

# MICROSATELLITE INSTABILITY TESTING





# **MSI IN CLINICAL RESEARCH**

## FROM PROMISING CANDIDATE TO CLINICAL PRACTICE

### MSI EVALUATION AS A RESEARCH MARKER FOR IMMUNE-CHECKPOINT THERAPY RESPONSE

In recent years, immune-checkpoint inhibitors (ICIs) have revolutionized the treatment of patients with advanced cancer and ICIs have become a strong pillar in cancer treatment<sup>(1)</sup>. However, understanding the molecular biological background is still required when considering the best indication for ICI<sup>(2)</sup>. In this context several studies have demonstrated that MSI status, as a surrogate for a defective mismatch repair system (dMMR), is a positive predictor for the response to immune-checkpoint inhibitors<sup>(3,4,5)</sup>. Furthermore, since 2017 several IC-therapies have been approved by the US Food and Drug Administration (FDA) or EMA considering the tumor's MSI status.

As the MSI status is a useful surrogate marker, clinical researchers are currently investigating MSI and its implications for predicting the response to immune checkpoint blockade in a variety of tumor entities<sup>(6)</sup>.

### **INVESTIGATION OF MSI IN ENDOMETRIAL CARCINOMA**

The Cancer Genome Atlas Research Network (TCGA) performed an integrating genomic, transcriptomic, and proteomic characterization of endometrial carcinoma. Exome sequence analysis revealed four groups of tumors<sup>(7)</sup>.

- Group 1 carcinomas have somatic inactivating hotspot mutations in the POLE exonuclease domain and a very high mutational burden (ultramutated).
- Group 2 includes endometrioid carcinomas which are microsatellite instable (MSI) (hypermutated), and present frequently with MLH-1 promoter hypermethylation and high mutation rates.
- Group 3 tumors are typically endometrioid carcinomas with low copy number alterations, and a low mutational burden, while lacking POLE mutations and MSI-H.
- Group 4 (Serous-like or copy-number high) show a low mutation rate, nearly universal TP53 mutations, and have a highly unfavorable prognosis.

Clinical researchers are now attempting to bring the TCGA molecular-based classification into clinical practice<sup>(8)</sup>.

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Levine, D., The Cancer Genome Allas Research Network. Integrated genomic characterization of endometrial carcinoma. Nature 497, 67–73 (2013). Concin N, Matias-Guiu X, Vergote I, et al ESGO/ESTRO/ESP guidelines for the management of patients with endometrial carcinoma International Journal of Gynecologic Cancer Published Online First: 18 December 2020.

# INTRODUCING THE MODAPLEX MSI SOLUTION

DETERMINE MSI-H WITH MODAPLEX MSI ANALYSIS KIT







## ANALYSIS USING A STANDARDIZED INDIVIDUAL MSI-H ASSESSMENT

- Simultaneous analysis of proven dinucleotide and mononucleotide markers
- Obtain results through intuitive result analysis

## **REPORT RESULTS WITH CONFIDENCE**

- Verified with artificial material and tested on FFPEderived CRC and EC sample material
- Include forensically accepted (human insertion/ deletion polymorphism) marker as sample mix-up and contamination control
- Comprehensible MSI-H detection using optimal 10ng genomic DNA

## **ENHANCE LABORATORY EFFICIENCY**

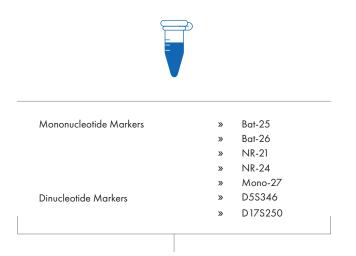
- Setup as straightforward as setting up a PCR
- Modaplex Result Analyzer (Moda-RA) software enables the researcher to interpret markers intuitively

# **MODAPLEX MSI ANALYSIS KIT**

## PROVEN MARKER TO EMPOWER STANDARDIZED MSI-H ASSESSMENT

The combination of five mononucleotide markers (Bat-25, Bat-26, NR-21, NR-24, and Mono-27) and two dinucleotide markers (D5S346 and D17S250) allows the individual MSI status determination. This microsatellite panel has been proven to be highly sensitive and specific in providing results on MSI status across various tumor entities, such as endometrial carcinoma and colorectal cancer<sup>(1,2)</sup>.

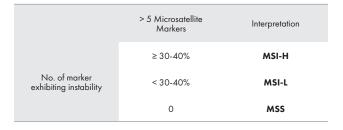
## **CHOOSE YOUR INDIVIDUAL STANDARDIZED MSI-H ASSESSMENT OPTION**







## MSI status assessment using mononucleotide- and dinucleotide-repeat marker<sup>(3)</sup>



### MSI status assessment using five quasimonomorphic mononucleotide-repeat marker<sup>(4)</sup>

	5 Mononucleotide Markers	Interpretation
	≥ 3	MSI-H
No. of marker exhibiting instability	≤2	MSI-L
	0	MSS

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# **EVALUATION OF INSTABILITY**

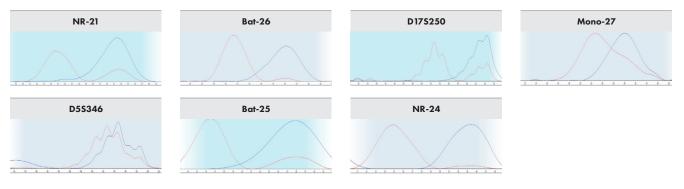
## EXPERIENCE INTUITIVE ASSESSMENT THROUGH AUTOMATIC PEAK-OVERLAY

Recently, certain testing procedures are available to determine the accurate MSI-H status such as PCR-based-, NGS-based- or fully automated approaches. Interestingly, several studies have demonstrated that different methods produce concordant results 95% - 100% when compared to each other<sup>(1,2,3)</sup>.

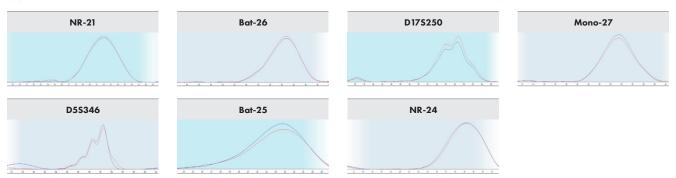
Regardless of the method, instability assessment requires careful estimation and knowledge of the technical limitations. For PCRbased systems, experience means to be knowledgeable on peak alterations and shift events. With Modaplex Result Analyzer Software, users can evaluate these peak alterations and shift events intuitively through an automated overlay of allele-peaks from tumoral and normal tissue.

## VISUAL ASSESSMENT TO EVALUATE THE "INSTABILITY" OF EACH MARKER

Typical results from a MSI-H FFPE colorectal cancer sample, 10ng; cancer tissue (red); normal adjacent tissue (blue)



Typical results from a MSS FFPE colorectal cancer sample, 10ng; cancer tissue (red); normal adjacent tissue (blue)



#### REFERENCES

A. Vanderwalde et al," Microsatellite instability status determined by next-generation sequencing and compared with PD-L1 and tumor burden in 11,348 patients", Cancer Medicine, vol. 7, no. 3, pp. 746-756, 2018

Y.F. Wong et al. "Detection of microscatellities of Microscatellities insbability in endowering in a dynamic and a

## MODAPLEX MSI ANALYSIS KIT FEATURES

## GENERATE RESULTS WITH CONFIDENCE

The Modaplex MSI Analysis Kit enables the users to generate reliable MSI results. For this purpose, the kit is designed and developed according to customer requirements, ensuring a robust, safe, and flexible assay.



### TAILORED TO LABORATORY REQUIREMENTS

To address the limitations of poor quantity and quality of DNA in a formalin-fixed, paraffinembedded environment, the Modaplex MSI assay has been verified on CRC and EC FFPE samples.



### **RELIABLE RESULTS**

The MSI assay is endowed with a comprehensive control concept. It comprises internal controls like a migration size standard as well as external positive and negative controls. In addition, the assay includes a forensically accepted marker serving as the sample-mix up and contamination control.

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## **APPLICABLE FOR LOW SAMPLE AMOUNTS**

The Modaplex MSI assay is suitable for MSI-H detection using small sample inputs. Optimized to be used with a DNA input of 10ng, the assay is suitable for the widespread application on FFPE-derived sample material.

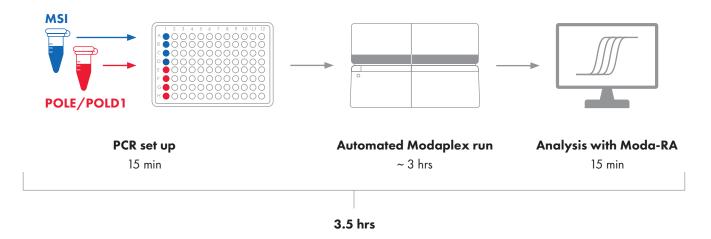
# **MODAPLEX MSI WORKFLOW**

## ENHANCE LABORATORY EFFICIENCY



The Modaplex MSI Analysis Kit is designed for use with the Modaplex instrument, a multiplex PCR bench-top system. It combines qPCR with capillary electrophoresis (CE) in an automated process. Therefore, users can individually combine tests for the purposes of mutational analysis, gene expression, copy number variation, gene fusion, and miRNA, among many others.

### **STREAMLINE LABORATORY OPERATIONS WITH A COMMON PROTOCOL**



Because Modaplex assays will make use of a universal PCR program, you can perform these assays simultaneously through a workflow which is simple as setting up a PCR. Thus, microsatellite fragment analysis and POLE / POLD1 mutation detection can be individually combined and flexibly performed in a single Modaplex run in less than 3.5 hours.

## **ORDER INFORMATION**

Product	Cat. no.	Application
Modaplex MSI Analysis Kit	85-10701-0050*	RUO
Modaplex Instrument	00-04901-0001	

For research use only. Not for use in diagnostic procedures. \*Not available in Italy and the USA

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