



Comprehensive Tumor Profiling

A better guide for
Molecular Intelligence





Why Caris?

With the promise of precision medicine becoming a reality, molecular profiling has become standard of care for many cancer types – and required for certain therapies (companion dx). More than ever, oncologists need a trusted tumor profiling partner to provide reliable, high-quality molecular information to guide more precise and individualized treatment decisions.

Caris Molecular Intelligence® identifies key molecular features of cancer and provides clinical insights to aid oncologists in personalizing cancer therapies for their solid tumor cancer patients. Caris Molecular Intelligence helps oncologists:

- **navigate among therapies with potential benefit;**
- **identify therapies that may not have been considered;**
- **determine drugs with potential lack of benefit (avoiding unnecessary toxicities and costs); and**
- **match patients to clinical trials.**

The Most Experienced Comprehensive Tumor Profiling Provider



Unmatched Experience

- 2,850,000+ tests completed
- 240,000+ clinical cases performed



Immunotherapy Diagnostics Expertise

- PD-L1 by IHC, including TPS and CPS scoring methods
- Mismatch Repair (MMR) proteins by IHC: MLH1, MSH2, MSH6, PMS2
- Microsatellite Instability (MSI) by NGS
- Tumor Mutational Burden (TMB) by NGS



Rigorous Quality Standards

- CAP, CLIA, NYSDOH, ISO15189 accredited
- 66,000-square-foot clinical laboratory in Phoenix, AZ
- 38,000 square-foot R&D facility in Phoenix, AZ
- 115,000-square-foot blood-based laboratory in Irving, Texas (to open in 2021)
- Staffed by: bioinformaticians, oncologists, molecular geneticists, pathologists and PhD scientists



Limited Tissue Capabilities

- Tumor enrichment via microdissection
- Multiple reflex options to alternative technologies/methods



Rapid Turnaround Time

- 8-14 calendar days

→ www.CarisLifeSciences.com



Tumor Profiling

Comprehensive tumor profiling with Caris Molecular Intelligence® assesses DNA, RNA and Proteins with multiple technologies (NGS DNA-WES & RNA-WTS, PyroSeq, IHC, ISH) to reveal a more complete molecular blueprint to guide precise and individualized treatment decisions.



DNA

Next-Generation Sequencing
(Mutations, Indels &
Copy Number Alterations)

Whole Exome
~22,000 genes

- ✓ 200+ clinically relevant genes
- ✓ 700+ genes sequenced at average 500x and balance of exome at average 200x
- ✓ ~250,000 exonic/ intronic/ intergenic SNPs

All solid tumors



RNA

Next-Generation Sequencing
(Fusions & Variant Transcripts)

Whole Transcriptome
~22,000 genes

- ✓ 60 million read counts
- ✓ Gene fusions and variant transcripts
- ✓ Novel translocation detection independent of intronic breakpoint or fusion partner

All solid tumors



Protein

Immunohistochemistry

Up to 15
Clinically Relevant IHCs
(Optimized across
~25 Tumor Types)

- ✓ Multiple FDA approved CDx assays for different disease types (per label)
- ✓ Controls on every IHC slide
- ✓ 4 µm cuts to preserve tissue

All solid tumors

Caris Molecular Intelligence offers two different options
for tumor profiling, MI Profile™ and MI Tumor Seek™.

MI PROFILE™

MI TUMOR Seek™

Whole Genome (NGS-DNA)	✓	✓
MSI	✓	✓
TMB	✓	✓
LOH	✓	✓
HRD	Coming Soon	
Whole Transcriptome (NGS-RNA)	✓	✓
IHC, ISH (Protein)	✓	Not Included
TAT	8-14 days	



Caris MAI™ – Molecular Artificial Intelligence

A revolutionary platform to identify unique “molecular signatures” using artificial intelligence, machine learning and multivariate analysis across the Caris clinical-molecular database to push the boundaries of innovation in precision medicine.



Genomic Prevalence Score

- Intended for cancer of unknown primary (CUP), or cases with atypical clinical presentation or clinical ambiguity.
- Provides a cancer type similarity assessment by comparing genomic (DNA) and transcriptomic (RNA) characteristics of the patient's tumor against other tumors in the Caris database.

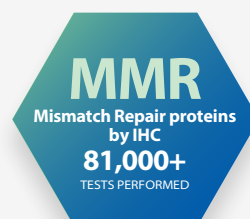
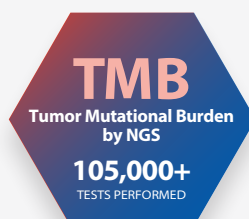
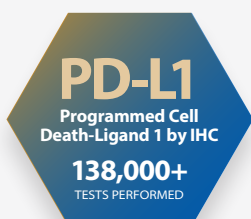


Frontline Therapy Predictor

- Intended as an aid in gauging a patient's likelihood to benefit from FOLFOX chemotherapy (in combination with bevacizumab) as the first-line chemotherapy regimen in metastatic colorectal adenocarcinoma.
- Outcomes data from the MI FOLFOXai validation demonstrated a ~50% improvement of overall survival.



Immunotherapy Diagnostics Expertise

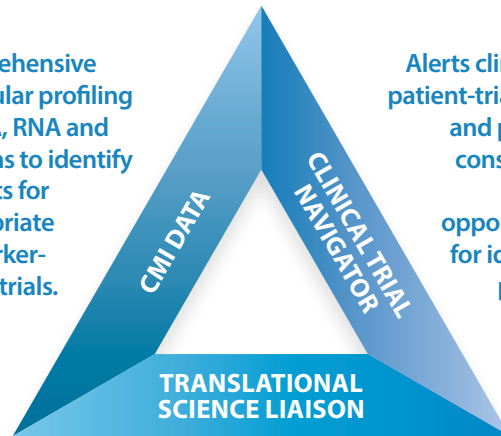


By harnessing the body's immune system to detect and destroy tumor cells, immune checkpoint inhibitors are rapidly ushering in a new era of precision medicine. Identify patients more likely to respond to immunotherapies with reliable molecular information based on validated assays for MMR, MSI, PD-L1 and TMB in our state-of-the-art laboratory.



MI Trials provides:

- Comprehensive
molecular profiling
of DNA, RNA and
proteins to identify
patients for
appropriate
biomarker-
driven trials.**



Alerts clinician of patient-trial match and provides consultation on trial opportunities for identified patients.




MI Report


Contains
important
biomarker and
associated
therapy results

Provides important information about drug/ biomarker associations and comments from Caris pathologists and/or molecular geneticists

Lists disease-relevant biomarker information, as well as other altered biomarker results



Final Report



Patient

Name: Patient, Test
Date of Birth: XX/Mon/19XX
Sex: Male
Case Number: TN19-XXXXXX
Diagnosis: Mucinous adenocarcinoma

Specimen Information

Primary Tumor Site: Transverse colon
Specimen Site: Liver
Specimen ID: ABC-1234-XYZ
Specimen Collected: XX-Mon-2019
Completion of Testing: XX-Mon-2019

Ordered By

Ordering Physician, MD
 Cancer Center
 123 Main Street
 Springfield, XY 12345, USA
 1 (123) 456-7890

High Impact Results

BIOMARKER	METHOD	RESULT	THERAPY ASSOCIATION	BIOMARKER LEVEL*
Mismatch Repair Status	IHC	Deficient	<div>BENEFIT</div> <div>BENEFIT</div> <div>LACK OF BENEFIT</div> <div>BENEFIT</div>	Level 1
MSI	NGS	High		
BRAF	NGS	Mutated, Pathogenic Exon 15 p.V600E		Level 2
ERBB2 (Her2/Neu)	CISH	Amplified		Level 3A
			lapatinib, pertuzumab, trastuzumab	Level 3A

* Biomarker reporting classification: Level 1 - highest level of clinical evidence and/or biomarker association included on the drug label; Level 2 - strong evidence of clinical significance and is endorsed by standard clinical guidelines; Level 3 - potential clinical significance (3A - evidence exists in patient's tumor type; 3B - evidence exists in another tumor type).


Important Note

This patient has a potential NCI-MATCH Trial-eligible result. Please see Clinical Trial see page 6

Additional Results

CANCER TYPE RELEVANT BIOMARKERS		
Biomarker	Method	Result
NTRK1	RNA-Seq	Fusion Not Detected
NTRK2	RNA-Seq	Fusion Not Detected
NTRK3	RNA-Seq	Fusion Not Detected
Tumor Mutational Burden		
ERBB2 (p=...)		

CANCER TYPE RELEVANT BIOMARKERS	
Biomarker	Result
PTEN	IH
OTHER FINDINGS	
Biomarker	Result




Navigate among therapies with potential benefit (noted in green)


- Identify therapies with potential lack of benefit (noted in red)

Utilize the Clinical Trials Connector to match biomarkers with open clinical trials through MI Portal




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