

# Plentiplex™ MSI

Made by PentaBase

## READY-TO-USE QUICK GUIDE

Plentiplex™ MSI for Detection of Microsatellite Instability in Cancer patients



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## 1. INTRODUCTION

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This quick guide is meant for experienced users. Please review the full “Instructions for use” carefully before using the kit. The PlentiPlex™ MSI assays offer fragment length analysis of 5 micro-satellite loci, recommended by the Bethesda and Hamelin guidelines, as well as our own PentaBase panel, enabling evaluation of MMR status on a fragment analyzer (Sanger sequencing instrument).

## 2. CONTENT

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PlentiPlex™ MSI assays are supplied as ready-to-use assays in either 1, 12 or 60 reactions in pre-aliquoted 0.2 ml PCR strips or tubes suitable for most standard 96-well PCR instruments. All reagents needed for the PCR amplification of extracted genomic DNA are pre-loaded into the strips or tubes.

## 3. STORAGE AND STABILITY

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The unopened product is stable at -20°C for a minimum of 9 months, but no longer than the expiry date.

**Important:** Keep frozen until use and thaw at room temperature. Avoid repeated freeze/thaw cycles.

## 4. DNA EXTRACTION

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- Use FFPE, fresh frozen or any other suitable biopsy material
- Genomic DNA (gDNA) can be extracted using any valid gDNA extraction kit
- Follow the instructions for gDNA extraction recommended by the kit supplier
- Determine the quantity and quality of gDNA prior to real-time PCR. Do not use gDNA of a low quality
- Use 5-200 ng gDNA per reaction/tube

## 5. KIT PREPARATION

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Thaw one strip or tube per sample per analysis and spin down before removing lids gently

## 6. SETUP

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When using the Bethesda panel, it is mandatory to include a germline (non-tumor) patient gDNA sample for comparison with the corresponding patient tumor sample. Analysis of patient tumor samples with the Hamelin’s and PentaBase panel can be carried out using either a patient-matched or non-matched germline DNA sample as reference.

- Add 5 µL extracted patient DNA (1-40 ng/µL) per tube
- Gently mix with a pipette, seal vial and spin down
- Place strips or tubes in a PCR instrument and run the protocol shown in Table 1

Table 1. PlentiPlex® MSI PCR protocol

Protocol	Temperature	Time	Number of cycles
Hold	94°C	600 sec.	1
Cycling	94°C	50 sec.	10
	55°C	50 sec.	
	72 °C	50 sec.	
Cycling	89°C	30 sec.	25
	55°C	30 sec.	
	72 °C	30 sec.	

Hold	72 °C	10 min.	1
Hold	10 °C	∞	1

After PCR, dilute PCR amplicons 1:10 with DNase free water

- Mix 1 µL of diluted DNA template with 9 µL of formamide and 0.25 µL of size standard DNA ladder in plasticware compatible with DNA fragment/sequence analyzer configurations
- Place in a heating block at 95°C for 120 secs
- Place immediately on ice bath at 4°C for at least 10 min
- Evaluate amplicon size applying standard protocols on DNA sequence analyzer

## 7. NOTES

- All test components should be stored as described in the “Instructions for use” (storage section)
- Do not mix reagents from different lots
- Always spin down before opening the lids

## 8. DATA ANALYSIS

Micro-satellite loci are evaluated individually by comparing the length of amplicons obtained with the reference gDNA to those observed when applying the patient’s tumor-derived gDNA. Both shorter and longer micro-satellite amplicons are indicative of instability. When using the Bethesda panel and the Hamelin and PentaBase panel with patient germline (non-tumor) gDNA as reference, with five micro-satellites analyzed, patients can be categorized as MSS (0 markers exhibiting instability), MSI-L (1 marker being unstable) or MSI-H (2 or more loci being unstable). When using the Hamelin and PentaBase panel with non-matched wild type gDNA as reference, patients can be categorized as MSS (0 markers exhibiting instability) or MSI-H (3 or more loci being unstable). When only 1 or 2 of the 5 loci exhibit micro-satellite length alteration(s) it is recommended to rerun the sample preferably with inclusion of patient-matched germline (non-tumor) gDNA as reference. If a rerun with non-matched wild type gDNA as reference produces the same results, the sample is considered MSS.

Typical size ranges are described in Tables 2-4.

**Table 2: MSI Bethesda Panel loci identification**

Marker	Detected in channel	Label	Typical size range (bp)
BAT25	Blue	PentaGreen	121-127
BAT26	Red	PentAltRed	122-128
D2S123	Blue	PentaGreen	197-227
D5S346	Yellow	PentAltOrange	96-132
D17S250	Green*	PentAltYellow	147-169

\*Risk of crosstalk between green and yellow channel, so that green signal can be observed in yellow channel.

**Table 3: MSI Hamelin Panel loci identification**

Marker	Detected in channel	Label	Average size (bp)
BAT25	Blue	PentaGreen	120
BAT26	Red	PentAltRed	124
NR-21	Green*	PentAltYellow	103
NR-22	Blue	PentaGreen	142
NR-24	Green*	PentAltYellow	132

\*Risk of crosstalk between green and yellow channel, so that green signal can be observed in yellow channel.

**Table 4: MSI PentaBase Panel loci identification**

Marker	Detected in channel	Label	Average size (bp)
BAT25	Blue	PentaGreen	120
BAT26	Red	PentAltRed	124
MONO-27	Red	PentAltRed	148
NR-22	Blue	PentaGreen	142
NR-24	Green*	PentAltYellow	132

\*Risk of crosstalk between green and yellow channel, so that green signal can be observed in yellow channel.

Please refer to the “Instructions for use” for examples of Micro-satellite instability.

## 9. TROUBLESHOOTING

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This short troubleshooting guide may assist in solving most frequent encountered problems that can occur. Please refer to the “Instructions for use” for further troubleshooting.

- If there is low or no signal, try to rerun the PCR and/or re-extract gDNA from the sample
- Extra signals in the yellow channel can be due to cross signal from primers labelled with PentaAltYellow. Please disregard signals in the yellow channel above 140 base pairs.

The full version of the “Instructions for use” can be found at [www.pentabase.com](http://www.pentabase.com).  
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