Comprehensive Genomic Sequencing to Identify Somatic Alterations in Solid Tumors

Introducing **PanCancerlQ[™]**, our next-generation sequencing (NGS) service that enables comprehensive genomic profiling of tumor samples.

The Eurofins Viracor new **PanCancerIQ**TM service uses the Illumina TruSight Oncology 500 (TSO500) assay system, combined with clinical interpretation through utilization of a knowledgebase developed by MD Anderson Cancer Center via a partnership with Philips.



PanCancerIQ supports identification of the four main classes of alterations known to drive cancer growth: mutations, insertions and deletions (indels), copy number variations (CNV), and gene fusions.

In addition, the assay accurately measures key current immuno-oncology biomarkers: microsatellite instability (MSI) and tumor mutational burden (TMB).

The assay also identifies oncogenic driver events that predict response or resistance to treatments, enabling the rapid confirmation and validation clinically relevant mutations, helping clients accelerate their clinical development.

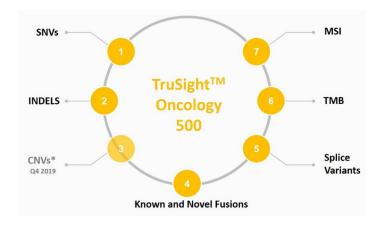
Now you can interrogate the oncogenome with unparalleled breadth and sensitivity, using our integrated genomic solutions along with many other complementary cell-based assay services to accelerate your oncology research and drug development programs.

Complete Genomic Sequencing Solution with Comprehensive Analytics

- Expertise in interrogating FFPE samples
- Optimal sensitivity and reproducibility
- Accurate and sensitive detection of diseaserelated, and therapy relevant genomic alterations
- Bioinformatics expertise for custom analysis and clinical interpretation
- CAP/CLIA validated assay

Genomic Profiling for Clinical Development of Precision Medicine

TSO500 is a hybrid capture-based approach targeting 523 oncology-relevant genes, including biomarkers that are both FDA-approved and in the NCCN guidelines.



The assay leverages unique molecular indices to enable accurate, ultra-low frequency detection of SNVs and Indels. This comprehensive cancer panel interrogates relevant cancer biomarkers with as little as 40 ng of DNA and 20 ng RNA, demonstrating robust analytical performance. In addition to variant calls, the assay reports a tumor mutational burden (TMB) score and microsatellite instability (MSI) status.

PanCancerIQ[™] Specifications Description and Sample Requirements: FFPE (DNA and RNA) 523 genes for DNA variants 55 genes for RNA variants Variants called: SNVs InDels **CNVs** Fusion **Splice Variants** IO signatures (TMB, MSI) Panel size: 1.94 Mb DNA; 358 kb RNA Sample input: 40 ng DNA 20 ng RNA FFPE Sections: 5 um, min 20% cellularity, <20% necrosis **FFPE** slides 10-20 slides

Our expertise in targeted FFPE sequencing, combined with an industry leading NGS platform and high value bioinformatics analysis, provides the ability to rapidly confirm and validate study relevant mutations to accelerate clinical research on novel anti-cancer therapeutics.

PanCancerIQ [™] Performance				
Whole panel CAP/CLIA validated for FFPE				
Sensitivity:				
VAF limit of detection (LOD): Analytical Sensitivity (LOD):	As low as 1% for >= 1000X coverage			
	Minimum DNA for TMB, MSI, CNV: 20 ng			
	Minimum DNA for SNVs, Indels: 40 ng			
	Minimum RNA: 20 ng			
Analytical Specificity:				
Accuracy:	PPA >=98%; PPV >=95%			
	TMB: 85% - 99% (sample dependent)			
	MSI: 96% - 99%			
	CNV: PPA >=90%			
	RNA: PPA >=90%			
Precision:	Small variants: PPA > 95%			
	MSI, TMB: %CV < 30% or St Dev < 4			
	Structural variants: PPA > 95%			

Driven by science, the innovative, consultative approach of our BioPharma team enables clients to overcome obstacles and get faster results. Our new PanCancerIQ assay can help:

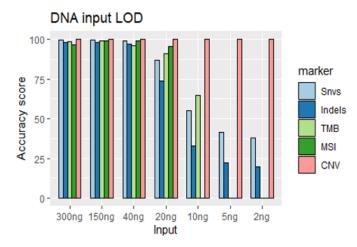
- Screen patients for enrollment
- Stratify patient cohorts,
- Provide biomarker-driven trial optimization
- Optimize clinical trial design and development decisions
- Assess the efficacy of targeted anticancer therapies
- Accelerate the implementation of precision oncology and guide the better use of targeted drugs

Genomic Profiling for Clinical Development of Precision Medicine

Eurofins Viracor BioPharma offers:

- Rapid and accurate genetic analysis of clinical FFPE tissue
- Genomic services to support translational research, and clinical development of precision medicine
- Bioinformatics expertise for analysis, interpretation, and biomarker assay development
- Extensive global testing network
- Improved limits of detection and coverage depth

High Sensitivity and Accuracy of DNA Variants

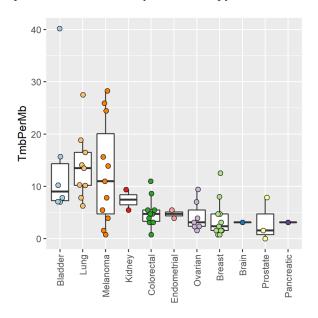


Lower Limit of Detection: 1% VAF at $\geq\!1000\text{X}$ coverage and 40 ng DNA

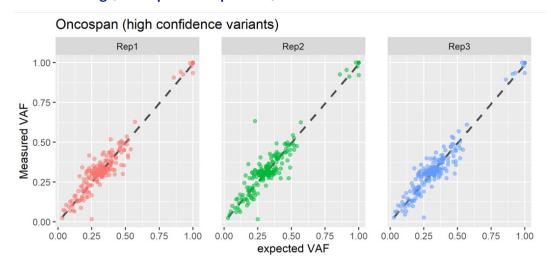
Custom Bioinformatics Services

- Clinical interpretation
- Time to event analysis
- Predictive modeling and classification
- Data linkage to clinical endpoints
- Mutational signature analysis
- Pathway analysis
- Open source and proprietary algorithm development

Analysis of TMB in multiple cancer types:



Reproducible Variable Calling (3 independent operators):

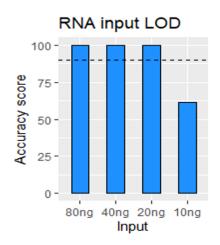


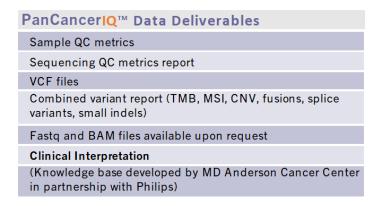
OncoSpan is a well-characterized, cell line-derived Reference Standard containing 386 variants across 152 key cancer genes.

Sensitive and Accurate RNA Fusion Detection:

SeraCare RNA Fusion mix V4				
Fusion	80ng	40ng	20ng	10ng
CCDC6-RET	✓	✓	✓	×
CD74-ROS1	✓	1	✓	×
EGFR variant III	✓	1	✓	×
EGFR-SEPT14	✓	1	1	×
EML4-ALK	1	1	1	×
ETV6-NTRK3	✓	1	1	✓.
FGFR3-BAIAP2L1	✓	1	1	✓
FGFR3-TACC3	✓	✓	1	✓
KIF5B-RET	✓	1	1	✓
LMNA-NTRK1	✓	1	1	1
MET Exon 14 Skipping	✓	1	1	×
NCOA4-RET	✓	✓	✓	✓
PAX8-PPARG1	✓	1	1	✓
SLC34A2-ROS1	✓	1	1	×
SLC45A3-BRAF	✓	1	1	1
TFG-NTRK1	✓	1	1	1
TMPRSS2-ERG	✓	1	1	✓
TMP-NTRK1	✓	1	1	1

Seracare RNA mix V4 Reference (RNA Reference Standard with 18 clinically relevant RNA fusions)





Let our experts support your biomarker-driven clinical trial programs with the comprehensive NGS solution that provides proven utility.

Contact us today to discover how the Viracor Eurofins team can make the difference in your projects.

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