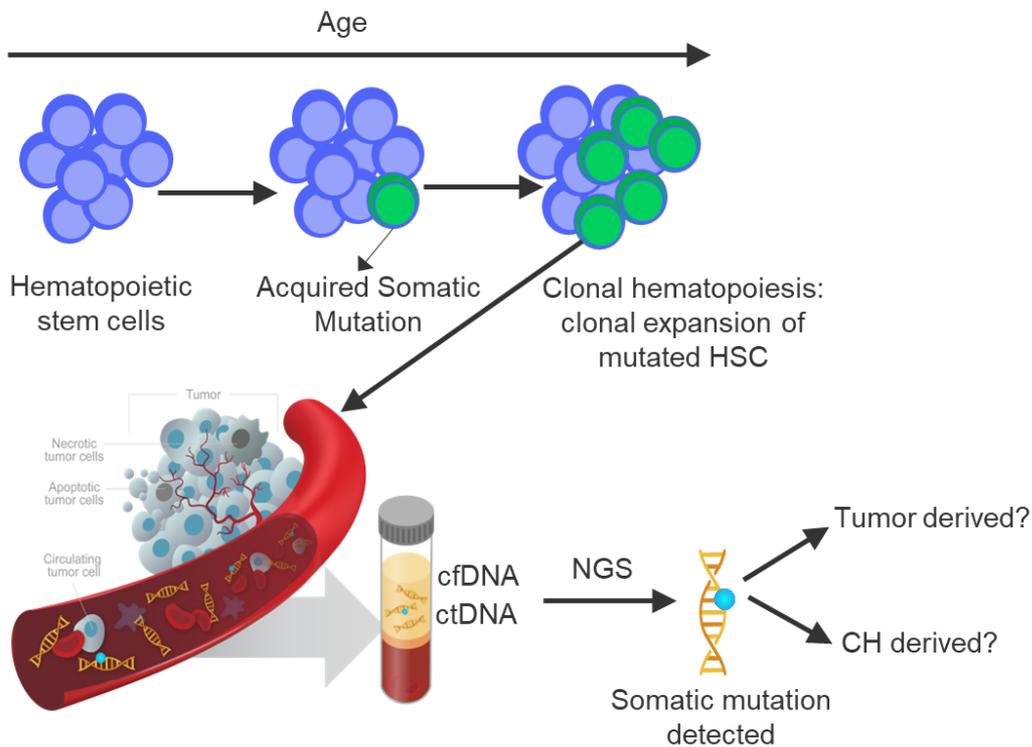
ctDNA=circulating tumor DNA; GIST=gastrointestinal stromal tumor; LBx=liquid biopsy; NSCLC=non-small cell lung cancer; SCLC=small cell lung cancer; TF=tumor fraction
 Husain H, Pavlick DC, Fendler BJ, et al. JCO Precis Oncol. 2022;6:e2200261

Sensitivity and Specificity With Liquid Biopsies is Higher in Samples with High ctDNA Fraction



Reflexing to tissue may be considered to confirm negative findings if the TF is low

Clonal Hematopoiesis Mutations Detected in Plasma From Liquid Biopsy Specimens Can Mimic Tumor-Derived Mutations



Clonal Hematopoiesis (CH)

- CH somatic mutations can occur in known oncogenes and are detected along with tumor-derived somatic mutations via NGS of liquid biopsy specimens
- Misclassification of CH mutations as tumor-derived (false-positive) may lead to inappropriate treatment decisions
- Cancer patients with CH are at risk of therapy-related myeloid neoplasms

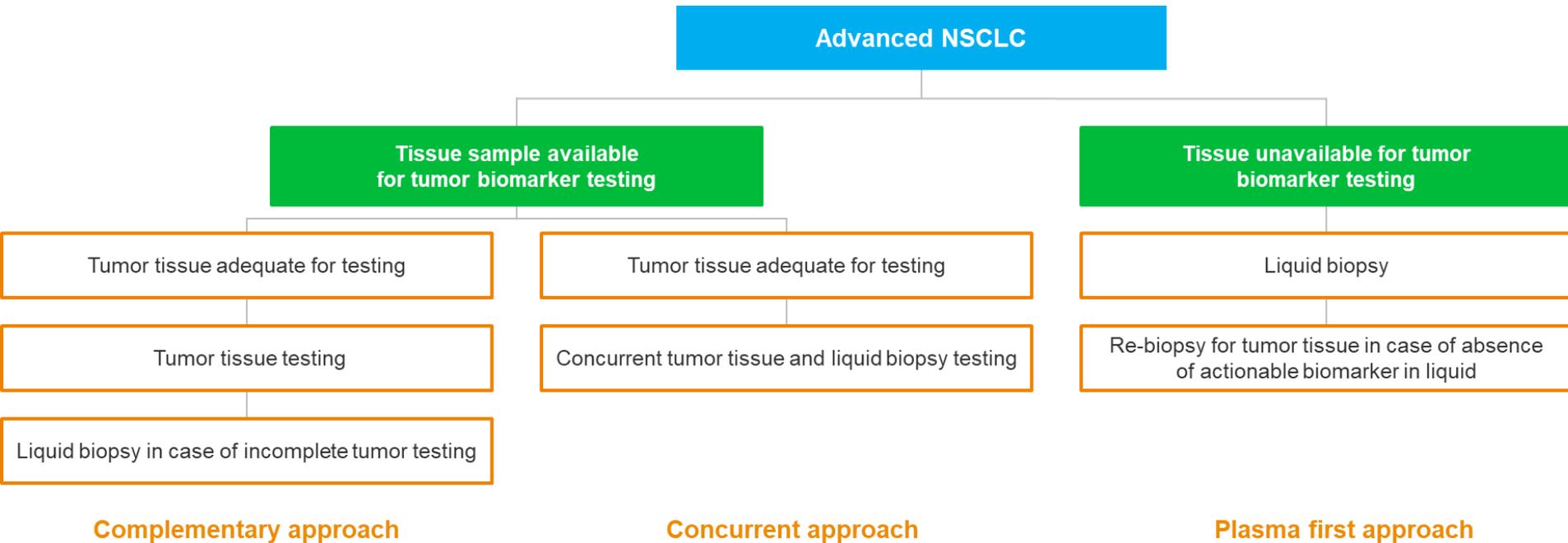


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Clinical utility of tissue & liquid biopsy specimens

Biomarker Testing for NSCLC

Guidance for Oncologists: International Association for the Study of Lung Cancer (IASLC)



Compromised Outcomes in Patients with NSCLC Treated Prior to the Report of an Actionable Oncogenic Driver (AOD) Alteration

Group A

(Comparator group)

379 pts treated after report of AOD

Group B

47 pts started treatment before report of AOD and switched to TKI within 35 days

Group C

84 pts started treatment before report of AOD and not switched to TKI within 35 days

	Group A (N=379)	Group B (N=47)	Group C (N=84)
OS			
Median (95% CI, months)	28.8 (23.3,34.6)	21.7 (12.2,NR)	15.3 (11.5,19.7)
Hazard ratio	Reference	1.12	1.62
P-Value	-	0.59	0.003
Time to next treatment			
Median (95% CI, months)	13 (10.5,15.5)	5.5 (3.4,8.9)	6.4 (4.9,8.7)
Hazard ratio	Reference	1.73	1.71
P-Value	-	0.002	0.0002

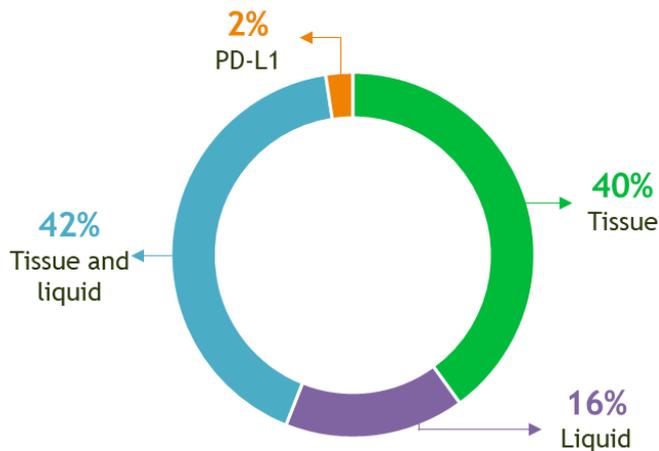


Outcomes are significantly compromised in patients harboring AOD but who are treated initially or even in patients quickly switched to TKI

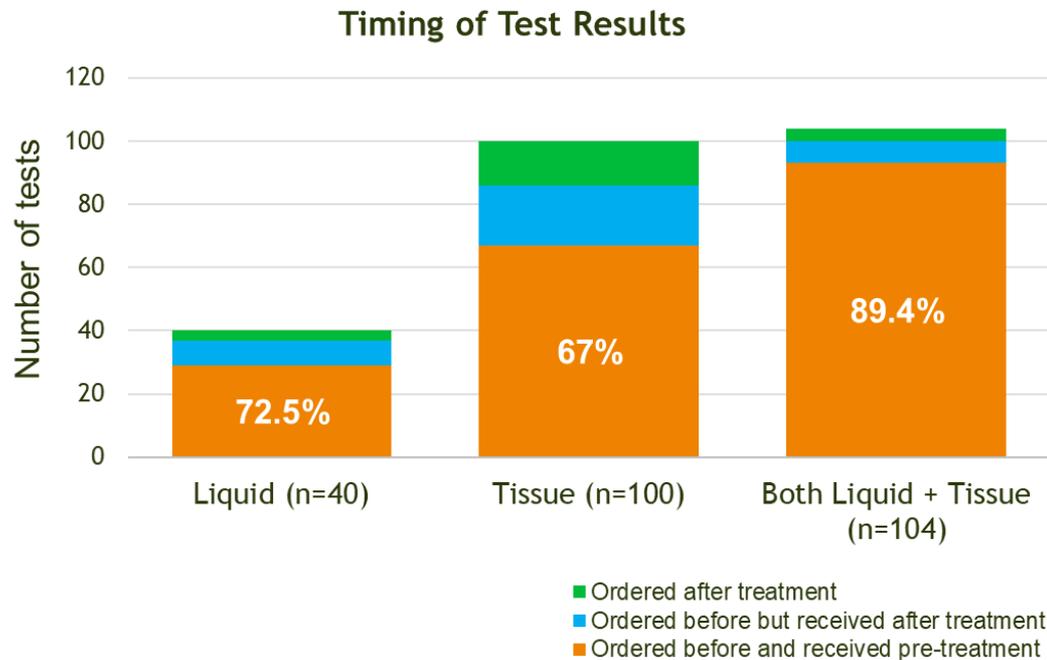
Time to next treatment: day1 line of treatment 1 to day 1 line of treatment 2 or apparent death; in group B is determined after the switch to TKI.
 NSCLC=non-small cell lung cancer; AOD=actionable oncogenic driver; TKI=tyrosine kinase inhibitor; OS=overall survival; CI=confidence interval
 Scott J.A., Lennerz J., Johnson M.L., et al. JCO Oncol Pract. 2023 Aug 9;OP2200611



Real-World Tissue and Liquid Biopsy Utilization in Advanced NSCLC in a Large Community US Practice



TAT (days)	Average	Median
Tissue (n=204)	16.7	11
Liquid (n=144)	7.9	7

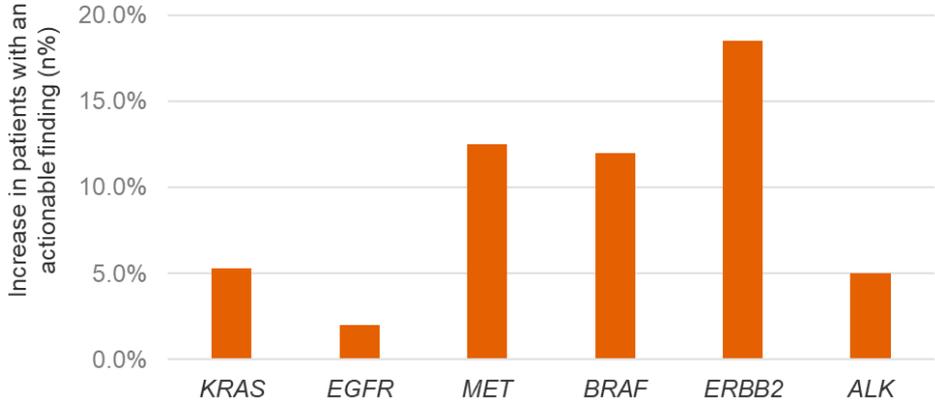


Concurrent Biomarker Testing With Tissue and Liquid Biopsies Leads to Increased Detection of Actionable Variants

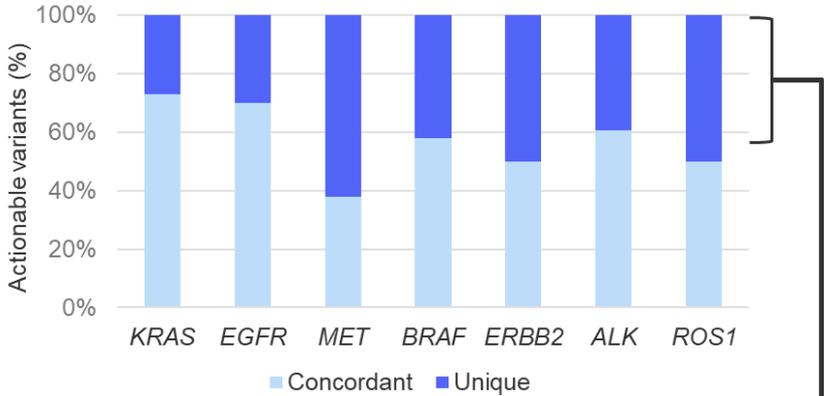
Actionable Variants in NSCLC (n=513)



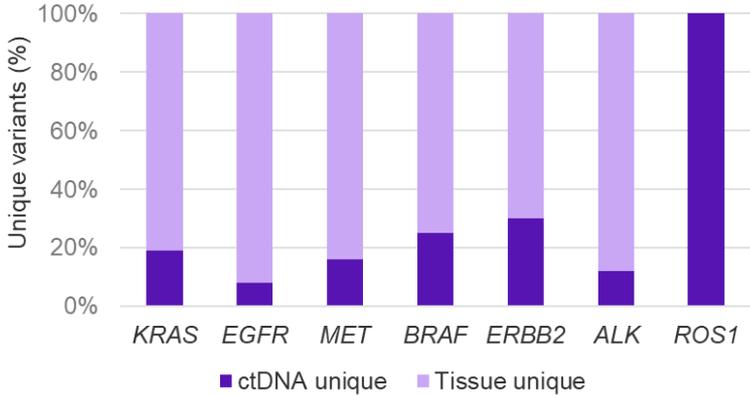
Increase in Patients With Actionable Variant



Actionable Variants Detected by Gene



Unique Variants Detected by Gene



NSCLC: Non-small cell lung cancer; ctDNA: circulating tumor DNA
 lams WT, Mackay M, Ben-Shachar R, et al. *JAMA Netw Open.* 2024;7(1):e2351700



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Coverage & Reimbursement

Biomarker Testing for NSCLC

Companion Diagnostics (CDx) Claims



are FDA approved by demonstrating analytical and clinical validity of the test through retrospective or prospective analyses, to accurately identify patients eligible to receive a targeted therapy for a defined biomarker



are supported by analytical validity of the test for each specific biomarker and a clinical study establishing either the **link between the result of that test and patient outcomes or clinical concordance to a previously approved CDx test**



Coverage of Tissue or ctDNA Testing in Advanced or Metastatic Cancer

Payer	CMP Coverage for tissue biopsy	Coverage for ctDNA	Allows for concurrent testing	Allows for repeat ctDNA testing
United Healthcare	CDx only	CDx OR <50 gene panels	Lung only	up to 3 times a year
Elevance (Anthem)	CMP for select tumors	CDx indications OR NSCLC, breast, and prostate*	No	Yes
Aetna	TMB only	Panels <50 genes for NSCLC	No	No
Cigna	CMP for select tumors	CDx only	No	No
Florida Blue	CDx only	CDX only	No	No
Horizon BCBS	CMP for select tumors	CDx only	No	No
BS CA	No limitations	covered for certain tumor types	Lung only	Not specified

Commercial Coverage Policy Examples



FDA Companion Diagnostic Testing medical coverage policy

CDx tests are proven and medically necessary when the oncology indication has a corresponding diagnostic test and biomarker on the US FDA list of cleared or approved Companion diagnostic devices. (Agnostic to tissue or liquid.)



Tumor Markers medical coverage policy

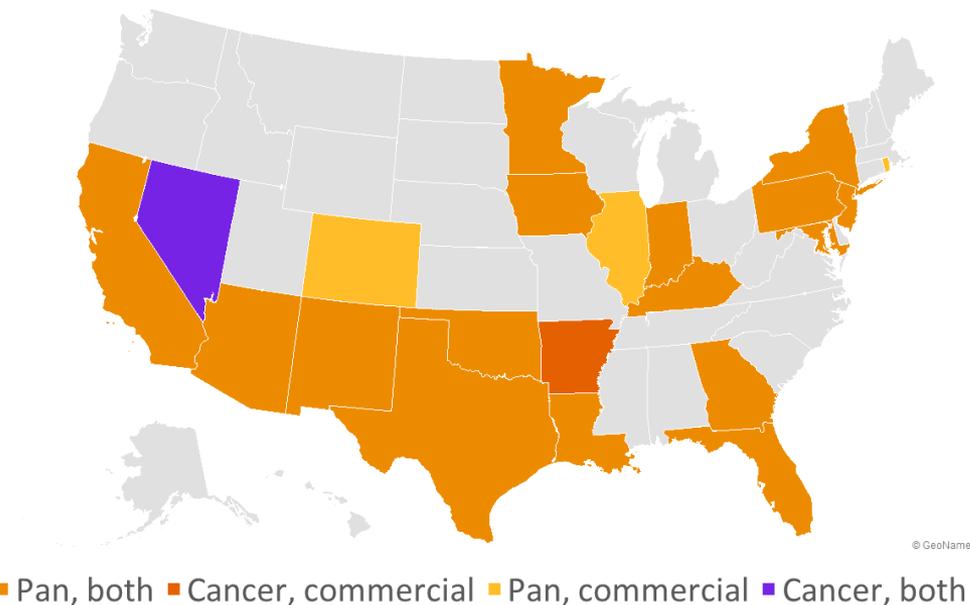
Circulating tumor DNA (ctDNA) (a type of liquid biopsy) for any indication (other than small panels, less than 50 genes, for non-small cell lung cancer), including, but not limited to, colorectal cancer, melanoma, ovarian cancer or prostate cancer are **considered experimental and investigational**.

ACS/CAN Biomarker Testing Bill Coverage in the US

Biomarker testing must be covered when the test is supported by medical and scientific evidence, including, but not limited to:

- Labeled indications for an FDA-approved or -cleared test;
- Indicated tests for an FDA-approved drug;
- Warnings and precautions on FDA-approved drug labels;
- CMS National Coverage or Medicare Administrative Contractor (MAC) Local Coverage of a test; or
- Nationally recognized clinical practice guidelines (and consensus statements).

Applicable to fully funded commercial insurers, Medicaid or both
"Pan" reflects coverage for all biomarkers not just cancer related



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Q&A

Thank you for attending!