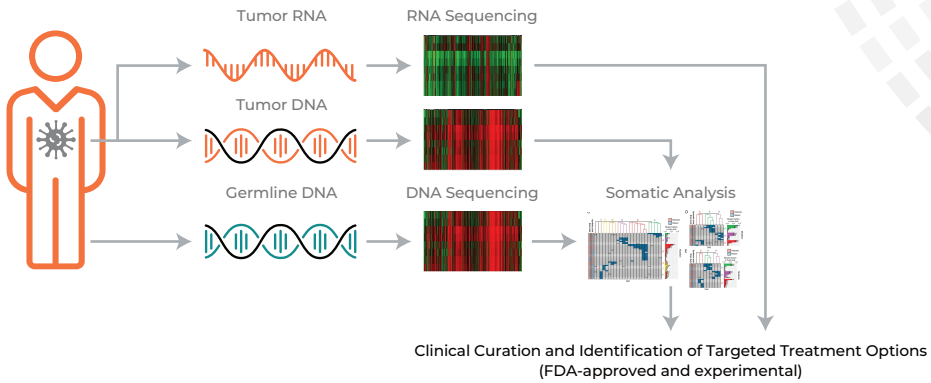
oncoExTra™ by EXACT SCIENCES

Covered by MEDICARE

Complete The Genomic Picture By Including DNA+RNA To Obtain The Most Actionable Insights For Therapy Selection

The OncoExTra™ test is an **ultra-comprehensive genomic profiling assay** that incorporates tumor whole-exome (DNA) and whole-transcriptome* (RNA) sequencing with paired tumor-normal analysis to identify alterations biomarkers in individuals diagnosed with advanced cancers. Findings are mapped to a knowledgebase of FDA-approved targeted treatment options as well as relevant clinical trial options.



WES (DNA) - Allows for comprehensive analysis of all protein-coding genes in a sample.

WTS (RNA) - Allows the identification of transcript variants and fusion genes that may be undetectable through conventional CGP tests which only employ DNA analysis.

Comprehensive Without Compromise

- The OncoExTra test interrogates ~20,000 genes.[3]
- IO signatures including tumor mutational burden (TMB) and microsatellite instability (MSI).
- 15 optional immunohistochemistry (IHC) stains† including PD-L1 (SP142, 22C3, SP263) and MMR (Mismatch Repair) proteins.
- Patient-matched tumor-normal sample to rule out benign variants.[3]

All About Actionability

- Reports clinically actionable mutations, copy number alterations, transcript variants/fusions through DNA and RNA analyses.
- FDA-approved therapies and clinical trial options based on the patient's results are also reported.
- In a clinical utilization study, at least one clinically actionable variant was identified in 83.9% of reports (1267/1509).[3]

*Whole-transcriptome sequencing with select variants reported in New York State
†IHC testing not currently available in New York State

According to one estimate, 20% of cancer morbidity occurs in tumors driven by translocations and gene fusions. Many of these alterations are actionable and may be missed by panel-based tests and WES alone.[1,2]

Case Study: OncoExTra™ detects fusion event in NSCLC

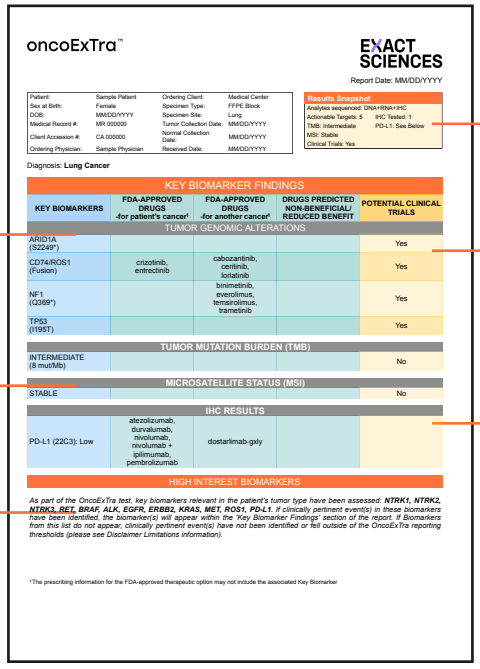
- A **64-Year-old- male** former smoker with a 50-pack year history presented to his primary physician with a 2-month history of fatigue, shortness of breath and unintentional weight loss.
- The patient previously had been offered lung cancer screening with a low dose CT scan of the chest but refused.
- A chest x-ray was ordered, and a **suspicious nodule was noted in the left lung upper lobe.** A CT of the chest demonstrated multiple nodules throughout both lungs suspicious for malignancy.
- CT guided lung biopsy **confirmed adenocarcinoma** consistent with pulmonary origin. PET/CT was completed and in addition to the lung lesions, 2 hypermetabolic hepatic lesions were also noted.
- **The OncoExTra test** was performed on the biopsy. It was discovered the **patient harbored ROS1 fusion.**
- **Tyrosine Kinase Therapy was initiated** and after **3 months repeat imaging demonstrated stable disease** throughout the chest and abdomen.

This case study is for educational purposes only and is not clinical, diagnostic, or treatment advice for any particular patient. Results and outcomes may vary. Providers should use their clinical judgment and experience when deciding how to diagnose or treat patients. Exact Sciences does not recommend or endorse any particular course of treatment or medical choice.

Targetable mutations and associated therapies

TMB & MSI Status

High Interest Biomarkers



oncoExTra™

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Report Date: MM/DD/YYYY

Patient:	Sample Patient	Ordering Client:	Medical Center
Sex at Birth:	Female	Specimen Type:	FFPE Block
DOB:	MM/DD/YYYY	Specimen Site:	Lung
Medical Record #:	MR 000000	Tumor Collection Date:	MM/DD/YYYY
Client Accession #:	CA 000000	Normal Collection Date:	MM/DD/YYYY
Ordering Physician:	Sample Physician	Received Date:	MM/DD/YYYY

Results Snapshot
Analytes sequenced: DNA+RNA+IHC
Actionable Targets: 5 IHC Tested: 1
TMB: Intermediate PD-L1: See Below
MSI: Stable
Clinical Trials: Yes

Diagnosis: **Lung Cancer**

KEY BIOMARKER FINDINGS

KEY BIOMARKERS	FDA-APPROVED DRUGS -for patient's cancer‡	FDA-APPROVED DRUGS -for another cancer‡	DRUGS PREDICTED NON-BENEFICIAL/ REDUCED BENEFIT	POTENTIAL CLINICAL TRIALS
TUMOR GENOMIC ALTERATIONS				
ARID1A (S2249*)				Yes
CD74/ROS1 (Fusion)	crizotinib, entrectinib	cabozantinib, ceritinib, lorlatinib		Yes
NF1 (Q369*)		binimetinib, everolimus, temsirolimus, trametinib		Yes
TP53 (I195T)				Yes
TUMOR MUTATION BURDEN (TMB)				
INTERMEDIATE (6 mut/Mb)				No
MICROSATELLITE STATUS (MSI)				
STABLE				No
IHC RESULTS				
PD-L1 (22C3): Low	atezolizumab, durvalumab, nivolumab, nivolumab + ipilimumab, pembrolizumab	dostarlimab-gxly		
HIGH INTEREST BIOMARKERS				

*As part of the OncoExTra test, key biomarkers relevant in the patient's tumor type have been assessed: **NTRK1, NTRK2, NTRK3, RET, BRAF, ALK, EGFR, ERBB2, KRAS, MET, ROS1, PD-L1**. If clinically pertinent event(s) in these biomarkers have been identified, the biomarker(s) will appear within the 'Key Biomarker Findings' section of the report. If Biomarkers from this list do not appear, clinically pertinent event(s) have not been identified or fell outside of the OncoExTra reporting thresholds (please see Disclaimer Limitations information).*

‡The prescribing information for the FDA-approved therapeutic option may not include the associated Key Biomarker

Snapshot of Key Findings

Clinical Trial Options

IHC Results†

To Learn More: OncoExTra.com | To Order: OncoExTra.com/order

References: 1. Drenner, Basu GD, Goodman LJ, et al. The value of comprehensive genomic sequencing to maximize the identification of clinically actionable alterations in advanced cancer patients: a case series. Oncotarget. 2021; 12:1836-1847. **2.** Nikanjam M, Okamura R, Barkauskas DA, Kurzrock R. Targeting fusions for improved outcomes in oncology treatment. Cancer. 2020; 126:1315-1321. **3.** White T, Szelinger S, LoBello J, et al. Analytic validation and clinical utilization of the comprehensive genomic profiling test, Oncotarget 2021;12: 726-739

Disclaimer: The OncoExTra test is not a FDA cleared or approved IVD device or companion diagnostic for the referenced biomarkers and FDA approved therapies.

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OncoExTra has been validated according to the guidelines set forth by the New York State Department of Health. Whole exome (DNA) events have been validated to include point mutations, indels, and copy number alterations, as well as MSI analysis and TMB calculation. Whole transcriptome (RNA) has been validated to report on select fusion genes and special transcripts.

†IHC testing not currently available in New York State

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