



Comprehensive Molecular Profiling *Tumor Tissue or Whole Blood*

Caris Life Sciences® performs comprehensive molecular profiling to assess biomarkers in both tissue and blood, helping to guide more precise and individualized treatment decisions for cancer patients.

TISSUE-BASED MOLECULAR PROFILING



DNA

Whole Exome Sequencing

SNVs, Indels, CNAs, Karyotyping, Viruses



RNA

Whole Transcriptome Sequencing

Gene Fusions, Gene Expression and Variant Transcripts

AVAILABLE WITH MI PROFILE™



Protein

Immunohistochemistry

Tumor-Relevant Protein Biomarkers

Caris GPSai™

Cancer type similarity assessment intended to help identify the tumor of origin by comparing molecular characteristics of the patient's tumor against 90 tumor categories in the Caris database.

Caris FOLFIRSTai™

Chemotherapy response predictor intended to gauge a mCRC patient's likelihood of benefit from first-line FOLFOX+BV followed by FOLFIRI+BV, versus FOLFIRI+BV followed by FOLFOX+BV treatment.

BLOOD-BASED MOLECULAR PROFILING



FROM PLASMA



Somatic Tumor

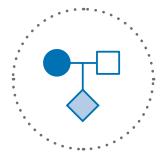
Tumor-derived mutations

FROM BUFFY COAT WBCs



Incidental CH

Non-tumor mutations (false-positives)



Incidental Germline

Hereditary variants



Shipper kits for specimen transportation to our lab is an important part of the molecular profiling process. Shipper kits and requisitions can be ordered on-demand via QR code.





MI Profile™ Comprehensive Testing (Tissue)

MI Tumor Seek Hybrid™ + IHCs and Other Tests by Tumor Type. Tissue-based Whole Exome and Whole Transcriptome Sequencing analysis, plus additional tumor-type relevant biomarker testing (IHC, ISH, etc.). Caris FOLFIRSTai™ is performed for mCRC cases and Caris GPSai™ is performed for CUP cases.

Biological Coverage

(DNA) SNVs, InDels, CNAs, Karyotyping, Viruses
(RNA) Gene Fusions, Gene Expression and Variant Transcripts
(Protein) 15+ Tumor-specific antigens

Variant Coverage

Somatic Tumor

Technologies

NGS IHC PyroSeq CISH

Next-Generation Sequencing

Whole Exome (DNA) Whole Transcriptome (RNA)

Genes & Depth

23,000+ 1500x

Genomic Signatures/Other

gLOH HRD MSI TMB HLA Genotype

Additional tumor-type relevant biomarker testing:

| Tumor Type | Immunohistochemistry (IHC) | Other |
|--|--|--|
| Bladder | Her2/Neu, PD-L1 (22c3) | |
| Breast | AR, ER, Her2/Neu, PD-L1 (22c3), PR, PTEN | |
| Cancer of Unknown Primary – Female | AR, ER, Her2/Neu, PD-L1 (SP142) | |
| Cancer of Unknown Primary – Male | AR, Her2/Neu, PD-L1 (SP142) | |
| Cervical | Her2/Neu, PD-L1 (22c3) | |
| Cholangiocarcinoma/ Hepatobiliary | Her2/Neu, PD-L1 (SP142) | |
| Colorectal and Small Intestinal | Her2/Neu, MMR, PD-L1 (SP142) | |
| Endometrial (all except uterine sarcoma) | ER, Her2/Neu, MMR, PD-L1 (SP142), PR | |
| Esophageal Cancer | Her2/Neu, PD-L1 (22c3) | |
| Gastric/GEJ | CLDN18, Her2/Neu, MMR, PD-L1 (22c3) | EBER (Chromogenic <i>in situ</i> Hybridization) |
| GIST | Her2/Neu, PD-L1 (SP142) | |
| Glioma | Her2/Neu | MGMT Methylation (Pyrosequencing) |
| Head & Neck | Her2/Neu, p16, PD-L1 (22c3) | EBER, HPV (Chromogenic <i>in situ</i> Hybridization), HPV reflex to confirm p16 result |
| Kidney | Her2/Neu, PD-L1 (SP142) | |

Specimen Quantity

20% Tumor (NGS) | 10 slides (NGS only) 25 slides (NGS+IHC)

Clinical AI

Caris FOLFIRSTai™ Caris GPSai™

Viruses

HPV 16 & 18 (Head & Neck, Anal, Genital, CUP)

EBV (Head & Neck, Esophagogastric Junction, Gastric Adenocarcinoma, CUP. If positive result in cancer type not listed here, EBER ISH reflex to confirm EBV result)

MCPyV (Merkel Cell, Neuroendocrine – Poorly Differentiated (High-Grade)/Large or Small Cell Carcinoma, CUP)

Chromosomal Alterations

+7/-10 and 1p19q co-deletion (glioma)

| Tumor Type | Immunohistochemistry (IHC) | Other |
|---------------------|---|-------|
| Melanoma | Her2/Neu, PD-L1 (SP142) | |
| Merkel Cell | Her2/Neu, PD-L1 (SP142) | |
| Neuroendocrine | Her2/Neu, PD-L1 (SP142) | |
| Non-Small Cell Lung | ALK*, Her2/Neu, PD-L1 (22c3, 28-8, SP142, SP263) | |
| Ovarian | ER, FOLR1 [†] , Her2/Neu [†] , PD-L1 (22c3), PR | |
| Pancreatic | Her2/Neu, PD-L1 (SP142) | |
| Prostate | AR, Her2/Neu, PD-L1 (SP142) | |
| Salivary Gland | AR, Her2/Neu, PD-L1 (SP142) | |
| Sarcoma | Her2/Neu, PD-L1 (SP142) | |
| Small Cell Lung | Her2/Neu, PD-L1 (22c3) | |
| Thyroid | Her2/Neu, PD-L1 (SP142) | |
| Uterine Sarcoma | ER, Her2/Neu, MMR, PD-L1 (SP142), PR | |
| Vulvar Cancer (SCC) | Her2/Neu, PD-L1 (22c3) | |
| Other Tumors | Her2/Neu, PD-L1 (SP142) | |

MMR = Mismatch Repair proteins: MLH1, MSH2, MSH6, PMS2

*ALK IHC only performed for NSCLC adenocarcinoma.

[†]FOLR1 IHC and HRD Status only performed for epithelial ovarian cancer.

Availability of certain assays and features may vary by location. Check website for complete details.



MI Tumor Seek Hybrid™ (Tissue)

Tissue-based Whole Exome and Whole Transcriptome Sequencing analysis. Caris FOLFIRSTai™ is performed for mCRC cases and Caris GPSai™ is performed for CUP cases.

Biological Coverage

(DNA) SNVs, InDels, CNAs, Karyotyping, Viruses
(RNA) Gene Fusions, Gene Expression and Variant Transcripts

Variant Coverage

Somatic Tumor

Technologies

NGS

Next-Generation Sequencing

Whole Exome (DNA) Whole Transcriptome (RNA)

Genes & Depth

23,000+ 1500x

Genomic Signatures/Other

gLOH HRD MSI TMB HLA Genotype

Specimen Quantity

20% Tumor (NGS) | 10 slides

Clinical AI

Caris FOLFIRSTai™ Caris GPSai™

Viruses

HPV 16 & 18 (Head & Neck, Anal, Genital, CUP)

EBV (Head & Neck, Esophagogastric Junction, Gastric Adenocarcinoma, CUP. If positive result in cancer type not listed here, EBER ISH reflex to confirm EBV result)

MCPyV (Merkel Cell, Neuroendocrine – Poorly Differentiated (High-Grade)/Large or Small Cell Carcinoma, CUP)

Chromosomal Alterations

+7/-10 and 1p19q co-deletion (glioma)



Caris Assure™ (Blood)

Blood-based Whole Exome and Whole Transcriptome Sequencing for pathogenic and likely pathogenic tumor-derived, incidental germline, and incidental CH variant detection.

Technology

Circulating Nucleic Acids Sequencing (cNAS)

Application

Biomarker Analysis
(including resistance mutations)

Biological Coverage

Plasma: cfDNA, cfRNA
White Blood Cells: gDNA, mRNA

Variant Coverage (pathogenic and likely pathogenic)

Somatic Tumor-Derived
Incidental Germline[†]
Incidental CH

Genes & Depth

23,000+
8,000x (raw average for clinically relevant genes)

Next-Generation Sequencing

Whole Exome
Whole Transcriptome

Alterations

SNV INDEL CNA Fusions

Genomic Signatures / Other

bTMB MSI HLA Genotype

Sample Quantity

2 Tubes Whole Blood

Performance in Advanced/Metastatic Patients

Compared to matched tissue collected within 30 days; based on ≥5 ng of cNAS input. Minimum reportable allele frequency is 0.1%.

Clinically Actionable SNV and INDEL:

Sensitivity 93.8% PPV 96.8%
Specificity >99.9%

Incidental Germline[†]:

Sensitivity >99% PPV >99%
Specificity >99%

[†]Not a replacement for comprehensive germline testing. Incidental pathogenic alterations detected in ACMG recognized cancer genes and others are reported. Negative results do not imply the patient does not harbor a germline mutation.

Caris Assure™ is intended for patients with previously diagnosed solid malignant neoplasms when tissue is not feasible and is to be used by qualified healthcare professionals. RNA results are intended for investigational purposes only. Not available in all locations. Check website for complete details.

Comprehensive Support

Caris is committed to providing the highest quality of support for physicians and patients. A dedicated multifunctional local team provides a wide range of support including case management, continuing education, investigator-led research support and assistance with molecular tumor boards.

Account Management

Coordinates case management, site training, logistics, online ordering/portal needs.

Strategic Accounts Management

Partners with senior leaders for broader, system-wide projects and contracting needs.

Customer Support

Facilitates case setup, ordering support (kits, questions), patient billing information, etc.

Account Implementation Team

Collaborates with hospital administration and laboratory for service agreements and contracting needs.



Molecular Oncology Specialists

Provides product updates and information, testing capabilities and issue management.

Precision Medicine Initiatives Team

Assists with technical lab workflows and efficiencies.

Molecular Science Liaisons (MDs/PhDs)

Supports technical, scientific and medical educational needs for precision medicine.

Precision Oncology Program Team

Implements EHR integrations and other electronic ordering and reporting initiatives.

Caris Molecular Testing – Complete Gene Coverage

As the pioneer in precision medicine, Caris was the first to provide WES and WTS for every patient. All molecular profiling orders include next-generation sequencing of 23,000+ genes. Listed below are the genes most commonly associated with cancer. Full gene search is available on CarisLifeSciences.com.

| | | | | | | | | | |
|----------|--------|---------|---------|----------|----------------------|--------|---------|---------|----------|
| ABL1 | BCR | CSF1R | FANCC | GNAQ | LZTR1 | MUTYH | PIK3CB | RAD51D | SOCS1 |
| ABL | BLM | CTNNA1 | FANCD2 | GNAS | MAML2 | MYB | PIK3R1 | RAD54L | SPEN |
| ACVR1 | BMPR1A | CTNNB1 | FANCE | H3F3A | MAP2K1 | MYC | PIK3R2 | RAF1 | SPOP |
| AIP | BRAF | CXCR4 | FANCF | H3F3B | MAP2K2 | MYCN | PIM1 | RASA1 | SRC |
| AKT1 | BRCA1 | CYLD | FANCG | HDAC1 | MAP2K4 | MYD88 | PKN1 | RB1 | SSBP1 |
| AKT2 | BRCA2 | CYP17A1 | FANCI | HIST1H3B | MAP3K1 | NBN | PMS1 | RELA | STAG2 |
| AKT3 | BRD3 | DDR2 | FANCL | HIST1H3C | MAPK1 | NF1 | PMS2 | RET | STAT3 |
| ALK | BRD4 | DICER1 | FANCLM | HNF1A | MAPK3 | NF2 | POLD1 | RHOA | STK11 |
| AMER1 | BRIP1 | DNMT3A | FAS | HOXB13 | MAST1 | NFE2L2 | POLD2 | RNF43 | SUFU |
| APC | BTX | EGFR | FAT1 | HRAS | MAST2 | NFKBIA | POLD3 | ROS1 | TERT |
| AR | CALR | EGFRV8 | FBXW7 | IDH1 | MAX | NOTCH1 | POLD4 | RPA1 | TET2 |
| ARAF | CARD11 | EGLN1 | FGFR1 | IDH2 | MED12 | NOTCH2 | POLE | RPA2 | TFE3 |
| ARHGAP26 | CASP8 | ELF3 | FGFR2 | INSR | MEF2B | NPM1 | POLQ | RPA3 | TFFB |
| ARHGAP35 | CBFB | EP300 | FGFR3 | IRF4 | MEN1 | NRAS | POT1 | RPA4 | THADA |
| ARID1A | CCND1 | EPHA2 | FGFR4 | JAK1 | MET | NRG1 | PPARG | RSPO2 | TMEM127 |
| ARID2 | CCND2 | ERBB2 | FGR | JAK2 | MET Exon 14 Skipping | NSD1 | PPP2R1A | RSPO3 | TMPRSS2 |
| AR-V7 | CCND3 | ERBB3 | FH | JAK3 | MGA | NTHL1 | PPP2R2A | RUNX1 | TNFAIP3 |
| ASXL1 | CD274 | ERBB4 | FLCN | KDM5C | MGMT | NTRK1 | PRDM1 | SDHA | TNFRSF14 |
| ATM | CD79B | ERCC2 | FLT1 | KDM6A | MITF | NTRK2 | PRKACA | SDHAF2 | TP53 |
| ATR | CDC73 | ERG | FLT3 | KDR | MLH1 | NTRK3 | PRKAR1A | SDHB | TRAF7 |
| ATRX | CDH1 | ESR1 | FLT4 | KEAP1 | MLH3 | NUMBL | PRKCA | SDHC | TSC1 |
| AXIN1 | CDK12 | ETV1 | FOXA1 | KIF1B | MPL | NUTM1 | PRKCB | SDHD | TSC2 |
| AXIN2 | CDK4 | ETV4 | FOXL2 | KIT | MRE11 | PALB2 | PTCH1 | SETD2 | U2AF |
| AXL | CDK6 | ETV5 | FUBP1 | KLF4 | MSH2 | PARP1 | PTEN | SF3B1 | VHL |
| B2M | CDKN1B | ETV6 | FYN | KMT2A | MSH3 | PBRM1 | PTPN11 | SMAD2 | WRN |
| BAP1 | CDKN2A | EWSR1 | GALNT12 | KMT2C | MSH6 | PCNA | RABL3 | SMAD4 | WT1 |
| BARD1 | CHEK1 | EXO1 | GATA3 | KMT2D | MSMB | PDGFRA | RAC1 | SMARCA4 | XPO1 |
| BCL2 | CHEK2 | EZH2 | GLI2 | KRAS | MST1R | PDGFRB | RAD50 | SMARCB1 | XRCC1 |
| BCL9 | CIC | FANCA | GNA11 | LCK | MTOR | PHOX2B | RAD51B | SMARCE1 | XRCC2 |
| BCOR | CREBBP | FANCB | GNA13 | LYN | MUSK | PIK3CA | RAD51C | SMO | YES1 |

Availability of certain assays and features may vary by location. Check website for complete details.

To order or learn more, visit www.CarisLifeSciences.com.
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Where Molecular Science Meets Artificial Intelligence.
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