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# IntelliPlex<sup>TM</sup> KRAS Mutation Plus Kit User Manual

REF

**82022 24 Reactions** 

RUO

For Research Use Only

PlexBio Co., Ltd.

6F-1, No. 351, Yangguang St., Neihu Dist., Taipei City, 11491, Taiwan

TEL: + 886-2-2627-5878 http://www.plexbio.com

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**IMPORTANT:** 

Read the instructions carefully prior to use

#### 1. INTENDED USE

The IntelliPlex KRAS Mutation Plus Kit, based on  $\pi Code^{TM}$  technology and PlexBio's instrument platform, is an in-vitro molecular assay intended for qualitative identification of 27 single nucleotide changes in exons 2, 3 and 4 of the *KRAS* gene using DNA samples derived from formalin-fixed paraffin-embedded (FFPE) tissues. This product is for research use only, and not for use in diagnostic procedures.

### 2. INTRODUCTION

Several cancers have elevated epidermal growth factor receptor (EGFR) activity, and EGFR and the EGFR signaling pathway are targets for the treatments of cancers such as metastatic colorectal cancer (mCRC) and non-small cell lung cancer (NSCLC). KRAS is a GTPase, tethered to cell membranes and functions as an effector molecule of the EGFR signaling cascade. Activating mutations in KRAS result in upregulation of the EGFR-mediated signaling pathway, leading to uncontrolled cellular proliferation.

Cetuximab and panitumumab are EGFR-targeting monoclonal antibodies approved for use in patients with mCRC. However, patients with CRC harboring KRAS mutations are unlikely to benefit from cetuximab or panitumumab therapy. Assessment of KRAS mutation status is therefore crucial for the treatment evaluation of patients with CRC. SelectAmp and  $\pi$ Code technology enables the multiplex, single-well detection of single nucleotide mutations of the KRAS gene from specimens containing large amounts of wild-type genomic DNA with significantly reduced sample requirement compared to conventional methods. The IntelliPlex KRAS Mutation Plus Kit identifies 27 nucleotide changes in exons 2, 3 and 4 of the KRAS gene (Table 1).

**Table 1. Mutations Detected** 

Gene	Exon Codon	Amino Acid Change	Nucleotide Change	COSMIC ID
KRAS		p.G12A	c.35G>C	522
		p.G12D	c.35G>A	521
		p.G12V	c.35G>T	520
	Exon 2 Codon 12	p.G12C	c.34G>T 516	516
	COUOTI 12	p.G12R	c.34G>C	518
		p.G12S	c.34G>A	517

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Gene	Exon Codon	Amino Acid Change	Nucleotide Change	COSMIC ID
		p.G13A	c.38G>C	533
		p.G13D	c.38G>A	532
	Exon 2	p.G13V	c.38G>T	534
	Codon 13	p.G13C	c.37G>T	527
		p.G13R	c.37G>C	529
		p.G13S	c.37G>A	528
	5 0	p.A59T	c.175G>A	546
	Exon 3 Codon 59	p.A59E	c.176C>A	547
	Codon 39	p.A59G	c.176C>G	28518
	Exon 3 Codon 61	p.Q61K	c.181C>A	549
		p.Q61E	c.181C>G	550
		p.Q61P	c.182A>C	551
		p.Q61R	c.182A>G (CAA>CGA)	552
	Codon or	p.Q61L	c.182A>T (CAA>CTA)	553
		p.Q61H	c.183A>C	554
		p.Q61H	c.183A>T	555
	Exon 4	p.K117N	c.351A>C	19940
	Codon 117	p.K117N	c.351A>T	28519
	- 4	p.A146T	c.436G>A	19404
	Exon 4	p.A146P	c.436G>C	19905
	Codon 146	p.A146V	c.437C>T	19900

# 3. TECHNOLOGICAL PRINCIPLES

The IntelliPlex KRAS Mutation Plus Kit utilizes two technologies, SelectAmp and  $\pi$ Code, to achieve high sensitivity multiplex mutation detection.

### SelectAmp Technology

SelectAmp technology enables mutation-specific multiplex PCR amplification by blocking amplification of wild-type sequences with Locked Nucleic Acid (LNA). The subsequent selective PCR amplification of mutated sequences increases assay sensitivity and specificity.

# **πCode MicroDisc**

 $\pi$ Code MicroDisc are manufactured to generate up to 85,000 distinct circular image patterns for multiplexing applications. Each  $\pi$ Code MicroDisc has a distinct circular image pattern, which corresponds to a specific capture agent conjugated to the surface of the disc.  $\pi$ Code tagged with different capture agents are pooled, enabling specific detection of multiple analytes in a one well reaction.

### **Detection Principle**

The test is based on five processes:

- 1. DNA extraction from formalin-fixed paraffin-embedded (FFPE) specimens
- 2. Mutation-specific multiplex PCR amplification
- 3. Hybridization of PCR amplicons with mutation-specific probe tagged  $\pi Code$  in a one-well reaction
- 4. Fluorescent labeling with streptavidin-phycoerythrin
- 5. Image pattern decoding and fluorescent signal detection by the PlexBio<sup>TM</sup> 100 Fluorescent Analyzer

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#### 4. WARNINGS AND PRECAUTIONS

- For research use only. Not for use in diagnostic procedures.
- This assay kit should only be used by qualified laboratory personnel.
- Separate, dedicated rooms and equipment for pre- and post- PCR process with unidirectional manner to avoid any contaminations are required.
- Pre-PCR process preparation should be operated in laminar flow hood to avoid contamination.
- Do not use a kit or reagent past its expiration date.
- Note that tumor samples are non-homogeneous in terms of genotype, and may contain non-tumor sections, which can cause false negative results.
- Reagent components have been diluted optimally. Further dilution of the component reagents is not recommended.
- Specimens should be handled as infectious material. Please follow universal precaution for safe use.
- Store assay kits and reagents according to the product label and instructions.
- Do not mix reagents from different lots.
- Dispose of unused reagents, specimens and waste according to applicable central/federal, state, and local regulations.
- Wear powderless gloves and do not touch and make any markings on the bottom of the plate at any time, as fingerprints and markings would interfere with decoding and signal acquisition.
- General laboratory precautions should be taken:
  - Do not pipette by mouth.
  - Wear protective clothing (e.g., disposable powderless gloves and laboratory coats) and eye protection.
  - Do not eat, drink or smoke in the laboratory.
  - Wash hands thoroughly after handling samples and reagents.
- The workspace, including racks and pipettes, should be thoroughly cleaned and wiped with 0.5% sodium hypochlorite solution followed by wiping with a 70% ethanol solution. A 1:10 dilution of household bleach will produce a 0.5% sodium hypochlorite solution.
- Material Safety Data Sheets (SDS) are available upon request from PlexBio Customer Service.

# 5. QUALITY CONTROL

The IntelliPlex KRAS Mutation Plus Kit contains a series of internal control  $\pi$ Code MicroDiscs that monitor the PCR amplification, SA-PE incubation procedure and background noise. Those controls must meet the specification in each test well with intensities above the cutoffs from the same run, or the run will be considered failed. The external controls (positive control and negative control) monitor the whole testing procedure to prevent false positive and false negative results. The test is considered invalid if any of the controls fail.

# 6. KIT COMPONENTS

The IntelliPlex KRAS Mutation Plus Kit contains sufficient reagents for up to 24 tests. The kit components include:

# (1) KRAS Plus KIT Reaction Mix

Ref. No.: 20188-R

**Quantity & Volume:** 1 vial, 240 μL/vial **Description:** For PCR amplification

Contents: MyFi 5X Reaction Buffer, MyFi DNA polymerase (Microbial), buffered solution containing MgSO₄ and

dNTPs

### (2) KRAS Plus KIT Primer Mix

**Ref. No.:** 20187-R

**Quantity & Volume:** 1 vial, 240 μL/vial **Description:** For PCR amplification

**Contents:** ~4 µM Primer (including biotin-labeled primers)

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# (3) KRAS Plus KIT πCode MicroDisc

Ref. No.: 20191-R

**Quantity & Volume:** 1 vial, 480 μL/vial **Description:** For PCR amplicon capture

Contents: πCode MicroDisc, Glycerol, Phosphate buffered saline, 0.1% Albumin- from bovine (Biological), <0.1%

EDTA and <0.1% Sodium azide

# (4) KRAS Plus KIT POS Control

Ref. No.: 20189-R

**Quantity & Volume:** 1 vial, 120 μL/vial **Description:** Assay positive control

Contents: Plasmid DNA containing KRAS exon 2 G12A sequences (Microbial), Tris-EDTA Buffer

# (5) NEG Control

Ref. No.: 20549-R

**Quantity & Volume:** 1 vial, 120 μL/vial **Description:** Assay negative control **Contents:** Nuclease-free water

# (6) SA-PE Solution

Ref. No.: 20302

Quantity & Volume: 1 bottle, 7 mL/bottle

Description: Streptavidin-phycoerythrin for fluorescent signal acquisition

Contents: Phosphate buffered saline, 0.5% Streptavidin-phycoerythrin, 1% Albumin- from bovine (Biological),

<0.1% Sodium azide

### (7) Hy Buffer

Ref. No.: 20547-R

Quantity & Volume: 1 bottle, 2.4 mL/bottle

**Description:** For hybridization

**Contents:** Saline-Sodium Phosphate-EDTA

**NOTE:** POS Control, NEG Control and Hy Buffer stand for positive control, negative control and hybridization buffer, respectively.

# 7. MATERIALS AND EQUIPMENT REQUIRED BUT NOT SUPPLIED

# Required products for compatibility with IntelliPlex kits:

- 96-well plate (PlexBio; Cat. No. 80025 or Greiner Bio-one; Cat. No. 655101)
- IntelliPlex<sup>™</sup> 1000 πCode Processor (PlexBio; Cat. No. 80033)
- PlexBio 100 Fluorescent Analyzer (PlexBio; Cat. No. 80000)
- U Tray (PlexBio; Cat. No. 80023)
- V Tray (PlexBio; Cat. No. 80024)
- DeXipher<sup>™</sup> RUO (Required: PlexBio; Cat. No. 80050)
- 10X Assay Wash Buffer (PlexBio; Cat. No. 80220)
- Deionized water for dilution of 10X Wash Buffer

# **Required components:**

- Qubit<sup>TM</sup> Fluorometer with dedicated quantitative reagents (Invitrogen; any models) or equivalent
- FFPE DNA extraction kit (Recommended: QIAamp DNA FFPE Tissue Kit, Qiagen; Cat. No. 56404) or equivalent (Stellar FFPE DNA Extraction Kit, PlexBio; Cat. No. 83025-R)
- Clean tubes for PCR reaction (Gunster; Cat. No. MB-P08A or equivalent)
- Dedicated micropipette
- Filter tips for micropipette
- Disposable powderless gloves

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- Vortex mixer
- Micro-centrifuge
- Thermocycler (Recommended: MiniAmp<sup>™</sup> Thermal Cycler, Applied Biosystems<sup>™</sup>; Cat. No. A37834 or equivalent)
- Industrial Computer (Recommended: PlexBio; Cat. No. 80002)

# 8. STORAGE, STABILITY AND TRANSPORTATION

### Storage

All kit components should be stored at 2-8°C.

### Stability

Do not use any kit that has expired. All unopened components are stable up to the expiration date on the label if handled and stored under the recommended conditions.

### **Transportation**

The shipping temperature for the kit is 2-8°C. If the kit package or components are incomplete, please contact PlexBio customer service (service@plexbio.com).

### 9. INSTRUMENT AND SOFTWARE

#### Instrument

Please refer to the instrument user manual for complete operation instructions (Thermocycler, IntelliPlex 1000  $\pi$ Code Processor and PlexBio 100 Fluorescent Analyzer).

#### **Software Installation**

The IntelliPlex KRAS Mutation Plus Kit has a designated Kit App and ENC file. The Kit App contains the  $\pi$ Code target assignments and the ENC file includes the lot number and expiration date. Please make sure you have the Kit App installed and the ENC file imported into DeXipher before your first assay run.

#### Kit App Installation

- 1. Visit www.plexbio.com and download the KRAS Plus Kit App.
- 2. Click on the "Installer" in the APP folder and follow the instructions to complete Kit App installation.

**NOTE**: The Kit App only needs to be installed once. Version updates will be notified by customer service.

# **ENC File Installation**

- Visit <u>www.plexbio.com</u> and download the **KRAS Mutation Plus Kit** ENC file. Each kit lot number will have a unique ENC file, so you will need to download a new ENC file each time you purchase a kit with a different lot number. Make sure to select the ENC file with the lot number that corresponds to your kit.
- 2. Save the ENC file to your computer.
- 3. Follow the PlexBio 100 Fluorescent Analyzer User Manual to import the ENC file.

# 10. SPECIMENS

The IntelliPlex KRAS Mutation Plus Kit has been validated to be used for formalin-fixed paraffin embedded tissues (FFPET). The nucleic acid of all samples must be extracted with a purification procedure supporting downstream PCR amplification.

# **NOTE FOR USE OF FFPET:**

- FFPET specimens may be stored ≤ 30°C for up to 12 months after the date of tissue collection and processing.
   The optimal tissue fixation time for test should be less than 72 hr.
- Only FFPET sections of 10-μm thickness containing at least 10% tumor content are to be used in the KRAS Mutation Test. Any specimen containing less than 10% tumor content should be macro-dissected prior to deparaffinization.
- Do not use stained FFPE specimens which could generate invalid and/or incorrect results.

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### Storage of Extracted DNA

Extracted DNA can be stored at 2°C to 8°C for immediate use (≤ 24 hours), or at -20°C for long-term (> 24 hours) storage. Do not subject the extracted DNA to repeated freeze/thaw cycles.

#### 11. BEFORE YOU START

- 1. Check that the Kit App has been installed and the lot specific ENC file has been imported to DeXipher.
- 2. Check that you have 20  $\mu$ L of extracted DNA ( $\geq$  0.5 ng/ $\mu$ L) ready for analysis.

# 12. ASSAY PROCEDURE

# Warning:

Read the instructions carefully and follow every step of the assay protocol correctly.

# 12.1 DNA Quantification

- 1. Quantify the extracted DNA using a Qubit Fluorometer with dedicated quantitative reagents (or equivalent) according to the manufacturer's protocol.
- 2. The DNA stock concentration should be  $\geq$  0.5 ng/ $\mu$ L to ensure optimal assay performance. Each PCR reaction uses 20  $\mu$ L of a 0.5 ng/ $\mu$ L DNA working stock (10 ng DNA input). Please prepare working stock for all samples before preparing PCR. DNA input amounts lower or higher than 10 ng per reaction are not recommended.

### 12.2 Multiplex PCR Amplification

- 1. Vortex mix each sample before use.
- 2. Spin down and keep samples on ice.
- 3. Prepare the PCR Reaction:

### For each PCR reaction:

KRAS Plus Reaction Mix	10 μL
KRAS Plus Primer Mix	10 μL
Sample/POS Control/NEG Control	20 μL
Total volume	40 μL

# NOTE:

- The amount of Reaction Mix and Primer Mix required for a Master Mix depends on the number of reactions. Always prepare a surplus.
- Both POS Control and NEG Control are required for test validity and report generation and must be included in each assay run.
- 4. Mix by tapping the tubes and spin down before placing the tubes on the thermocycler. Set up the PCR program conditions as below:

### **PCR Program Conditions**

Temp. (°C)	Time	Cycles
95	5 min	-
95	20 sec	
70	20 sec	36
60	60 sec	
4	Hold	-

**NOTE:** Ramp rate:  $3^{\circ}$ C/sec (ABI MiniAmp<sup>TM</sup> Cat. No. A37834).

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# 12.3 DNA Hybridization and SA-PE Reaction

1. **Prepare 1X Wash Buffer:** Transfer 100mL of the 10X Assay Wash Buffer (PlexBio; Ref: 80220) to the IntelliPlex 1000 πCode Processor 1L Wash Buffer bottle and add 900 ml deionized water. Mix by swirling.

**NOTE:** The prepared 1X Wash Buffer can be used for up to one week. Please always check the Wash Buffer is sufficient for assay runs. Additional 10X Assay Wash Buffer can be ordered from PlexBio (Ref. No: 80220).

### IntelliPlex 1000 πCode Processor Wash Buffer consumption:

Procedure	Wash Buffer Consumption (mL)
Self-test	50 mL
DNA & RNA program	150 mL
(1 lane, up to 8 tests)	
DNA & RNA program	220 mL
(3 lanes, up to 24 tests)	

2. Add 20  $\mu$ L  $\pi$ Code MicroDisc to 96 well plate: Mix by vortexing the KRAS Plus  $\pi$ Code for 10 seconds, then, by pipetting, add 20  $\mu$ L of the  $\pi$ Code to each well directly. Vortex the tube of  $\pi$ Code every four wells in between dispensing to ensure homogeneous suspension.

**NOTE**: Each amplified PCR products (including samples, POS and NEG control) should be added into wells lane wise, respectively in order of A1, B1...H1 and followed by A2, B2...H2 and so on.

- 3. Add 100 µL of Hy Buffer to each well.
- 4. Spin down the PCR products.
- 5. **Denature the PCR products** on the thermocycler by heating at **95°C for 7 minutes**, immediately cooled on ice/cooler or thermocycler to ensure the denatured status. Spin down before use. Use immediately (within 1 hour after denaturation).

**NOTE:** Pay attention to the lid temperature of thermocycler while taking out the denatured PCR products.

- Add 20 μL of the denatured PCR products to each well.
- 7. **Pipet the desired volume of SA-PE solution** into the V Tray in SA-PE tank. Please note that the dead volume of V Tray is **500** μL for up to 6 selected lanes or **800**μL if more than 6 lanes are selected. The minimum usage of SA-PE is **one lane (900** μL).

### **Calculation Example:**

For a 3-lane reaction, the required SA-PE solution volume is at least:

400 μL x 3 lanes + 500 μL(dead volume)= 1.7 mL

It is recommended to add extra solution volume into the V-tray to ensure sufficient dispensing volume.

**NOTE:** List of Required SA-PE Solution by Lane(s).

Number of Processed Lane(s)	Required SA-PE Solution (μL)
1	900
2	1300
3	1700
4	2100
5	2500
6	2900
7	3600
8	4000
9	4400

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Number of Processed Lane(s)	Required SA-PE Solution (μL)
10	4800
11	5200
12	5600

- SA-PE solution should be kept in the dark.
- <u>Do not</u> reuse the leftover SA-PE solution and V Tray tank. Replace a new V Tray with every assay run.
- 8. **Run hybridization and wash:** This assay uses the **DNA/RNA program** in the **Molecular Assay** window of the IntelliPlex 1000  $\pi$ Code Processor. Refer to the IntelliPlex 1000  $\pi$ Code Processor user manual and follow the instructions to run the built-in assay program (Homepage/ Molecular Assay/ Well Selection/ DNA/RNA/ Confirm procedure conditions/ Start Running). The plate will be ready for decoding once the program is finished.

#### NOTE:

- IntelliPlex 1000  $\pi$ Code Processor must be maintained properly and regularly.
- Do not open the door when the instrument is in operation.
- The kit contains sufficient reagents for 6 runs of tests (including POS and NEG controls) for a maximum of 24 tests.

# 12.4 Image Decoding and Fluorescent Detection

1. Follow the PlexBio 100 Fluorescent Analyzer User Manual to set up the read.

#### NOTE:

- PlexBio 100 Fluorescent Analyzer must be calibrated regularly (once per month).
- Check that the correct ENC file has been imported.
- 2. Launch DeXipher to run the Qualitative Assay.
- 3. Mark the wells for sample, positive and negative controls.
- 4. Enter sample information and assay name. Place the plate into the device with the correct orientation as shown on the screen.
- The raw data will be analyzed through the kit ENC to generate the mutation call report.

### NOTE:

A single run can include from 2 to 96 tests (including POS and NEG controls) per 96 well Microwell plate. When
running more than 24 specimens, multiple IntelliPlex KRAS Mutation Plus Kits of the same lot will be required.

# 13. DISCLAIMERS

### **Negative Test Result**

A negative test result means that the targeted mutation was not detected by the kit. Experimental errors or other causes may lead to false negative results. Interpretation of the results should consider these possibilities.

### **Positive Test Result**

A positive test result means that the targeted mutation was detected by the kit. Experimental errors or other causes may lead to false positive results. Interpretation of the results should consider these possibilities.

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#### 14. INTERPRETATION OF RESULTS

**Table 2. Interpretation of Result** 

Test Result	Reported Result	Interpretation	
Mutation Detected	Ex. A59T (Refer to Table 1 for details)	Targeted mutation detected	
Mutation Not Detected	None	Targeted mutation not detected	
Invalid Assay	Invalid	<ol> <li>Possible Causes:</li> <li>PCR Inhibition (presence of inhibitor in the sample)</li> <li>Improper stored reagents</li> <li>Low sample DNA input or quality</li> <li>Low πCode Disc Count (the πCode tube was not vortexed before pipetting)</li> <li>Reagent not added</li> <li>Failed Blank πCode Control</li> <li>Sample quality due to improper fixation process or storage condition</li> </ol>	

#### NOTE:

- All runs and specimen validation were performed by the dedicated KIT APP along with IntelliPlex 1000  $\pi$ Code Processor and PlexBio 100 Fluorescent Analyzer.
- In case of heterogeneity of samples or multiple mutations, only the dominantly detected mutation is reported. "Mutation Detected" indicates that the signal for at least one mutation site is greater than the cutoff value of the corresponding target. When multiple mutations are detected in a sample, only the one that exhibits the highest signal is reported.

# 15. ANALYTICAL PERFORMANCE

## Limit of Blank (LoB)

The limit of blank (LoB) values were determined by two operators performing 12 replicates of wild-type KRAS cell line (K562) across three days and four replicates of 12 wild-type KRAS FFPE specimens across three days on two reagent lots. Duplicates of another 43 wild-type FFPE specimens from different biobank and procurement year were also tested. Only "No Mutation Detected" results were observed in these wild type samples.

The cutoff values of each targeted mutations were then determined by 1.4-folds of the measured maximum analytical signal intensity values, respectively.

# Limit of Detection (LoD)

The limit of detection (LoD) was determined using a dilution series (ranging from 0.05-5%) containing different levels of mutant DNA (either from cell lines or mutant plasmid) blended in a background of wild type cell line (K562) DNA. Each dilution was tested with 21 replicates across three days per reagent lot across five operators and two reagent lots. The LoDs were determined based on a positive hit rate at 95% in PriProbit analysis (Table 3). The LoDs ranged from 0.36~3.07%.

Table 3. Limit of Detection (LoD)

Amino Acid Change	Nucleotide Change	LoD (% Mutation)
p.G12A	c.35G>C	0.82
p.G12D	c.35G>A	1.0
p.G12V	c.35G>T	1.5

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Amino Acid Change	Nucleotide Change	LoD (% Mutation)
p.G12C	c.34G>T	1.83
p.G12R	c.34G>C	0.49
p.G12S	c.34G>A	0.73
p.G13A	c.38G>C	1.29
p.G13D	c.38G>A	1.14
p.G13V	c.38G>T	1.58
p.G13C	c.37G>T	0.92
p.G13R	c.37G>C	1.71
p.G13S	c.37G>A	1.0
p.A59G	c.176C>G	1.56
p.A59T	c.175G>A	1.63
p.A59E	c.176C>A	1.4
p.Q61H	c.183A>C	1.02
p.Q61H	c.183A>T	1.44
p.Q61K	c.181C>A	1.1
p.Q61P	c.182A>C	0.48
p.Q61E	c.181C>G	0.64
p.Q61R	c.182A>G (CAA>CGA)	3.07
p.Q61L	c.182A>T (CAA>CTA)	2.01
p.K117N	c.351A>T	0.61
p.K117N	c.351A>C	0.57
p.A146T	c.436G>A	1.5
p.A146P	c.436G>C	0.36
p.A146V	c.437C>T	1.0

# **16. TROUBLESHOOTING**

The troubleshooting listed below addresses possible problem causes and solutions provided during assay procedures.

Problem	Possible Cause	Recommendations
	1. No plate inserted.	1. Confirm plate is inserted and repeat reading.
A1 37 1: 1 A	2. Plate inserted in wrong orientation.	2. Confirm orientation of plate and repeat reading.
No Valid Assay	3. No assay APP installed.	3. Install assay APP and repeat reading.
Assigned	4. No ENC file imported.	4. Import ENC file and repeat reading.
	5. Two or more lots of reagent used.	5. One reagent lot used at a time.

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Problem	Possible Cause	Recommendations
	<ol> <li>No POS Control or NEG Control added.</li> <li>DNase contamination.</li> </ol>	<ol> <li>Ensure POS Control and NEG Control are added.</li> <li>Ensure all operating procedures are followed correctly. Ensure work environment is free of DNase.</li> </ol>
Positive Control Fail / Negative Control Fail	3. Assay did not work.	Make sure all the assay procedures are followed correctly.
J J	<ul><li>4. Cross contamination between samples.</li><li>5. Wrong PC/NC wells chose.</li></ul>	<ul> <li>4. Clean all surfaces and equipment.</li> <li>Operate pre-PCR and post-PCR in the dedicated area and separate the equipment for use.</li> <li>5. Choose the correct PC/NC wells and repeat</li> </ul>
	DeXipher is unable to detect sufficient πCode Mic	reading. roDiscs numbers for mutation determination.
πCode MicroDiscs Count Fail	<ol> <li>πCode MicroDiscs are not proper dispersed in the well.</li> <li>Not enough πCode MicroDiscs added to well.</li> <li>Microbes exist in buffers.</li> </ol>	<ol> <li>Re-disperse the microplate using IntelliPlex 1000 Processor, and repeat reading.</li> <li>Ensure πCode MicroDiscs are well-mixed with proper amount added.</li> <li>Use freshly prepared wash buffer and deionized water for hybridization to reduce πCode MicroDiscs loss rate.</li> </ol>
	4. Instrument error or malfunction.	4. Contact PlexBio Customer Service.
	Performance of SA-PE is assessed by the SAPE Mo	nitor Control.
SA-PE Monitor Control Fail	<ol> <li>No SA-PE was added or insufficient SA-PE solution for dispensing.</li> <li>SA-PE solution inactivation.</li> </ol>	<ol> <li>Make sure all the assay procedures are followed correctly. Calculate sufficient SA-PE solution volume for dispensing. Repeat test.</li> <li>Ensure correct storage condition and minimize the light exposure. Do not use SA-PE past its</li> </ol>
	Incorrect tested lanes of microplate selected for SA-PE solution dispensing.	expiration date.  3. Repeat assay and make sure lanes selected correctly.
	"Background" is determined by measuring MFI of	an internal control that should not give a signal.
Blank Control Fail	<ol> <li>Wrong hybridization conditions.</li> <li>Residues of SA-PE solution in wells after hybridization.</li> <li>PlexBio 100 Fluorescent Analyzer is not calibrated.</li> </ol>	<ol> <li>Check correct hybridization program is selected.</li> <li>Ensure all buffers (Wash buffer and deionized water) on IntelliPlex 1000 Processor are freshmade and sufficient for washing procedures.</li> <li>Perform calibration on PlexBio 100 Fluorescent Analyzer.</li> </ol>
	4. Markings on plates.	4. Do not make any marking on plate.
	1. PCR inhibition exists.	Follow instructions of sample extraction carefully. Ensure required temperature ranges and centrifugation needs are complied. Ensure complete removal of ethanol.
Internal Control Fail	<ul><li>2. PCR procedures are not performed correctly.</li><li>3. DNase contamination</li></ul>	<ol> <li>Make sure all PCR procedures are followed correctly. Do not to use expired materials or mixed lots of reagents. Ensure storage conditions are correct.</li> <li>Ensure all the operating procedures are</li> </ol>
	4. Hybridization did not work.	followed correctly. Ensure work environment is free of DNase.  4. Make sure all the assay procedures are followed correctly. Ensure samples are freshly heatdenatured.

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Problem	Possible Cause	Recommendations		
Reference Gene Fail	Reference Gene monitors quality of tested sample and must be positive.			
	No Sample added or absence of human- derived DNA.	Ensure human-derived DNA samples are added.     Do not use artificial DNA as samples which may generate invalid results.		
	Insufficient sample input for assays or poor sample quality.	2. Quantify samples and check the sample input and quality. If still remains failed, ensure the collected samples meet specimen requirements. Retest with new samples if needed.		
	3. PCR inhibition exists.	3. Follow sample extraction instructions carefully. Ensure required temperature ranges and centrifugation needs are complied. Ensure complete removal of ethanol.		
	4. PCR procedures are not performed correctly.	4. Make sure all PCR procedures are followed correctly. Do not to use expired materials or mixed lots of reagents. Ensure storage conditions are correct.		

### 17. SYMBOLS

Symbol	Explanation	Symbol	Explanation
RUO	For research use only	REF	Catalog number
LOT	Batch number	(i	Consult instructions for use
***	Manufacturer	Ω	Use by Date
ł	Temperature limitation	س	Date of Manufacture
Σ	Contains sufficient for <n> tests</n>		

### **Notice to User**

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