

Gentier Real-Time PCR System

User Manual

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Suitable Instrument Model: Gentier 48E, Gentier 48S, Gentier 48R

Intended Use

The Gentier real-time PCR system (hereinafter referred to as **Gentier instrument**) is intended for performing rapid, accurate polymerase chain reaction (PCR), meanwhile real-time measuring nucleic acid signals from DNA-binding fluorescent dyes or labeled probes and converts them to comparative quantitative readouts of DNA or reverse transcribed RNA.










Special Declaration

Before installing and operating the Gentier instrument, please read this manual carefully, observe the warnings and non-recommended functions. Also bear mind the potential scope for misuse; it is advisable to draw attention to the possible consequences.





User Requirements

The Gentier instrument must only be used by laboratory professionals trained in laboratory techniques, who have carefully read this manual.

Safty Labels on Transport Package

Label	Description
	Fragile: the items inside are fragile, please handle with care.
	This Side Up: indicates the upward side of the transport package.
	Keep Dry: keep the transport package away from rain or any liquid.
	IVD Instrument: the instrument belongs to In Vitro Diagnostic equipment.
	Temperature Limit: indicates the temperature limits for the storage and transportation of package.
	Humidity Limit: indicates the humidity limits for the storage and transportation of package.
	Atmospheric Pressure Limit: indicates the atmospheric pressure limits for the storage and transportation of package.
	Max 4 Packages: it is prohibited to stack more than 4 layers for the storage and transportation of packages.
	Recycle: indicates the packaging materials are recyclable.

Safty Labels on Gentier Instruement

Label	Description
	Biohazard: indicates that a certain area of the Gentier instrument can be easily contaminated with biological reagents or samples. Reminding user to timely disinfect this area, keep necessary precautions during operation and take corresponding protective measures at the same time.
	High Temp.: indicates that a certain area of the Gentier instrument may produce high temperature. Reminding user to pay attention and be careful for burns.
	Moving Part: indicates that a certain moving part of the Gentier instrument may cause personal injury. Reminding user to proceed with appropriate caution.
	IVD Equipment: The Gentier instrument belongs to In Vitro Diagnostic equipment.



European Community Representative: indicates the authorized representative of this instrument in European.



CE Mark: indicates this instrument is in conformity with the essential health and safety requirements set out in European directives.



Serial Number: indicates the serial number of this instrument.



Manufacturer: indicates the manufacture of this instrument.



Caution: indicates the "caution" of this instrument .



Manufacture Date: indicates the manufacture date of this instrument.






Consult Instructions for Use: indicates the consult instructions for the use of this instrument.



Separate Collection for this Equipment: indicates that this instrument is reusable and not contaminated at the end of the instrument life.

Symbols Used in User Manual

Symbol	Description
	Caution: reminding users to pay attention to a certain operation. Operating the Gentier instrument in any manner unspecified in this manual may result in instrument damage or abnormal function.
	Reminding: providing important information regarding the Gentier instrument operation and successful application, including the information explained in further detail elsewhere in this manual.
	Prohibit: prohibiting users from a certain dangerous operation. Otherwise it may result in instrument damage or abnormal function, even constitute a personal injury hazard.


Conventions Used in User Manual


Convention	Meaning
Ordered list	Procedure steps must be performed following the list order.
(Double) Click A	Click A on the application software.
Click A > B	Click B in menu A on the application software.
Press A	Press A key on the instrument system software.
<i>Italic + Bold</i>	Refers to the instructions/options of the application software.
< Italic + Bold >	Refers to the keys/icons of the application software.
Bold	Refers to the instructions/options of the instrument system software.
< Bold >	Refers to the keys/icons of the instrument system software.
[]	Refers to the keys on computer keyboard.
<i>italic + Blue</i>	Indicate the reference chapter.

Safety and Regulatory Compliance


The operation, maintenance and repair of Gentier instrument shall strictly follow the safety specifications listed in this section and through this manual. The design of Gentier instrument has fully considered its biological contamination protection, electrical safety protection and mechanical motion protection. Non-observance of the instructions or performing any operations not stated herein may affect the safety protection provided, and may also destroy the safety standards of design and manufacture as well as the expected application scope of the Gentier real time PCR system.

XATL Co., Ltd. will not be responsible for any possible consequence caused by either not read or violate the instructions mentioned herein.


 **Caution:** please carefully read this manual before operating the Gentier instrument. Incorrect understanding or operations may result in instrument damage or inefficiency usage, laboratory damage, even personal injury.

 **Reminding:** please pay attention to the descriptions with 'Caution', 'Reminding', 'Prohibit' symbols, and the safety labels on the instrument and transport package.


I. General Instrument Safeties and Precautions


 **Caution:** no person except the **XATL Co., LTD** professional engineers are allowed to open the instrument, to replace any component or to debug the Gentier instrument.




 **Caution:** do not drop or damage the Gentier instrument, please handle it with care.

 **Caution:** in case of any following conditions, immediately cut off the power supply and contact the distributor or manufacturer to ask for professional engineer for processing.




- Any liquid has entered into the Gentier instrument;
- Abnormal sound or smell appears while the Gentier instrument is running;
- Gentier instrument is soaked with water or rain;
- Obvious functional changes of the Gentier instrument.

 **Prohibit:** never handling or move the Gentier instrument while it is running.







 **Caution:** for protection against overheating hazards, the openings on the instrument are designed for ventilation. Please do not block these openings nor cover the instrument with dust cover and other materials while it is running.






-  **Caution:** Gentier instrument installation and transportation should be performed by professional engineer or under professional guidelines.
-  **Caution:** do not open the top lid while the instrument is running, this may break the biological safety and electromagnetic radiation protection measures of the instrument.
-  **Caution:** do not force to place unmatched consumables into the sample block.

II. Personal Safeties and Precautions






-  **Caution:** Gentier instrument is heavy, please adopt appropriate tools and methods, or cooperate with other people to complete the lifting or moving of the instrument. Move or lift the instrument in an improper way may result in bodily injury, pain, or instrument damage.
-  **Prohibit:** never touch the plug, the power cord or the power switch with wet hands.
-  **High Temp.:** do not directly touch the sample block and the hot lid while the instrument is still running, they may generate enough heat to cause serious burns. Please wait the sample block to return to idle temperature before opening the top lid.

III. Electrical Safeties and Precautions

-  **Prohibit:** the voltage of Gentier instrument can cause harms to human body, please cut off the power supply before opening the instrument shell, and it is prohibited to replace any part of the instrument while it is electrified.
-  **Caution:** Gentier instrument should be properly grounded, any damage of the internal or external grounding path may cause dangerous.
-  **Caution:** in case of electric leakage, immediately unplug the Gentier instrument and stop using.
-  **Caution:** please unplug the power cord before moving the Gentier instrument.
-  **Reminding:** under normal circumstances, please use the instrument attached power cord. If the original power cord is destroyed, please substitute it with an equal one.
-  **Caution:** the power grid environment of Gentier instrument must have ground wire.

-  **Caution:** for protection against electric shock hazards, the Gentier instrument must be grounded properly. The power cord provided is a standard three-pin plug, please plug it into an appropriate three-wire grounded receptacle for operation safety.
-  **Caution:** please check the power connection carefully. Hold the power plug when you plug the power cord and make sure the power plug is perfectly inserted into the socket, do not pull the power cord to pull out the plug.
-  **Caution:** please keep the power cord away from heater or other high temperature objects. Please do not put anything on the power cord and keep it away from places where people move around.
-  **Caution:** the fuse tube type of the Gentier instrument is F10AH250V, located in the fuse tube box near the power outlet at the rear of the instrument. Use improper fuse tube may lead to circuit system damage. Please check and make sure that the fuse tube has been properly installed before switch on the instrument.
-  **Caution:** before replacing the fuse tube, please cut off the instrument power supply, unplug the power plug and use a screwdriver to pry open the fuse box. Then substitute the old fuse tube with an equal one.

IV. Environmental Safeties and Precautions

-  **Caution:** Gentier instrument is for indoor use only, the room should be well ventilated and without corrosive gas.
-  **Prohibit:** never run the Gentier instrument in places that have or may have flammable and explosive gas.
-  **Prohibit:** spraying liquid on electrical parts may cause a short circuit and result in fire, do not use sprays in vicinity of the Gentier instrument.
-  **Reminding:** the working environment temperature of Gentier instrument should be between 10°C~30°C, and the relative humidity should be between 20%~85%.
-  **Reminding:** the working environment of the Gentier instrument should be under normal atmosphere (the altitude should below 2000m).

V. Biological Safety and Precautions



Biohazard: the sample objects of the Gentier instrument are nucleic acids, please consider all samples are with potential biohazard, and please take applicable safety protection measures and wear appropriate protective goggles, clothing and gloves while processing the samples.



Biohazard: in case of any liquid overflows during the operation, please immediately disinfect the contaminated area with appropriate detergent to avoid the spreading of contaminant, prevent the laboratory or instrument from biohazards.



Biohazard: please comply with the local or national applicable regulations to complete the disposal of waste samples and contaminated materials.



Biohazard: user should consider the abandoned Gentier instrument as biological contaminated material and comply with the local or national applicable regulations to complete the instrument disposal. Before disposal or recycle the instrument, please completely clean and disinfect the instrument.



Biohazard: user should only use reagents and consumables that are within their expiration date.

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A. Overview

1. Application Fields

The Gentier real-time PCR system is intended for performing rapid, accurate polymerase chain reaction (PCR), meanwhile real-time measuring the nucleic acid signals from DNA-binding fluorescent dyes or labeled probes and converts them to comparative quantitative readouts of DNA or reverse transcribed RNA.

The Gentier real-time PCR system can be used in medical institution laboratories and clinical laboratories for infectious pathogens (such as viruses, bacteria, mycoplasma, chlamydia, etc.) detection; or multiple tumor marker tests of neoplastic diseases, etc. For scientific research, it could be used for the fluorescence quantitative or qualitative analysis of genetic materials DNA/RNA in fields of immunology, molecular biology, forensic science, genetics, archeology, zoology, phytology, etc.

2. Product Constitute and Instrument Structure

2.1 Product Constitute

The Gentier real-time PCR system is mainly consisted of control system, power system, temperature control system, detection system, man-machine interface system and instrument shell, etc.

Among them, the temperature control system includes Peltier heating module, sample block and hot lid, etc; the detection system includes LED excitation light source, photodiodes and light filter, etc; the man-machine interface system includes data acquisition and analysis software, which is mainly responsible for real-time data collection, fluorescence diagram formation, data processing and diagram analysis, in order to quantify or characterize the target nucleic acid and obtain other test report information, etc.

2.2 Instrument Structure

The structure of Gentier instrument is shown in figure A-1.



Figure A-1. Gentier Instrument Structure

1&11. Top lid	2&10. Top lid handle	3. Radiator air vent
4. Power air vent	5. Power socket	6. Fuse tube box
7. Power switch	8&12.USB interface	9. Internet access
13.Hot lid	14.Sample block	15. Touch screen

3. Product Introduction

3.1 Instrument Parameters and Characters

3.1.1 Instrument General Parameters

► **Instrument Specification:**

Dimension: 400mm(L) x 260mm (W) x 260mm(H);

Weight: 11kg;

► **Package Specification:**

Dimension: 500mm(L) × 350mm (W) × 360mm (H);

Weight: 13kg;

► **Power Specification:**

Voltage: AC 100-240V;

Frequency: 50-60Hz;

Rated power: 600VA;

► **Communication Specification:**

Network port: TCP/IP protocol;

Ethernet connection;

► **Application Environment:**

Temperature: 10°C ~30°C;

Relative humidity: 20%~85%, non-condensing;

Atmospheric pressure: 85.0kPa~106.0 kPa;

Altitude: below 2000m;

► **Storage and Transportation Enviroment:**

Temperature: -20°C ~55°C, with transport package;

Relative humidity: less than 93%;

► **Running Noise:**

While the instrument is running, the maximum noise does not exceed 65 decibel.

3.1.2 Instrument Technical Parameters

Thermal Parameters

► **Temperature Accuracy:** $\leq 0.1^{\circ}\text{C}$;

► **Temperature Uniformity:** $\pm 0.1^{\circ}\text{C}$;

► **Temperature Precision:** $\leq 0.1^{\circ}\text{C}$;

► **Max Heating and Cooling Ramp:** 8°C/s

Optical Parameters

► **Excitation Light Source:** LED light sources;

► **Detect System:** photodiodes;

Optical Channel:	Dyes	Instrument Models		
		48E	48S	48R
1	FAM, SYBR Green I, etc.	✓	✓	✓
2	VIC, HEX, TET, JOE, etc.	✓	✓	✓
3	ROX, Texas Red, etc.	✓	✓	✗
4	Cy5, etc.	✓	✓	✗

Detective Parameters

► **Throughputs:** simultaneously detect 48 samples;

► **Repeatability:** $CV \leq 0.5\%$;

► **Linear Correlation:** $|r| \geq 0.999$, within the scope of no less than five magnitudes concentration gradients.




Man-machine Interactive System Parameters

► **Touch Screen:** with built-in 7.0' inches full color touch screen, providing more simple operation. The instrument can run alone without the control computer. The stand-alone running saves more space and energy.


► **Main Control Computer:** the recommend configurations of main control computer are listed as below, customs are allowed to purchase by themselves.

- CPU: 2.4GHz, dicaryon;
- Internal Storage: $\geq 2G$;
- Hard Disk: 500G hard disk is recommended;
- Operating System: applicable for Windows XP, Windows 7, Windows 8, etc.

► **Network Control:** one computer can be connected with 10 Gentier instruments at the same time, user can remote control the experiment and truly realize the network centralized control.

-  **Caution:** the main control computer of the Gentier instrument is not designed for online use, connect it to internet may cause risks of computer virus infection or hacker attacks. **XATL Co., LTD.** will not be responsible for any damages caused by.
-  **Caution:** it is not recommended to install other software on the main control computer of the Gentier instrument. Otherwise there may be potential risks of software module conflict, and may also influences the results reliability.
-  **Caution:** **XATL Co., LTD.** will not provide anti-virus software. Therefore, if necessary, please take preventive measures to prevent the main control computer from virus.

3.1.3 Instrument Characters



- ▶ **Touch Screen Operation:** Gentier instrument possesses a built-in 7.0'-inches colored touch screen, very easy for operation. It can run independently from the computer;
- ▶ **Independent Temperature Control:** Gentier instrument adopts independent temperature control technology and high repeatability temperature zone. Truly realize the optimization of PCR;
- ▶ **Power-off Protection:** Gentier instrument possesses instantaneous power-off protection function which can protect all configuration settings in case of power off and the interrupted experiment can continue after power on.
- ▶ **Multiple PCR Step Modes:** Gentier instrument provides multiple PCR step modes, such as touchdown step, long step, gradient step, melting step and so on.
-  **Reminding:** the Gentier 48S instrument model cannot support gradient PCR step mode.
- ▶ **Remote Running:** user could edit experiment settings on control computer and command the connected Gentier instrument, truly realize the remote running control and real-time monitoring.
- ▶ **Lis System:** Gentier instrument can be connected to hospital Lis system.

3.2 Software Characters

- ▶ **Software Interface:** wizard-style interface, intuitive layout and program setting, very easy for operation.

- ▶ **Software Language:** with switchable multiple languages. Default languages are Chinese and English.
- ▶ **Multiple Functions:** Gentier software possesses multiple analysis functions that can adapt to a variety of experimental requirements. Such as absolute quantification analysis, relative quantification analysis, melting curve analysis, high resolution melting (HRM) analysis, genotyping analysis, end point fluorescence analysis, etc.
- ▶ **Data Transmission:** data can be transmitted between the Gentier instrument and the control computer. In the process of experiment, Gentier instrument will real-time transmitting the experiment data to the control computer.
- ▶ **Data Storage:** Gentier instrument can store over 1000 experiment setting /experiment data files.
- ▶ **Program Setting:** each Stage can contain 99 steps, and the maximum cycle number is 99. User can set the program for low temperature long-term preservation after the experiment.
- ▶ **Report Template Design:** the unique universal report function is designed by **XATL Co., LTD.** for user to customize their own report form, more suitable for the personal needs of all user.

3.3 Reagent Specifications

- ▶ **PCR Reagent:** open platform for all qualitative and quantitative PCR reagent.
- ▶ **Dye:** FAM/SYBR Green I, VIC/HEX/TET/JOE, ROX/Texas Red, Cy5, etc.
- ▶  **Reminding:** Gentier 48R instrument model cannot support ROX/Texas Red, Cy5 or similar fluorescence reagent.
- ▶  **Biohazard:** user should only use reagents and consumables that are within their expiration date.

3.4 Consumable Specifications

- ▶ 0.2mL single PCR tube.
- ▶ 0.2 ml 8-strip PCR tube.

B. Gentier Instrument Installation

1. Unpacking Instructions

The Gentier transport package is as shown in figure B-1. The Gentier instrument and its accessories are well preserved in a carton case. In order to prevent the collision and oscillation during transportation, the Gentier instrument is sealed with plastic bag and well supported by protective foams.



Figure B-1. Package Details

1.1 Unpacking Steps

The unpacking of the Gentier transport package mainly includes 4 steps, as shown in figure B-2. The specific operation method is as follow.

- 1st Step:** cut off the packing belt and unseal the transpot package.
- 2nd Step:** take out the instrument accessories and remove the protective foam on top of the instrument..
- 3rd Step:** hold the Gentier instrument on both sides and take it out of the carton.
- 4th Step:** remove the plastic bag and place the instrument on steady plane.



Figure B-2. Unpacking Steps

- ❗ **Reminding:** in order to prevent the formation of condensation water, please do not open the transport package until it reaches the room temperature.
- ❗ **Reminding:** please check the package integrity before open it. In case there is any damage, collision or water mark, please contact the transport department or our company.
- ❗ **Reminding:** please check the instrument and accessories according to the packing list and ensure that all components are present and intact, report any missing items to **XATL Co., LTD.**
- ❗ **Reminding:** please fill in the relevant content on installation information feedback table after acceptance, and send it back to our company for document and warranty.
- ⚠ **Caution:** please reserve the original packing materials for future use. The transport package of Gentier instrument is designed to reduce the instrument damage and ensure its transportation safety. Adopt other packaging materials will break the warranty, and **XATL Co., LTD.** will not be responsible for damages as consequences of improper packaging that incurred during the transportation. Please also keep instrument related documents provided by our company for future use too.

2. Working Environmental Requirements

The Gentier instrument is for indoor use only.


2.1 Instrument Space Requirements


- 1) The Gentier instrument should be placed in the room with low humidity (between 20%~85%RH) and appropriate temperature (between 10°C~30°C). The room should be well ventilated and without corrosive gas.
- 2) The Gentier instrument should be placed on steady lab work bench and leave enough space for placing the attachments, cables and reagent bottles, etc.
- 3) Please keep the Gentier instrument away from heat sources (directly sunshine, heater, stoves) and water sources (such as water pool, water tube etc.).
- 4) The working environment of Gentier instrument should be without electromagnetic interference, vibration and high frequency wave electrical equipment.
- 5) For single Gentier instrument running, the distance between adjacent objects and the surrounding ventilation openings shall be no less than 30cm.
- 6) For protection against overheating hazards, the openings on the Gentier instrument are designed for ventilation. Please do not block or cover these ventilation openings while it is running.
- 7) Do not place the Gentier instrument on soft surface, the instrument base may sink into the soft surface and block up the air inlet beneath.
- 8) Do not place the Gentier instrument close to the wall or pile up other goods at the rear of the instrument, this may affect heat dissipation.


2.2 Instrument Power Requirements


- 1) The power grid environment of Gentier instrument should possess ground wire and the instrument should be properly grounded.
- 2) Please ensure the power strip of Gentier instrument has 3~4 three-phase plugs in order to meet the demands for the instrument, control computer and printer.


- 3) The power specifications of Gentier instrument are listed in *"A-3.1 Instrument Parameters and Characters - 3.1.1 Instrument General Parameters"*, use improper power may damage the circuit system and cause fire, it is recommended to use UPS power supply.

 **Reminding:** before connecting the AC power supply, please ensure the consistent of the Gentier instrument required voltage and the power supply voltage (allowable deviation $\pm 10\%$). And make sure that the rated load of receptacle is no less than the requirement of the Gentier instrument.

 **Reminding:** if the power supply system of the Gentier instrument working environment is unstable, please do not connect other electrical equipment at the same power circuit, and it is required to equip with over 600VA AC regulated power supply.

 **Prohibit:** spraying liquid on electrical parts may cause a short circuit and result in fire, do not use sprays in vicinity of the instrument.

 **Caution:** do not put anything on the power cord and keep it away from places where people move around. Hold the power plug when you plug the power cord and make sure the power plug is perfectly inserted into the socket, do not pull the power cord to pull out the plug.

 **Reminding:** under normal circumstances, please use the instrument attached power cord. If the original power cord is destroyed, please substitute it with an equal one.


3. Instrument Installation

3.1 External Device Connection

Please plug the corresponding power cords of Gentier instrument, control computer and printer to the power supply respectively. Then please confirm the connections of external devices, such as the control computer, display, keypad, mouse and printer, etc. The external device connection is completed.

3.2 Application Software Installation

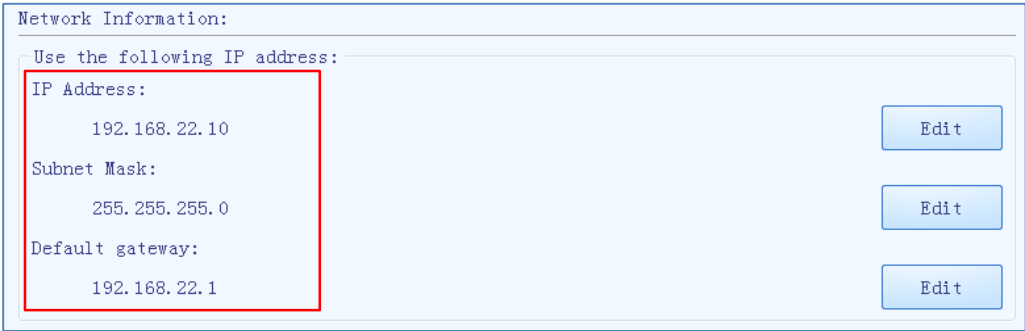
The application system software is required on the control computer for commanding the Gentier instrument and analyzing the experimental data. Before starting up the application software, user should install it on the control computer. The specific operations are as follow:

- 1st step:** switch on the control computer and close its firewall.
- 2nd step:** place the Gentier software CD in computer's CD drive.
- 3rd step:** the application software installation interface will automatically pop up, please follow the prompt to install the software.
- 4th step:** double-click the MED-TL  icon on the desktop to startup the Gentier application software.

3.3 Computer Network Setting

The network address of the control computer should be set according to the default network information of the Gentier instrument. So the control computer can connect the identifiable Gentier instruments within LAN.

- 1st step:** please first check the network information of Gentier instrument, which is displayed on the network information window of the instrument software, as shown in figure B-3a. For detailed operation, please refer to [E-2.4 Gentel Settings – 2.4.2 Configuration Sub-tab – Network Information](#).
- 2nd step:** please open the **Control Panel > Network and Sharing Center > Local Area Connection > Properties > Internet Protocol Version 4 (TCP/IPv4)** on the control computer, as shown in figure B-3b.
- 3rd step:** set the IP address, subnet mask and default gateway of control computer according to the default network information of the Gentier instrument .



Network Information:

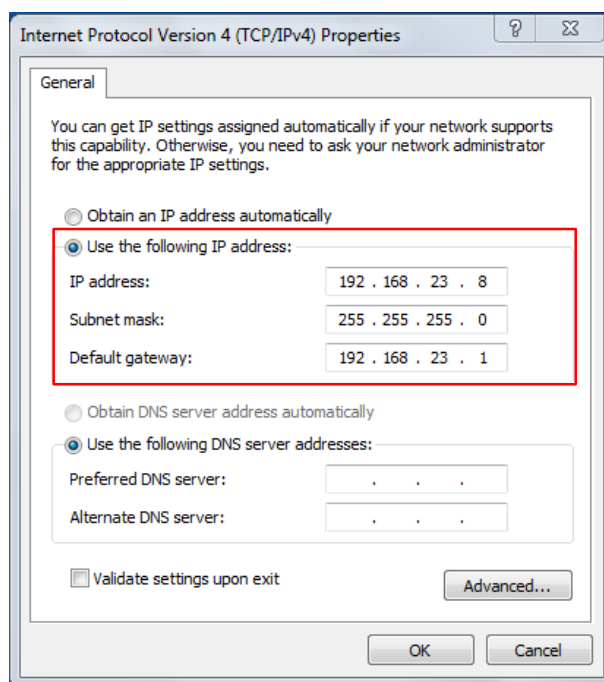
Use the following IP address:

IP Address:	192.168.22.10	Edit
Subnet Mask:	255.255.255.0	Edit
Default gateway:	192.168.22.1	Edit

Figure B-3a. Network information window - default network information

vs

Figure B-3b. Control computer network setting



- **IP address:** the first 3 octets of control computer IP address should be identical to the Gentier instrument IP address.
- **Subnet mask:** the control computer subnet mask should be identical to the Gentier instrument subnet mask.
- **Default gateway:** the control computer default gateway should be identical to the Gentier instrument default gateway.

4th step: click [**OK**] to confirm the control computer network setting.

5th step: after the control computer network setting, user can connect the identifiable Gentier instruments within LAN, for detailed operations, please refer to [D-3.1 Menu Bar - Tool \(T\) submenu- Instrument Management](#).

C. The Preparation before Experiment

1. Instrument Self Inspection

The Gentier instrument possesses self-inspection function, it is recommended to let the instrument conduct the self-inspection before running any experiment, to ensure that it can work normally.


1st Step: switch on the power switch of Gentier instrument.

2nd Step: Gentier instrument will automatically conduct the self-inspection after power on, which will inspect instrument version, whether electric system is in normal working state, whether power supply is normal and query initialize motor positions, etc.

3rd Step: after the self-inspection, the Gentier instrument will enter the standby state.

 **Caution:** before power on the Gentier instrument, please ensure:

- The external devices of the Gentier instrument are properly connected.
- The power supply of Gentier instrument is properly connected.

 **Caution:** in case the Gentier instrument fails to pass the self-inspection, do not slap or shake it, please contact the distributor or **XATL Co., LTD.**

2. Reagent Preparation

1st Step: please follow the operation instructions of PCR kit to prepare the PCR reagent.

2nd Step: please add the sample and PCR reagent into the suitable consumables, if needed, please seal the consumables .

3. Instrument Loading

1st Step: please hold the top lid handle to lift up and open the top lid, the sample block will be present in front of the user.

2nd Step: place the consumables that contain the sample and PCR reagent mixture on the sample block and close the top lid.

3rd Step: select or edit an experiment program and then start running.

D. Application Software Operation

The software operation instructions of the Gentier real-time PCR system mainly including the introduction of basic software functions and operation descriptions, such as user account management, new experiment creation and edition, real-time monitoring and experiment data analysis, etc.

Among all the 48 throughputs Gentier instrument models, Gentier 48E instrument model possesses the most comprehensive functions. Therefore, the Gentier 48E instrument model is taken as an example for the function and operation description in this user manual, the instrument and software interface figures presented herein are all based on Gentier 48E instrument model. For different Gentier instrument models, the software interface will display the corresponding features and instrument model names, this user manual will specify the differences of software interfaces or functions.

1. Startup Application Software


After successfully installing the application software on the control computer, user can double click the  icon on the desktop or click the application software file on the start menu to startup the Gentier real-time PCR application software. The Welcome screen of the application software is as shown in figure D-1.



Figure D-1. Welcome screen of the Gentier real-time PCR application software

After starting the application software, the Welcome screen will automatically switch to the startup interface and pop up quick start bar on this interface, as shown in figure D-2.

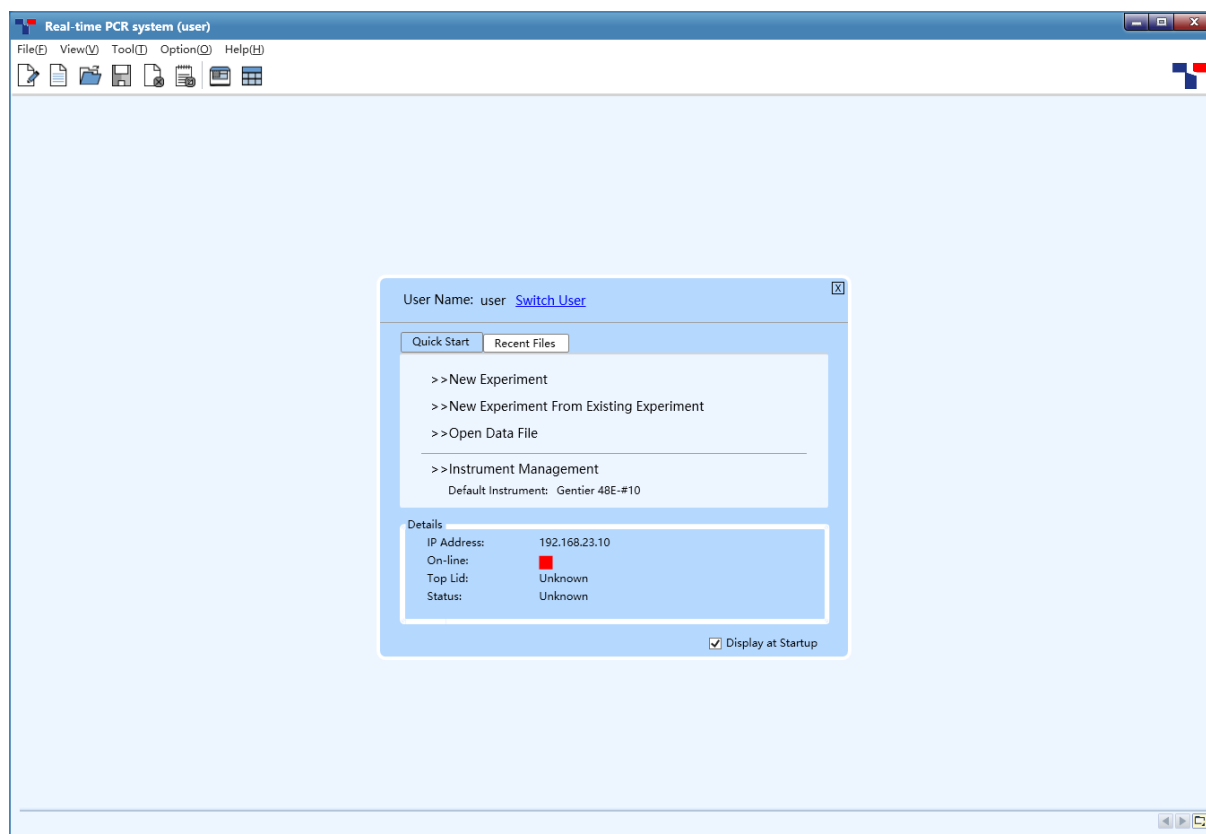


Figure D-2. Startup interface of application software

2. Quick Start Bar

The quick start bar of application software is as shown in figure D-2.1. At this interface, user can login, create new experiments, open data file, set default instrument, view instrument information and conduct other operations.

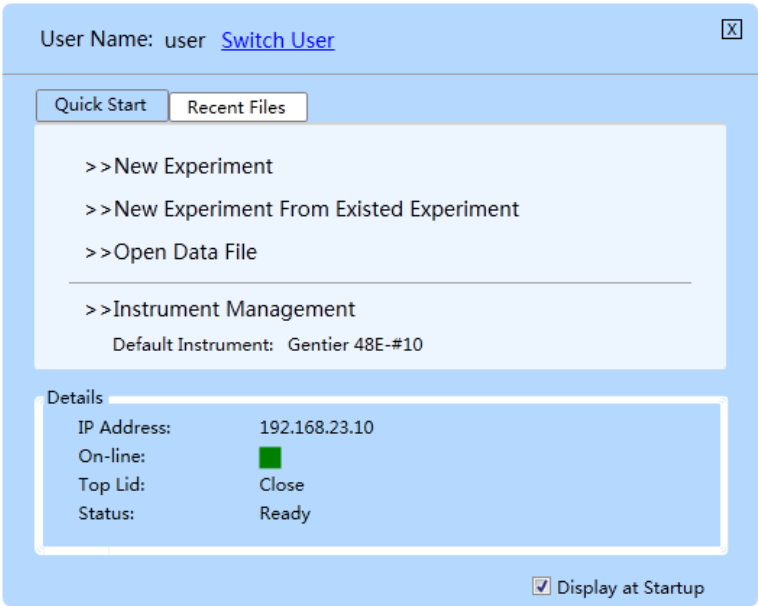


Figure D-2.1 Quick start bar

Quick Start Bar Introduction & Parameter Descriptions

- ▶ The current **User Name** is displayed on the top of quick start bar and user can click < **Switch User** > to change the current user account.
- **Login:** user can click < **Switch User** > and enter the registered account name in the input box; or select a registered account name in the drop-down menu, and then click < **Login** > to log in as the current user account, as shown in figure D-2.1a.
- ❗ **Reminding:** the application software provides two user names by default: **user** and **admin**.



Figure D-2.1a Quick start bar - Login

- **Register New User Account:** user can click < **Switch User** > and select **Add User** option in the drop-down menu of **User Name** input box; then the application software will automatically pop up the add user dialog box, as shown in figure D-2.1b. Please enter a new user account name in the **User Name** input box and click < **OK** > to register a new user account.

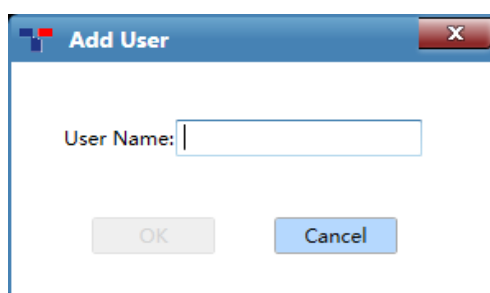


Figure D-2.1b Quick start bar - Add User

- ▶ **Quick Start** tab: includes four shortcut keys: < **New Experiment** >, < **New Experiment from Existed Experiment** >, < **Open Data File** > and < **Instrument Management** >.
- Click < **New Experiment** > and the application software will automatically pop up the new experiment dialog box, as shown in figure D-2.1c. User can manually enter a name in the **Experiment Name** input box and click < **New** > to create a new experiment file.

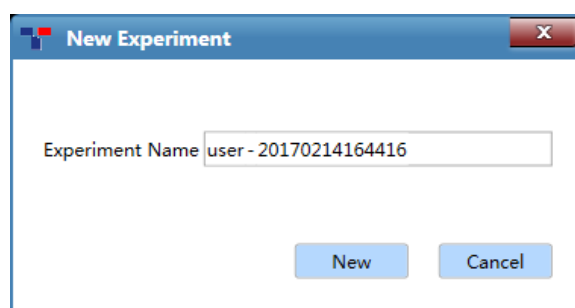




Figure D-2.1c Quick start bar - New Experiment

- ❗ **Reminding:** the application software defaults to name the current new experiment with the logged in user account name and experiment creation time.
- ❗ **Reminding:** the experiment name can consist of numbers, letters, horizontal lines, underlines, or Chinese characters, but cannot contain special characters.
- Click < **New Experiment from Existed Experiment** > and the application software will automatically pop up the choose experiment file window; user can choose an pre-existed experiment file and click < **Open** >; then the application software will create a new experiment file with same experiment settings as the selected experiment file.
- Click < **Open Data File** > and the application software will automatically pop up the open experiment file window; user can select an experiment data file from the save path and click < **Open** > to view and analysis the experiment data file.

- Click < **Instrument Management** > and the application software will automatically pop up the instrument management interface; user can manage all instruments within the local area network (LAN). For operation details, please refer to [D-3.1 Menu Bar - Tool \(T\) submenu-Instrument Management](#).
 - ▶ **Recent Files** tab: display the recent experiment or data files and user can directly click the file name to open the file.
 - ▶ **Details**: the details of the default instrument are displayed at the bottom part of quick start bar, including **IP Address**, **On-line**, **Top Lid** and **Status** informations; when user open the application software again after setting the default instrument, the application software will automatically connect the default instrument and display its status.
 - ▶ User can check the **Display at Startup** check box to decide whether to open the quick start bar automatically when starting the application software.
 - ▶ User can click the  icon on the upper left corner of quick start bar to close the quick start bar. To open the quick start bar again, please click **View > Quick Start [Ctrl + G]** in the menu bar of application software's main interface or click <  **Quick Start** > icon in the toolbar to open the quick start bar.
-

3. Main Interface

After closing the quick start bar, the application software will automatically enter the main interface, which consists of menu bar, tool bar and operational area, as shown in figure D-3.



Figure D-3. Main Interface

3.1 Menu Bar

The menu bar of application software includes: **File (F)**, **View (V)**, **Tool (T)**, **Option (O)** and **Help (H)** five submenus.

File (F) submenu: includes nine options.

Options and Functional Description

1. **New Experiment (N)**: create a new experiment.
2. **New Experiment from Existed Experiment**: user can select a saved experiment file and create a new experiment file with same experiment settings as the selected experiment file.
3. **New Experiment from PANA**: Gentier instrument can be connected with PANA nucleic acid workstation and the Gentier application software can establish experiment according to the PCR reaction system established by the PANA nucleic acid workstation.

4. **Open Data File...:** user can select an experiment data file from the save path and click < **Open** > to view and analysis the experiment data file.
 5. **Recent Files:** user can click the experiment name from the list to open the selected data file.
 6. **Close Experiment:** close an opened or created experiment file.
 7. **Save...:** the experiment file is saved to the default file path.
 8. **Save As...:** user shall specify another file save path and save the experiment file.
 9. **Export Raw Data ...:** user can choose a path to export the experiment raw data file.
 10. **Export All Data Sheets to Excel ...:** user can choose a path to export the experiment data as an Excel file.
 11. **Report Manager...:** user can edit and manage the experiment report information after the experiment.
 12. **Exit:** exit and close the application software.
-

View (V) submenu: includes three options.

Options and Functional Description

1. **Quick Start:** open the **Quick Start** bar. For detailed functions, refer to [D-2 Quick Start Bar](#).
2. **Show Toolbar:** user can decide whether to **Show Toolbar** on the main interface of application software. For detailed functions, refer to [D-3. Main Interface - 3.2 Tool Bar](#).
3. **Show Instrument Information:** user can decide whether to **Show Instrument Information** on the main interface of application software.
 - If user check the **Show Instrument Information** option in the **View** submenu, the instrument information area will be displayed on the top right corner of main interface to remind user the current status of the connected Gentier instrument, as shown in figure 3.1.


On-line:  Loading Platform: Close Status: Ready

Figure D-3.1 View Submenu - Instrument Information Area

- **On-line:** display the connection status between the application software and the instrument: ■ indicates disconnected status; ■ indicates connected status;
- **Loading Platform:** display the status of Loading Platform, which is *Close* or *Open*;
- **Status:** display the running state of the connected instrument, which is *Ready* or *Running*;

Tool (T) submenu: include three options.

Options and Functional Description

1. **Instrument Management:** Click **Instrument Management** option in **Tool** submenu and the application software will automatically pop up the instrument management interface, which consists of these parts: **Instrument List**, **Details** tab and **File Transmission** tab, as shown in figure D-3.2. User can manage all instruments within the LAN on this interface.

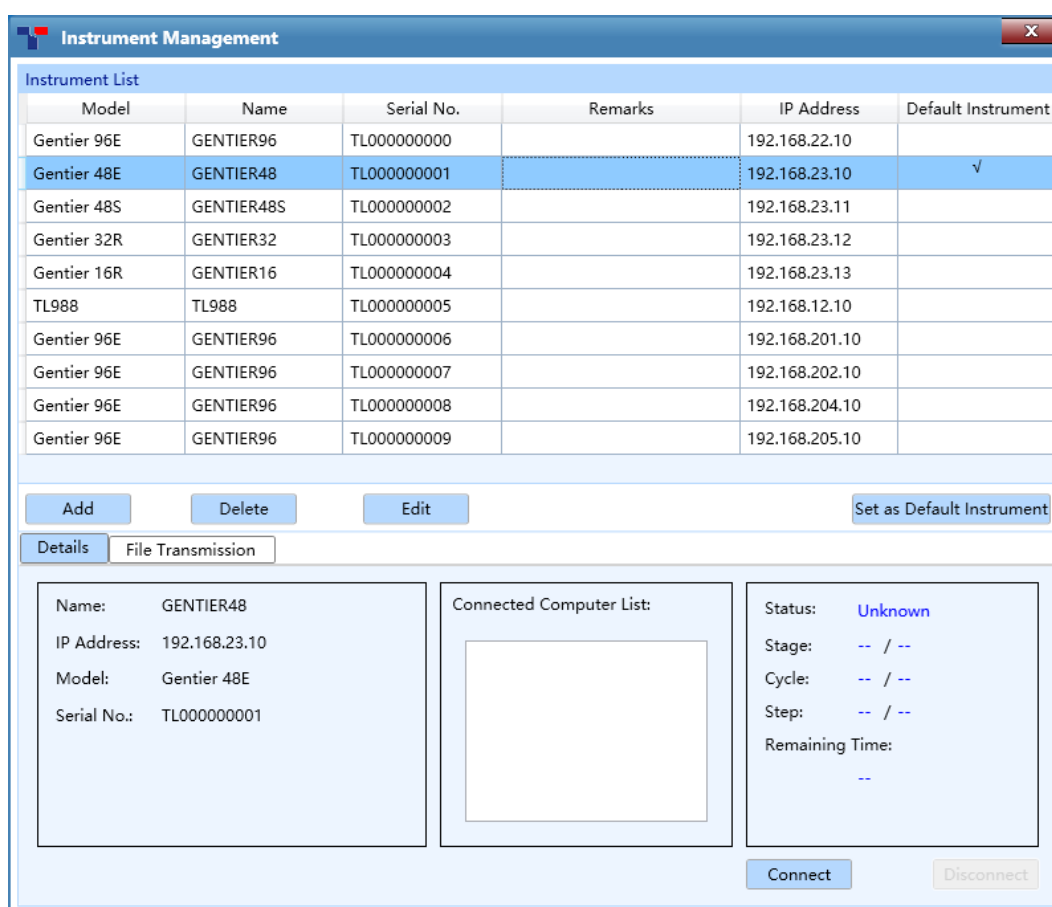


Figure D-3.2 Tool submenu - Instrument Management interface

► **Instrument List:** displays relevant information of instrument connected, including **Model**, **Name**, **Serial No.**, **Remarks**, **IP Address** and whether the instrument is set as the **Default Instrument**. There are four function keys under the list: **< Add >**, **< Edit >**, **< Delete >** and **< Set as Default Instrument >**.

a. Click **< Add >** and the add instrument window will automatically pop up, as shown in figure D-3.2a.

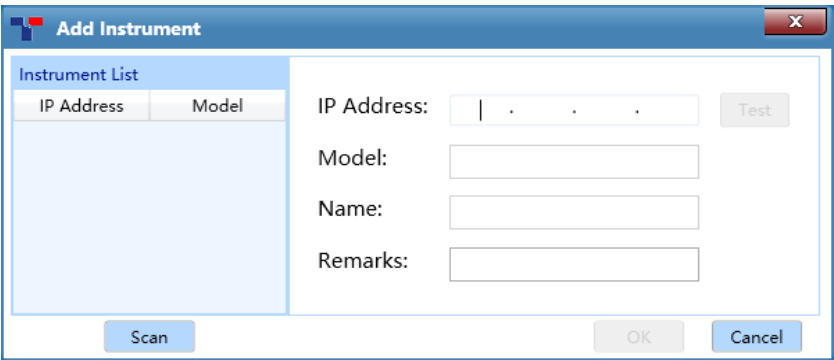


Figure D-3.2a Instrument Management interface - Add Instrument window

- User can click **< Scan >** and the application software will automatically scan all identifiable Gentier instrument within LAN and display their **IP Address** and **Model** in **Instrument List**.
- User can double click a certain instruemnt IP address in the **Instrument List** and the selected IP address will be displayed in the **IP Address** input box on the right; click **< Test >** and the application software will test the selected IP address; if the test passes, the selected instrument can be connected, then click **< OK >** to add the current instrument.
- If user already know the IP address of a certain instrument, you can also manually enter the address in **IP Address** input box and click **< Test >**; if the test passes, the selected instrument can be connected, then click **< OK >** to add the current instrument.

b. Click **< Edit >** and the Edit Instrument window will be automatically popped up; user can change the remarks of instrument in the **Remarks** input box, as shown in figure D-3.2b.

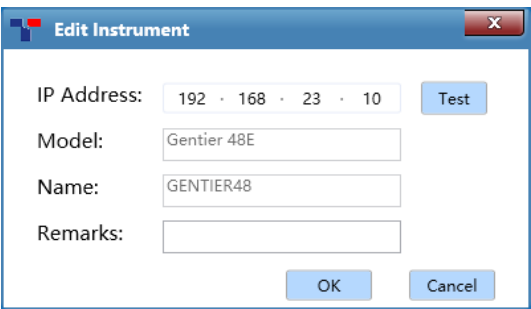


Figure D-3.2b Instrument Management interface - Edit Instrument window

c. Click < **Delete** > to delete the instrument selected in the list.

d. Select any instrument in the **Instrument List** and click < **Set as Default Instrument** > to set it as the default instrument, it will be automatically connected after startup of application software.

► **Details** tab: includes three information areas and two function keys, as shown in figure D-3.2c.

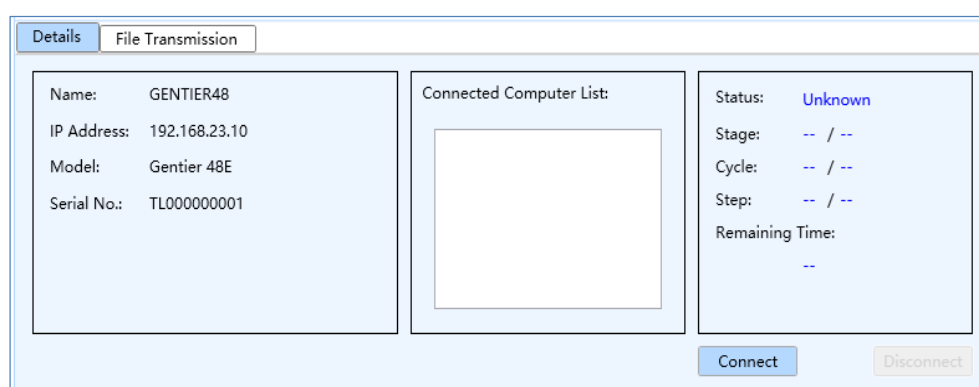


Figure D-3.2c Instrument Management interface - Details tab

a. Information Areas:

- Instrument Information Area: display the relevant information of instrument selected in the **Instrument List**, such as **Name**, **IP Address**, **Model**, **Serial No.** and **Version No.** of instrument components, etc.
- Computer information Area: display the IP address list of the **Connected Computer** within LAN.

❗ **Reminding:** the IP address of the **Connected Computer** should be set according to the default IP address of Gentier instruemnt.

- Running information Area: display the real-time **Status** of connected instrument, such as **Stage**, **Cycle**, **Step** and **Remaining Time**.

b. Function keys:

- Click < **Connect** > to connect the instrument selected in the **Instrument List**.
- Click < **Disconnect** > to disconnect the connected instrument.

- **File Transmission** tab: to transmit the file between the control computer and the connected Gentier instrument, as shown in figure D-3.2d.

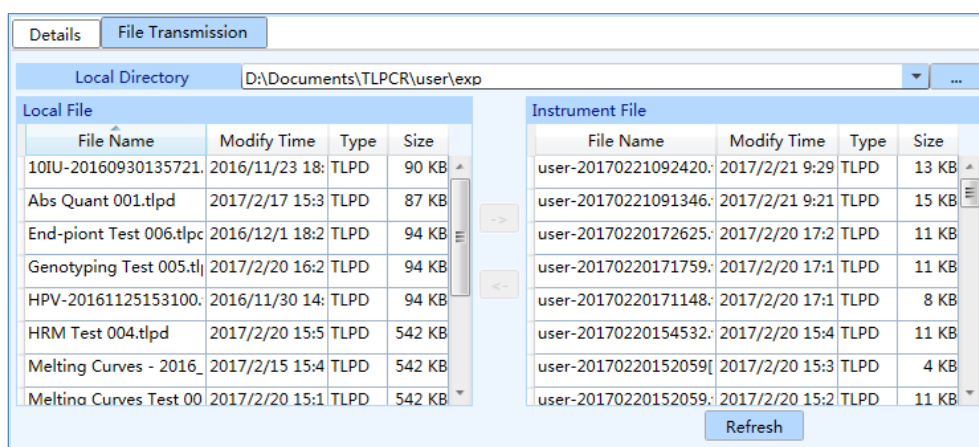


Figure D-3.2d Instrument Management interface - File Transmission tab

- a. **Local Directory**: user can select a local experiment file directory or experiment data file directory from the drop-down list of **Local Directory**.
 - b. **Local File**: after selecting a **Local Directory**, the **Local File** list will display all experiment files and data files in this directory.
 - c. **Instrument File**: display all experiment files and data files saved in the connected instrument.
 - Click < → > to download the selected **Local File** to the connected instrument.
 - Click < ← > to upload the selected **Instrument File** to the control computer and save in the **Local Directory** selected by user.
 - Click < **Refresh** > to refresh the displayed **Instrument File** list.
2. **T_m Calculator**: Click **T_m Calculator** option in **Tool** submenu and the application software will automatically pop up the T_m Calculator interface, as shown in figure D-3.3.

Figure D-3.3 Tool submenu - T_m Calculator interface

Interface Introduction and Functional Description

- ▶ User can click **A T G C** four keys to input the forward primer sequence in **Forward Primer** input box;
- ▶ User can click **A T G C** four keys to input the reverse primer sequence in **Reverse Primer** input box;
- ▶ User can input the salt concentration value in **Salt Concentration** input box by using the keyboard or **▲ ▼** keys;
- ▶ User can click **< Calculate >** and the application software will automatically calculate **Fwd Primer T_m value**, **Rev Primer T_m value**, **Avg T_m value** and **Annealing Temperature**.

3. **Research Report:** Click **Research Report** option in **Tool** submenu and the application software will automatically pop up the research report interface, as shown in figure D-3.4.

! Reminding: user shall create an experiment before entering the research report interface.

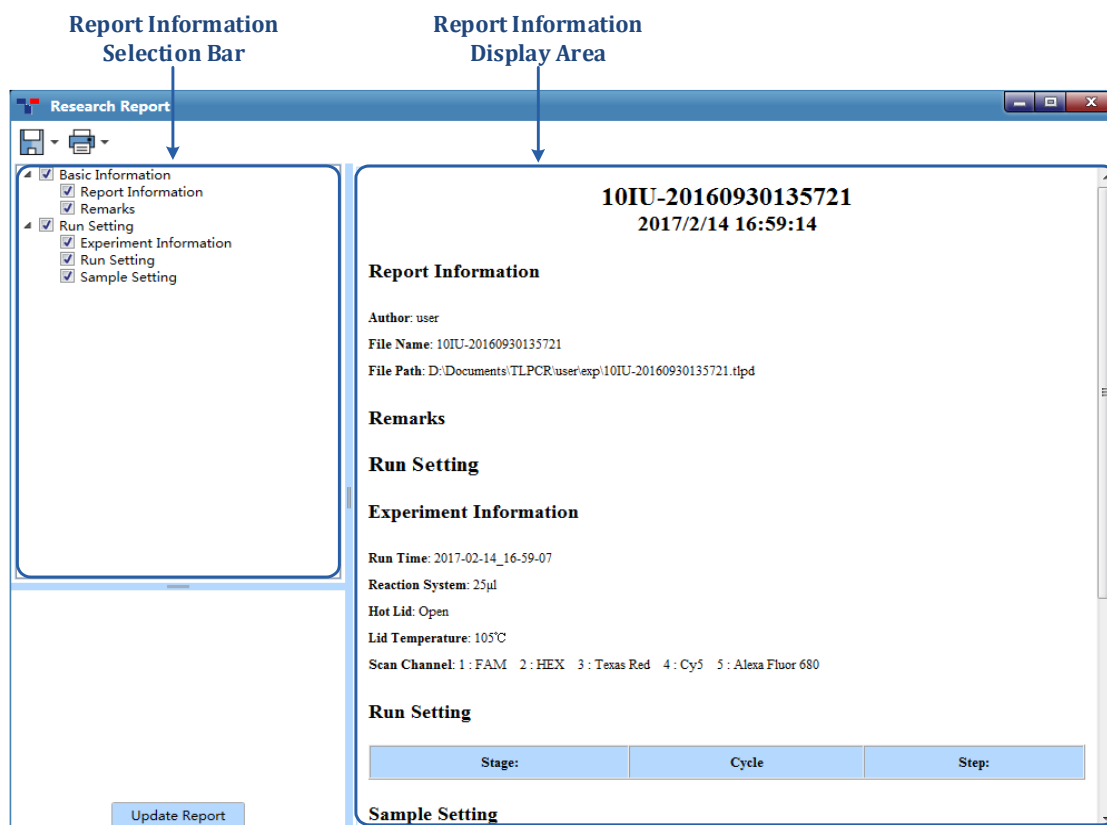




Figure D-3.4 Research Report interface - Research Report

Interface Introduction and Functional Description

- ▶ **Report Information Selection Bar:** user can check the relevant information check boxes in the report information selection bar to determine the information to be displayed and printed in research report.
- ▶ **Report Information Display Area:** display the information selected in the report information selection bar.
- ▶ **Update Report:** user can click < **Update Report** > to refresh the information in the research report.
- ▶ **Save Research Report:** after editing the research report, please click <  **Save** > icon to choose a path for saving the research report and choose a file format from the drop-down list after the icon to save the research report as a PDF or html file.
- ▶ **Print Research Report:** After editing the research report, user could click <  **Print** > icon to choose a printer and select **Print** option from the drop-down list after the icon to print the current research report, or select **Print Preview** option from the drop-down list after the icon to preview the research report.

Option (O) submenu: includes three options.

Options and Functional Description

1. **Configuration Management:** Click **Configuration Management** option in the **Option** submenu and the application software will automatically pop up the configuration management interface, as shown in figure D-3.5a.

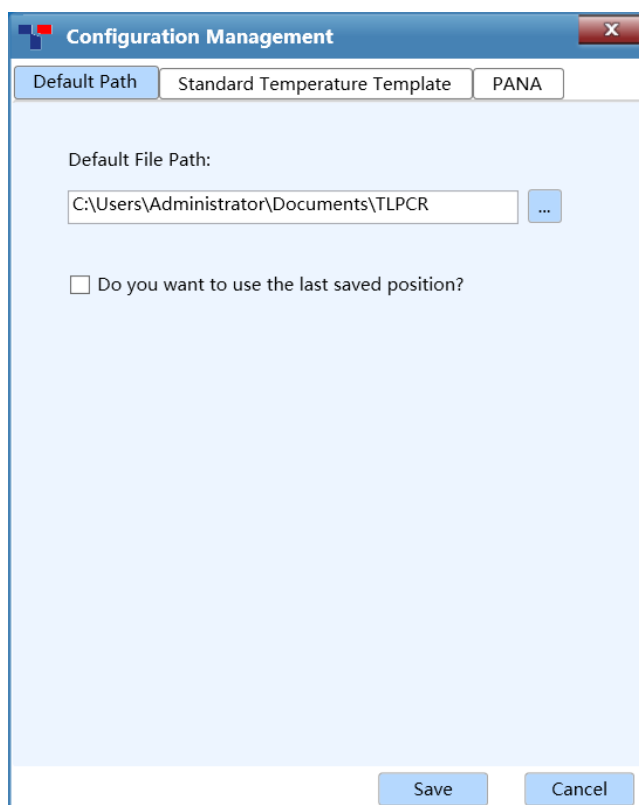



Figure D-3.5a Option submenu - Configuration Management interface - Default Path tab

Interface Introduction and Functional Description

- **Default Path** tab: the configuration management interface displays the **Default Path** tab by default, as shown in figure D-3.5a.
 - User can set the default path for saving run parameter template, sample parameter template and other experiment files in **Default Path** tab. User can manually enter the path name in **Default File Path** input box; or click  icon and select a path from the pop-up Browse Folder.

- User can check ***Do you want to use the last saved position*** check box to determine whether to use the same default file path for saving experiment file or template next time. If the ***Default File Path*** is successfully set and user want to save the experiment file to another path, please click ***File > Save As...*** and select another file save path, then click ***< Save >*** to save the experiment to the selected file directory.
- ***Standard Temperature Template*** tab: 7 predefined standard temperature templates are displayed in this tab, as shown in figure D-3.5b. These templates are provided by the application software for temperature program stage settings.

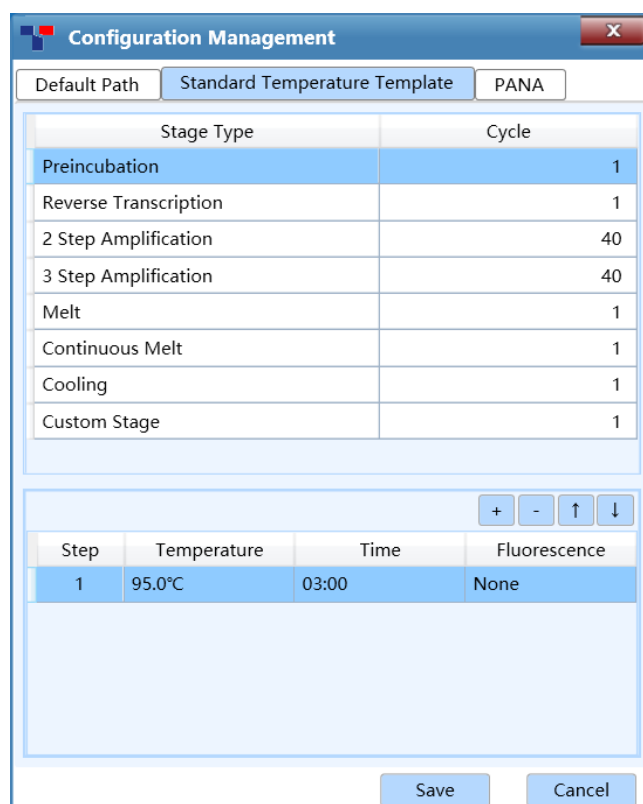


Figure D-3.6 Configuration Management interface - Standard Temperature Template tab

- User can modify the predefined ***Standard Temperature Template*** in this tab, change the corresponding stage and step settings. For detailed information, please refer to ***D-3.3 Operation Area - 3.3.1 Run Setting***.
- ***PANA tab***: Gentier instrument can be connected with PANA nucleic acid workstation of our company, and experiment can be created according to the PCR reaction system established by PANA nucleic acid workstation.

- User can check the PANA connection check box in the PANA label and fill in the IP address box below to ask for the IP address of the PANA full-automatic nucleic acid workstation to be connected. Finally, click test to check whether the PANA instrument can be connected, , as shown in figure D-3.5c.

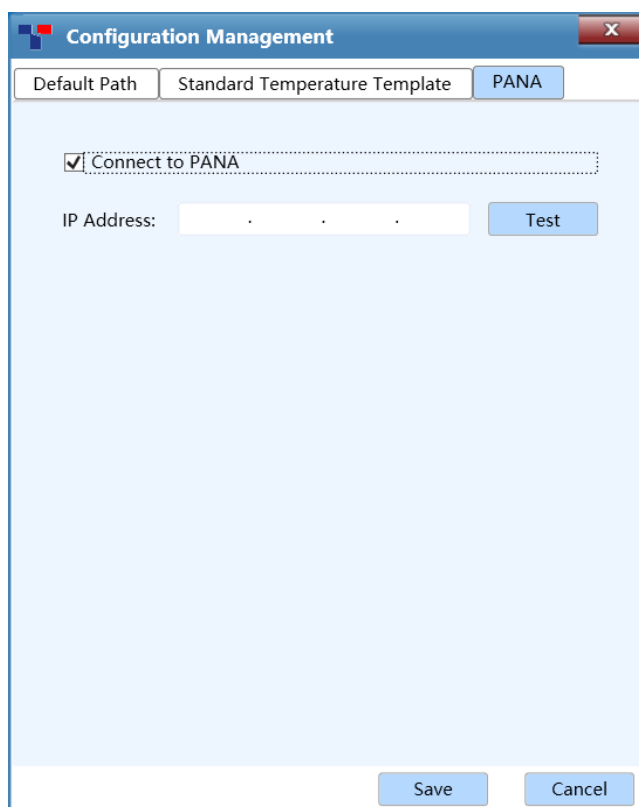


Figure D-3.5c Configuration Management interface - PANA tab

2. **User:** click **User** option in the **Option** submenu and the application software will automatically pop up the user interface, as shown in figure D-3.6a.

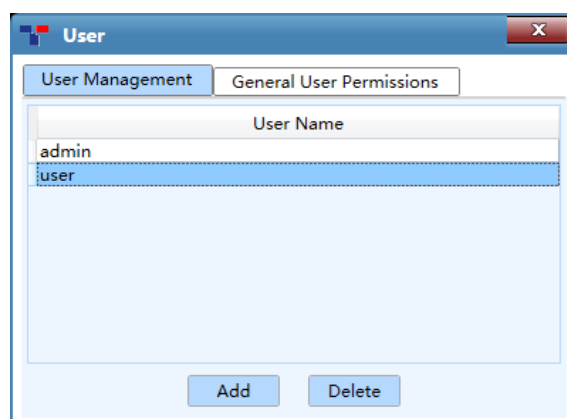


Figure D-3.6a User interface - User Management tab

- ▶ **User Management** tab: if user login as **admin** user account (the password is "admin"), user can manage general user accounts in this tab.
 - < **Add** >: **admin** user can add general user accounts in **User Management** tab.
 - < **Delete** >: **admin** user can delete general user accounts in **User Management** tab.
- ▶ **General User Permissions** tab: as shown in figure D-3.6b.

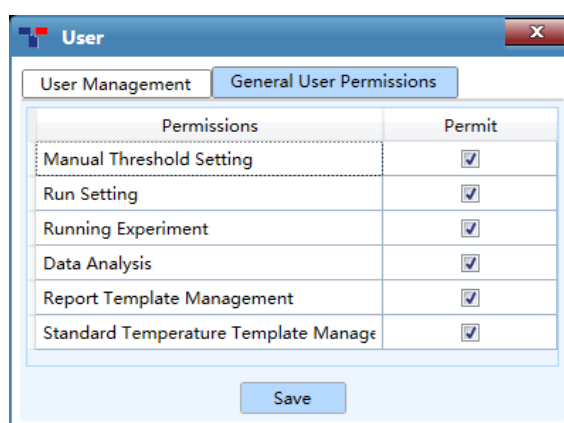


Figure D-3.6b User interface - General User Permissions tab

- a. If user login as **admin** user account, user can manage general user permissions.
 - **Admin** user can set the general user permissions by check relevant **Permissions** check box in the **Permit** column.
 - **Admin** user can click < **Save** > to save the current general user permission settings.
 - b. If user login as general user account, user can view the general user permissions in this tab.
3. **Choose Language**: click **Choose Language** option in the **Option** submenu and the application software provide two optional software languages:
 - **Chinese**: check **Chinese** option and the software language will switch to Chinese after restarting the application software.
 - **English**: check **English** option and the software language will switch to English after restarting the application software.

Help (H) submenu: includes three options.

Options and Functional Description

1. **User Manual**: open the Gentier Real- Time PCR System User Manual.

2. **Home Page:** visit the home page of Drawell.
3. **About:** display the application software version and copyright information, as shown in figure D-3.7.

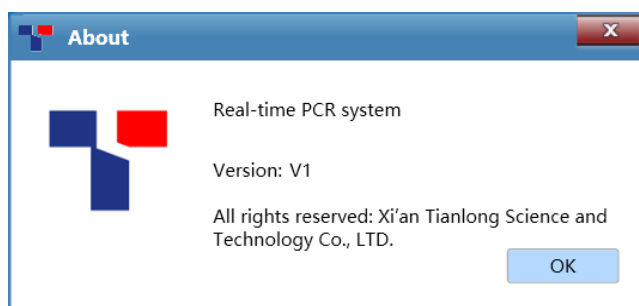










Figure D-3.7 Help submenu - About interface

3.2 Tool Bar

The Tool Bar of the application software consists of eight commonly used function icons.

Options and Functional Description

- ▶ <  **New Experiment** >: create a new empty experiment.
- ▶ <  **New Experiment from Existed Experiment** >: user can select an existed experiment file and create a new experiment file with same experiment setting as the selected experiment file.
- ▶ <  **Open Data File** >: open a data file for viewing or analyzing.
- ▶ <  **Save Experiment** >: the experiment file is saved to the default file path set in configuration management interface.
- ▶ <  **Close Experiment** >: close an opened experiment file.
- ▶ <  **Export** >: after experiment, user can choose a path to export the raw experiment data as an Excel file.
- ▶ <  **Instrument Management** >: click to enter the instrument management interface. For detailed operation, please refer to [D-3.1 Menu Bar - Tool \(T\) submenu- Instrument Management](#).

- ▶ <  **Quick Start** >: click to open the quick start bar. For detailed operation, refer to [D-2 Quick Start Bar](#).

3.3 Operation Area

After creating a new experiment file or open an experiment file, the operation area on the main interface of the application software will be activated, as shown in figure D-4.

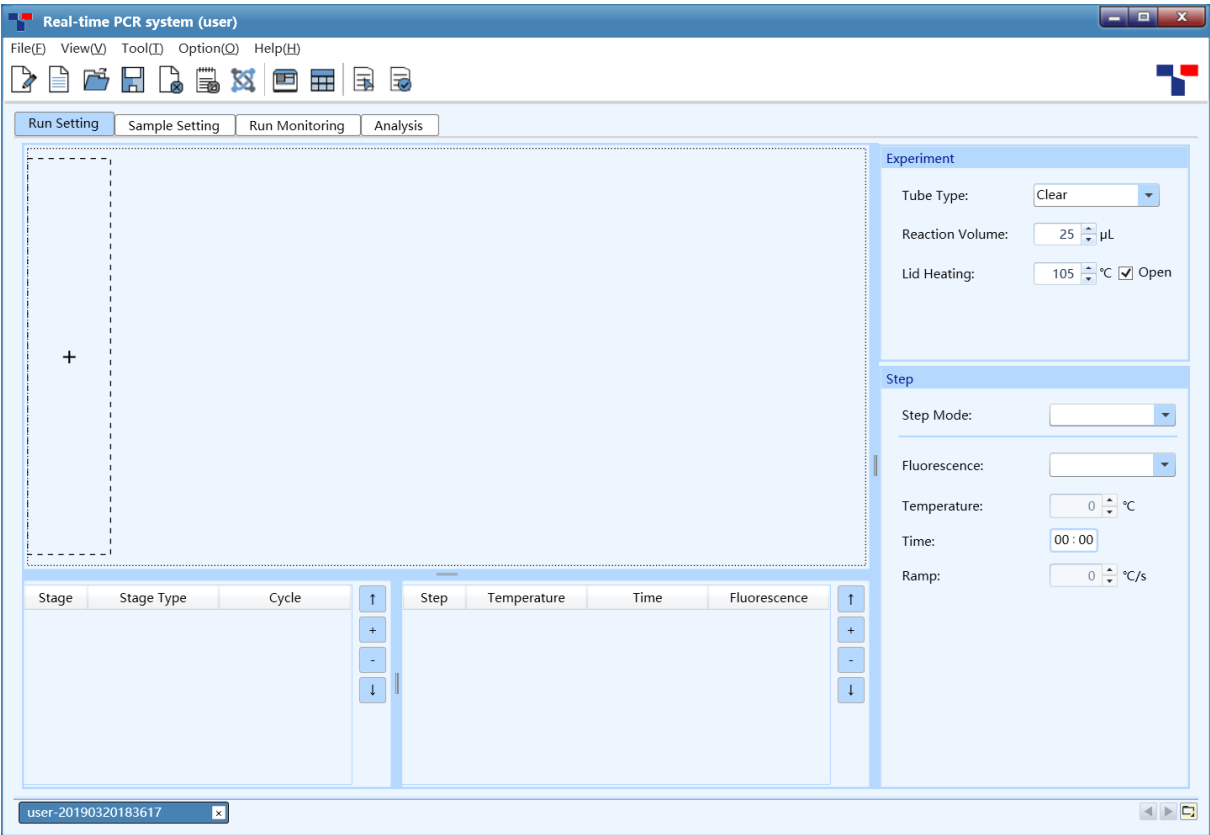









Figure D-4. Operation area on main interface

1. The experiment file created or opened will be displayed in the experiment file tab at the bottom of operation area and user can select or close any experiment file in this tab.
-  **Reminding:** if several experiment files are created or opened at the same time, user can click  and  icon on the right side of the experiment file tab to view all experiment files; or click  icon to view the list of all experiment files.



2. The operation area is consisted of four tabs: **Run Setting**, **Sample Setting**, **Run Monitoring** and **Analysis**. User can select different tabs for relevant operations to realize complete experiment workflow.

 **Reminding:** when user select the different tab in the operation area, different function icons will be added accordingly in the tool bar of application software.




- **Run Setting** tab: when the **Run Setting** tab is selected on the main interface, two additional function icons will be shown in the tool bar on the run setting interface.


- <  **Choose Run Parameter Template** >: click this icon in the tool bar and the application software will automatically pop up choose run parameter template window; user can choose a saved run parameter template set the current experiment based on the selected template.
- <  **Save Run Parameter Template** >: click this icon in the tool bar and the application software will automatically pop up save run parameter template window; user can choose a path and save the parameters of current experiment as a template.

- **Sample Setting** tab: when the **Sample Setting** tab is selected, two additional function icons will be shown in the tool bar on the sample setting interface.

- <  **Choose Sample Parameter Template** >: click this icon in the tool bar and the application software will automatically pop up choose sample parameter template window; user can choose a saved sample parameter template and set current experiment based on this template.
- <  **Save Sample Parameter Template** >: click this icon in tool bar and the application software will automatically pop up save sample parameter template window; user can choose a path and save the sample parameters of current experiment as a template.

- **Analysis** tab: when the **Analysis** tab is selected, four additional function icons will be shown in the tool bar on the analysis interface.

- <  **New Analysis** >: after the completion of experiment, user can click this icon in the tool bar to create a new analysis method based on the current experiment data; the application software provides user with six analysis methods: **Abs Quant**, **Rel Quant**, **Melting Curve**, **High Resolution Melting**, **Genotyping** and **End Point Fluorescence**.
- <  **Analysis Setting** >: after the completion of experiment, user can click this icon in the tool bar to set the relevant parameters for the current experiment data analysis method.
- <  **Delete Analysis** >: user can click this icon in the tool bar to delete the selected analysis method.

- <  **Export Lis** >: after the completion of experiment, user can click this icon in the tool bar to export the current experiment data; user can choose the export file format (.csv/.txt/.xlsx) from the drop-down list.

3.3.1 Run Setting

The operation area on main interface displays **Run Setting** tab by default, as shown in figure D-5.

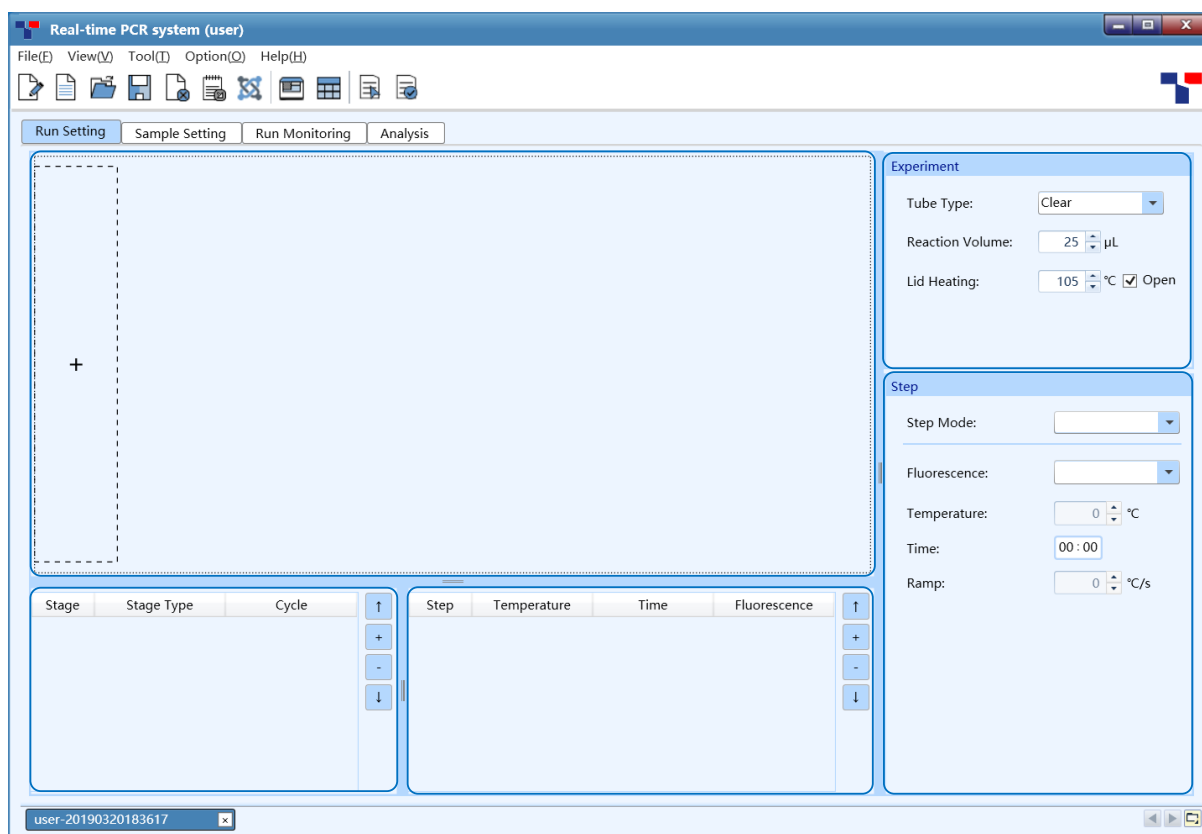


Figure D-5. Run Setting interface

Run Setting

The Run Setting interface consists five parts: temperature program area, stage and step lists, experiment and step editing areas.

1. **Temperature program area**: user can view the temperature program of the current experiment, or edit the temperature program stages and steps in this area, as shown in figure 5.1.

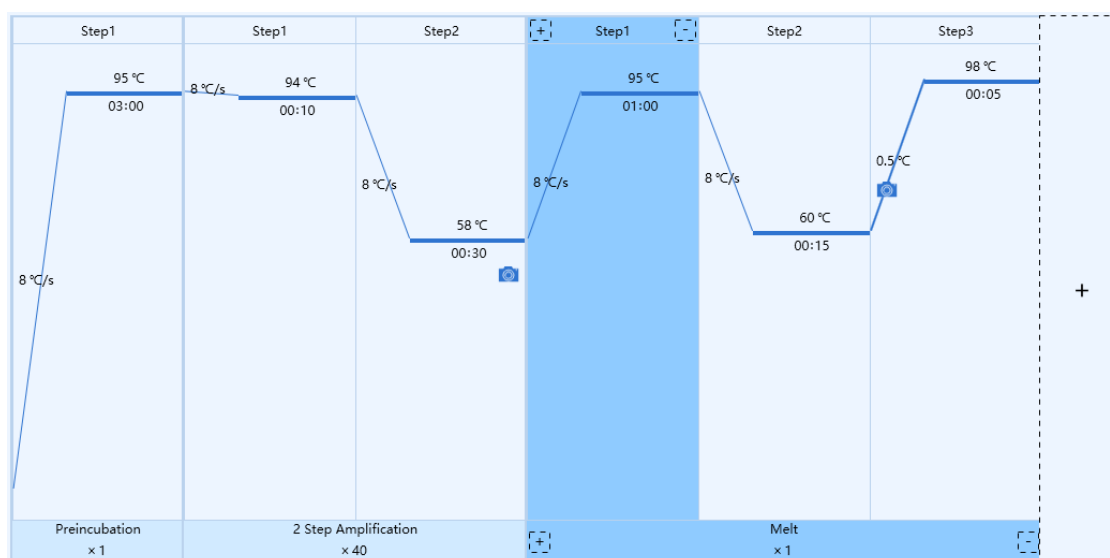





Figure D-5.1 Temperature program area – Example temperature program

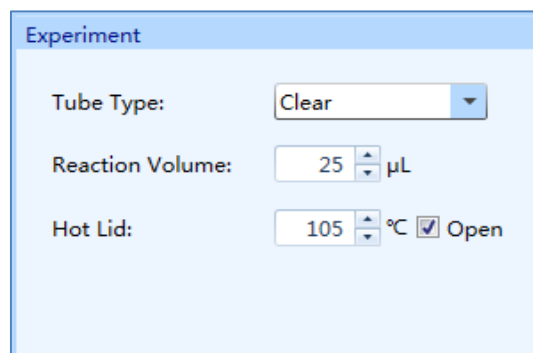
Introduction to Temperature Program Area

- Stages are longitudinal separated; if a stage contains several steps, these steps are separated by solid blue line;
- The corresponding stage type and cycle number are displayed below each stage box and the corresponding step number is displayed above the step box;
- Inside each step box, the corresponding temperature of the current step is shown above the solid blue line and corresponding time of the current step is shown below the solid blue line;
- The solid blue line between two stages/steps represents the temperature ramp, or increment or readings of melting step;
- If user decide to read the fluorescence at a certain step,  icon will be displayed in the corresponding step box.

 **Reminding:** user can hover mouse over the temperature and time of a certain step in the temperature program area, and the application software will display the temperature setting details of this step.

 **Reminding:** user can double click any step box in the temperature program area and edit the current step in the pop-up **Step Setting** dialog box.

2. **Experiment Editing Area:** user can edit the relevant parameters of the current experiment in the experiment editing area, as shown in figure D-5.2.



The 'Experiment' window contains three settings:

- Tube Type:** A dropdown menu currently showing 'Clear'.
- Reaction Volume:** A numeric input box showing '25' followed by a unit selector set to 'μL'.
- Hot Lid:** A numeric input box showing '105' followed by a unit selector set to '°C' and a checked checkbox labeled 'Open'.

Figure D-5.2 Experiment editing area



- ▶ **Tube Type:** user can choose the consumable type of current experiment from drop-down list of **Tube Type**, including **Clear**, **White** and **Frosted**.
- ▶ **Reaction Volume:** user can enter the reaction volume for current experiment in the **Reaction Volume** input box.
- ❗ **Reminding:** the reaction volume setting range is 0μL - 100μL.
- ▶ **Hot Lid:** user can enter the hot lid temperature for current experiment in the **Hot Lid** input box. User can also check the **Open** check box to determine whether to use the hot lid heating function
- ❗ **Reminding:** the hot lid temperature setting the range is 40.0°C - 110.0°C

3. Stage list: as shown in figure D-5.3.

Stage	Stage Type	Cycle	
1	Preincubation	1	↑
2	2 Step Amplification	40	+
3	Melting	1	-
			↓

Figure D-5.3 Stage list





Stage List Introduction



- ▶ **Stage:** display the number of each stage.
- ▶ **Stage Type:** display the type of each stage.
- ▶ **Cycle:** display the cycle number of each stage. User can double click this number, manually enter or use  and  keys to set the cycle number of relevant stage.


Reminding: user can also double click the cycle number below the stage box in the temperature program area to complete the cycle setting.

Reminding: the cycle number setting range is 1-99.

► User can add or delete any stage according to the experiment requirements or adjust the sequence of each stage.

- User can click <  **Add** > icon to add a new temperature stage.
- User can click <  **Delete** > icon to delete the selected stage.
- User can click <  **Up** > icon to move up the selected stage.
- User can click <  **Down** > icon to move down the selected stage.

Reminding: User can also click  icon in the dotted box of temperature program area or click <  **Insert Stage** > icon below any stage box to add a new stage.

Reminding: User can also click <  **Delete Stage** > below any stage box in the temperature program area to delete the stage.

► If user add a new stage, the application software will automatically pop up a stage type window, as shown in figure D-5.4.

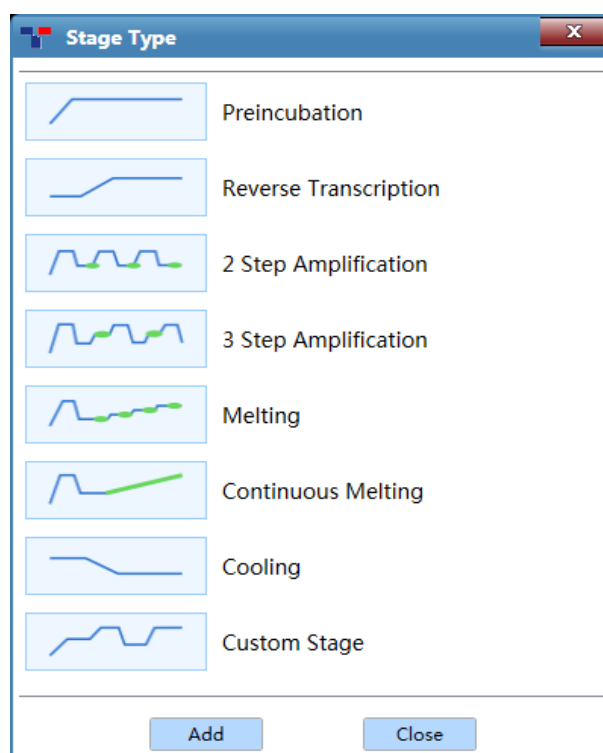



Figure D-5.4 Stage Type window

Introduction to Stage Type Window

- ▶ The application software provides the following stage types: **Preincubation**, **Reverse Transcription**, **2 Step Amplification**, **3 Step Amplification**, **Melting**, **Continuous Melting** and **Cooling**. User can choose the corresponding predefined stage in the stage type window and click **< Add >** to add the selected stage; or double click any stage to add directly.
- ▶ User could select **Custom Stage** and click **< Add >** to define the stage setting according to specific experiment requirements.
- ▶ After adding a stage, please click **< Close >** to return to the run setting interface. The new stage will be added to the temperature program area and stage list.



 **Reminding:** at least one stage should be included in the temperature program.



4. **Step list:** as shown in figure D-5.5.




Step	Temperature	Time	Fluorescence
1	95.0°C	01:00	None
2	60.0°C	00:15	None
3	98.0°C	00:05	Reading

Figure D-5.5 Step list

Step List Introduction

- ▶ **Step:** display the number of each step;
- ▶ **Temperature:** display the target temperature of the current step.;
- ▶ **Time:** display the temperature holding time of the current step;
- ▶ **Fluorescence:** display whether to read fluorescence at the current step;
- ▶ User can add/delete the step according to the experiment requirements or adjust the sequence of each step.
 - User can click  **Add >** icon to add a new temperature step.
 - User can click  **Delete >** icon to delete the selected step.

- User can click <  **Up** > icon to move up the selected step.
- User can click <  **Down** > icon to move down the selected step.

 **Reminding:** user can also click <  **Insert Step** > or <  **Delete Step** > above any step box in the temperature program area to add or delete a temperature step.

5. **Step Setting:** double click any step in the step list or in temperature program area to edit the selected step, the application software will pop up the **Step Setting** dialog box for user to edit the selected step, as shown in figure D-5.6a.

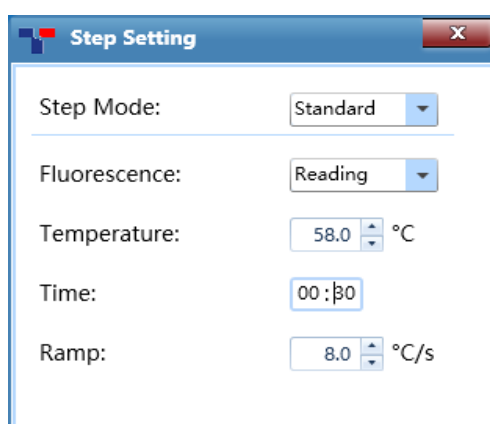




Figure D-5.6a Step Setting dialog box – Standard step mode


Introduction to Step Setting Dialog box

Step Mode: user can set the step mode for the selected step in the drop-down list of *Step Mode*, which including *Standard*, *Touchdown*, *Long* and *Gradient*.


Step Mode \ Model	Gentier 48E	Gentier 48S	Gentier 48R
<i>Standard</i>	✓	✓	✓
<i>Touchdown</i>	✓	✓	✓
<i>Long</i>	✓	✓	✓
<i>Gradient</i>	✓	✗	✓


- Standard Step Mode:** user can set the *Temperature*, *Time*, *Ramp* for the standard step and decide whether to read its *Fluorescence*, as shown in figure D-5.6a.
- **Fluorescence:** user can decide whether to read the fluorescence at current step from the drop-down list of *Fluorescence* according to the experiment requirements.



- **Temperature:** the temperature in °C, which is to be held for a defined time. User can use  and  keys, or manually enter the current step temperature in the **Temperature** input box.

 **Reminding:** the temperature setting range is 0.0°C -100.0°C.

- **Time:** the time for which the temperature is to be held. User can enter the current step time in the **Time** input box.

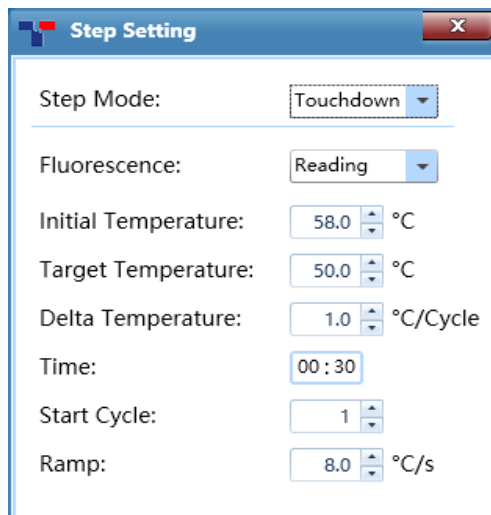
 **Reminding:** the time setting range is 1s~60min.

 **Reminding:** If the current step is the last step of temperature program, user can check the ☐ ∞ check box to set the current step time to infinite.

- **Ramp:** the rate of temperature change(°C) per second. User can use  and  keys, or manually enter the temperature change rate in the **Ramp** input box.

 **Reminding:** the ramp setting range is 0.1°C/s~8°C/s.

- II. **Touchdown Step Mode:** the touchdown step mode allows the temperature program to change the annealing step temperature from the initial temperature to the target temperature as the cycling proceeds. User could set the corresponding parameters in the **Step Setting** dialog box, as shown in figure 5.6b.



The image shows a 'Step Setting' dialog box with the following fields and values:

Parameter	Value	Unit
Step Mode:	Touchdown	
Fluorescence:	Reading	
Initial Temperature:	58.0	°C
Target Temperature:	50.0	°C
Delta Temperature:	1.0	°C/Cycle
Time:	00:30	
Start Cycle:	1	
Ramp:	8.0	°C/s

Figure D-5.6b Step Setting Dialog Box – Touchdown Step Mode

- **Initial Temperature:** the initial value of annealing temperature change range.
- **Target Temperature:** the target value of annealing temperature change range.
- **Delta Temperature:** the temperature change(°C) per cycle.
- **Start Cycle:** the cycle number after which the temperature change is started.

- ❗ **Reminding:** the initial and target temperature setting range is 0.0 °C - 100.0 °C.
- ❗ **Reminding:** the start cycle setting range is 1-max cycle number of the current stage.
- ❗ **Reminding:** the Delta Temperature setting range is 0.1°C- 5.0°C.

III. **Long Step Mode:** the long step mode allows the temperature program to change the elongation step temperature holding time from the initial time to the target time as the cycling proceeds. User could set the corresponding parameters in the **Step Setting** dialog box, as shown in figure 5.6c.

The screenshot shows a 'Step Setting' dialog box with a blue title bar and a close button (X). The settings are as follows:

Parameter	Value	Unit
Step Mode	Long	
Fluorescence	Reading	
Temperature	58.0	°C
Initial Time	00:30	
Target Time	02:00	
Delta Time	00:05	/Cycle
Start Cycle	1	
Ramp	8.0	°C/s

Figure D-5.6c Step Setting Dialog Box – Long Step Mode

- ▶ **Initial Time:** the initial value of the elongation time change range.
- ▶ **Target Time:** the target value of the elongation time change range.
- ▶ **Delta Time:** the time change per cycle.
- ▶ **Start Cycle:** the cycle number after which the time change is started.
- ❗ **Reminding:** the initial initial and target time setting range is 1s~60min.
- ❗ **Reminding:** the delta time setting range is 1s~10min.
- ❗ **Reminding:** the star cycle range is 1~ max cycle number of the current stage.

IV. **Gradient Step Mode:** the gradient step mode allows the sample block to adopt different temperatures. User can set the **Temperature Center** and **Temperature Offset** values for the current gradient step, as shown in figure 5.6d. The system will then automatically calculate the coresponding gradient temperatures.

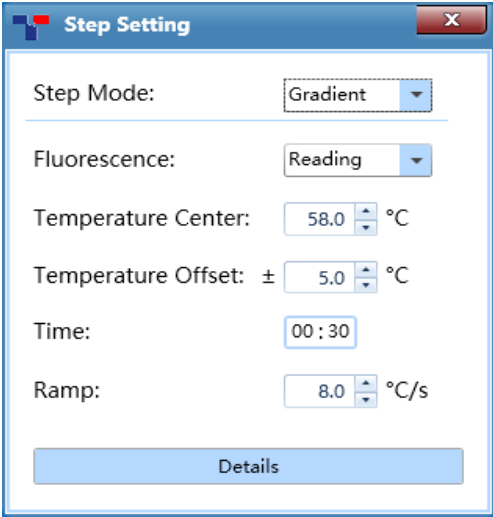





Figure D-5.6d Step Setting Dialog Box – Gradient Step Mode

-  **Reminding:** the gradient step mode is not available on the application software of Gentier 48S instrument model.
 -  **Reminding:** the temperature center setting range is 35.5°C - 99.5°C.
 -  **Reminding:** the temperature offset setting range is 0.5°C - 20.0°C.
- **Details:** User can click *< Details >* to the specific gradient temperatures of each column and the Gradient distribution, as shown in figure D-5.6e.

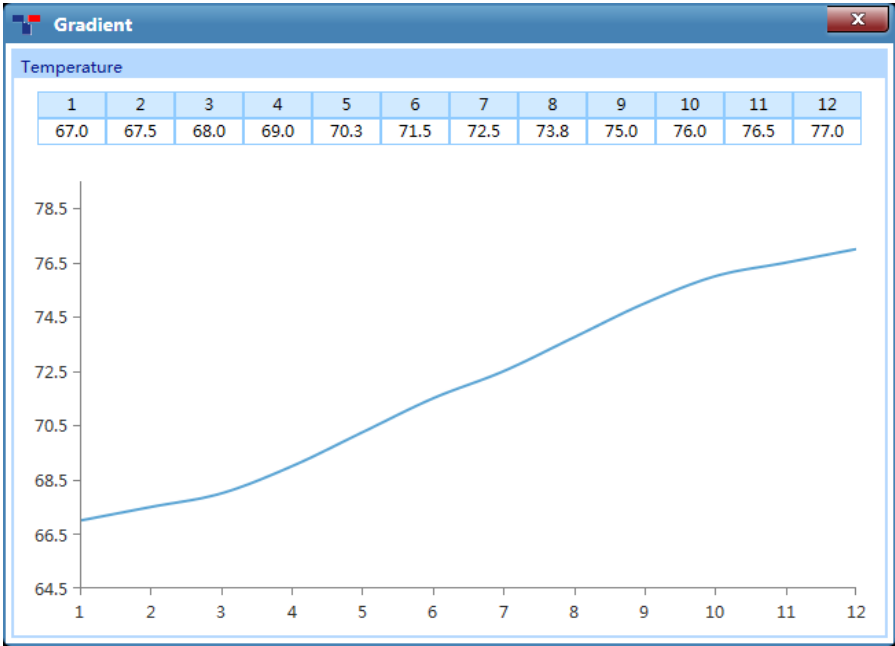


Figure D-5.6e Step Setting Dialog Box – Gradient Temperature Details

- V. **Melting**: melting stage allows the system to read the fluorescence signals after each temperature increment. If the current step is the last step of melting stage, user can set the temperature **Increment** in the **Step Setting** dialog box for the current step, as shown in figure 5.6f.

The 'Step Setting' dialog box for the Melting Step contains three input fields: 'Temperature' set to 98.0 °C, 'Increment' set to 0.5 °C, and 'Time' set to 00:05. Each field has up and down arrow buttons for adjustment.

Figure D-5.6f Step Setting Dialog Box – Melting Step

- **Increment**: the temperature change(°C) after which the system will read the fluorescence. User can use and keys, or manually enter the temperature increment in the **Increment** input box.

Reminding: the temperature increment setting range is 0.1°C - 5.0°C.

- VI. **Continuous Melting**: continuous melting stage allows the system to read the fluorescence more frequently. If the current step is the last step of continuous melting stage, user can set the fluorescence reading times per °C for the current step in the **Step Setting** dialog box, as shown in figure 5.6g.

The 'Step Setting' dialog box for Continuous Melting Step Mode contains two input fields: 'Readings' set to 10 Readings/°C and 'Temperature' set to 97.0 °C. Each field has up and down arrow buttons for adjustment.

Figure D-5.6g Step Setting Dialog Box – Continuous Melting Step Mode

- **Readings**: the fluorescence reading times per °C. User can use and keys, or manually enter the fluorescence reading frequency in the **Readings** input box.

Reminding: the fluorescence reading frequency setting range is 2 readings/°C – 15 readings/°C.

3.3.2 Sample Setting

After finishing the experiment run setting, user can click **Sample Setting** tab to enter the sample setting interface, which is consisted of sample setting area and sample property editing area, as shown in figure D-6.

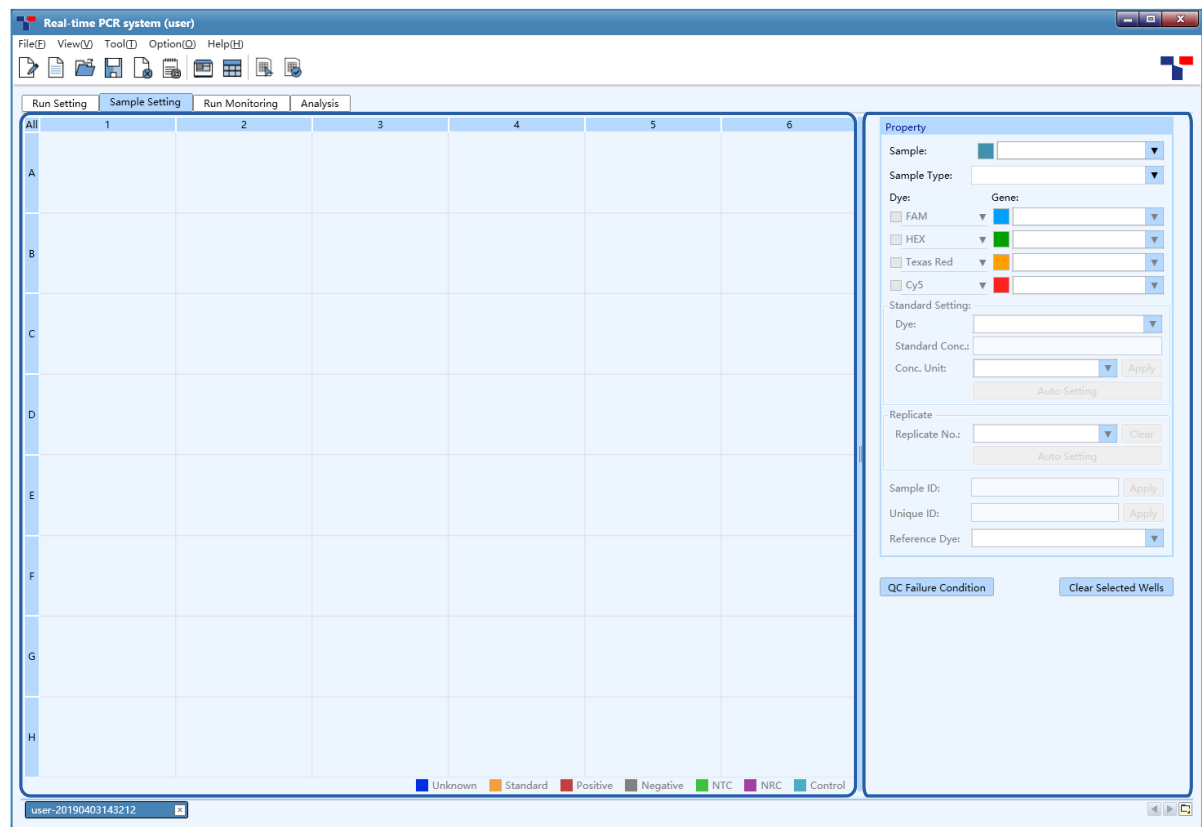


Figure D-6. Sample Setting interface

Sample Setting Operation

- 1. Sample Setting Area:** After entering the sample setting interface, user shall first choose a sample well in the sample setting area, as shown in figure D-6.1. The well distribution in sample setting area corresponds to the sample block well distribution in loading platform. There are 48sample wells in total arranged in 12 columns (1-6) and 8 rows (A-H).

All	1	2	3	4	5	6
A						
B						
C						
D						
E						
F						
G						
H						

Figure D-6.1 Sample Setting interface - Sample Setting area

Introduction to Sample Setting Area

- User can select the relevant sample well in the sample setting area and operations as follows:
 - Click and select a single well in sample setting area to.
 - Press **[Ctrl]** key on computer keyboard and click corresponding wells to select multiple discontinuous wells;
 - Press **[Ctrl]** key on computer keyboard and click corresponding column number to select multiple discontinuous columns of wells;
 - Press **[Ctrl]** key on computer keyboard and click corresponding row number to select multiple discontinuous rows of wells.
 - Click the left mouse button and drag the mouse in sample setting area to select multiple continuous wells;
 - Click the left mouse button and drag the mouse on the column number to select multiple continuous columns of wells, or
 - Click the left mouse button and drag the mouse on the row number to select multiple continuous rows of wells.
 - Click **<All>** on the top left corner of sample setting area to select all sample wells.

2. **Sample Property Setting Area:** after selecting one or more sample wells in the sample setting area, user can set the properties for the corresponding wells in the sample property setting area.
- **Sample Property Box:** user shall edit the *Sample*, *Sample Type*, *Dye* and *Gene* for the selected wells, as shown in figure D-6.2.

Property	
Sample:	 <input type="text"/>
Sample Type:	Unknown <input type="button" value="v"/>
Dye:	Gene:
<input checked="" type="checkbox"/> FAM <input type="button" value="v"/>	 <input type="text"/>
<input type="checkbox"/> HEX <input type="button" value="v"/>	 <input type="text"/>
<input type="checkbox"/> Texas Red <input type="button" value="v"/>	 <input type="text"/>
<input type="checkbox"/> Cy5 <input type="button" value="v"/>	 <input type="text"/>
<input type="checkbox"/> Alexa Fluor 680 <input type="button" value="v"/>	 <input type="text"/>
<input type="checkbox"/> FRET <input type="button" value="v"/>	 <input type="text"/>

Figure D-6.2 Sample Setting interface - Sample Property Box

- a. **Sample:** user can enter the relevant sample name for the selected well in *Sample* input box or select a sample name from the drop-down list;



Reminding: user can click the color block before the *Sample* input box to select a data analysis color for all samples with the current sample name.

- b. **Sample Type:** user can set the relevant sample type for the selected wells from the *Sample Type* drop-down list. The optional sample types and their corresponding colors in sample setting area are *Unknown*, *Standard*, *Positive*, *Negative*, *NTC*, *NRC*, and *Control*.

- c. **Dye:** user can check the relevant dye check box to determine the detection dyes for the selected wells.



Reminding: figure D-6.2 takes Gentier 48E/48S type instrument models as an example. The application software displays the corresponding dye of four channels; and the Gentier 48R instrument models will display the corresponding dye of two channels.

- d. **Gene:** user can enter the relevant gene name for the selected wells in *Gene* input box or select a gene name from the drop-down list;

Reminding: user can click the color block before the **Gene** input box to select a data analysis color for all samples with the current gene name.

- **Standard Setting:** if the sample type for the selected well is **Standard**, the **Standard Setting** box will be activated, as shown in figure D-6.3.

Figure D-6.3 Sample Setting interface - Standard Setting box

- a. **Dye:** user can set the dye for corresponding standard curve from the **Dye** drop-down list.

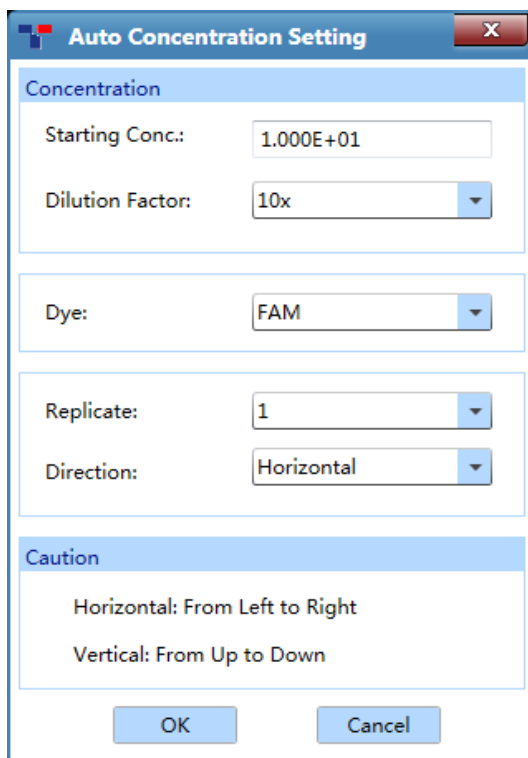
Reminding: please first set the detection dye for the selected standard samples in the sample property box, and then set the dye for the corresponding standard curve from the **Dye** drop-down list in the standard setting box.

- b. **Standard Conc.:** user can select a single standard sample and enter its standard concentration in **Standard Conc.** input box.

- c. **Conc Unit:** user can select a single standard sample and set its concentration unit from **Conc Unit** drop-down list; The application software provides two default concentration units: **IU/ml** and **Copies/ml**.

- d. After editing the standard concentration and concentration unit for the selected standard sample, please click **< Apply >** to confirm the relevant settings.

- e. For a series of standard samples diluted according to a certain dilution factor, user can select corresponding standard sample wells in the sample setting area and let the application software to automatically calculate and set the relevant standard concentration for this series of standard samples. Please click **< Auto Setting >** and the auto concentration setting window will pop up automatically, as shown in figure D-6.4.



The image shows a software window titled "Auto Concentration Setting". It contains several input fields and dropdown menus. The "Concentration" section has a "Starting Conc.:" field set to "1.000E+01" and a "Dilution Factor:" dropdown set to "10x". The "Dye:" dropdown is set to "FAM". The "Replicate:" dropdown is set to "1" and the "Direction:" dropdown is set to "Horizontal". Below these is a "Caution" section with text: "Horizontal: From Left to Right" and "Vertical: From Up to Down". At the bottom are "OK" and "Cancel" buttons.

Figure D-6.4 Sample Setting interface - Auto Concentration Setting Window

- User can set the starting concentration for a series of diluted standard samples in the **Starting Conc.** input box;
- User can select the concentration dilution factor for a series of diluted standard samples from **Dilution Factor** drop-down list;
- User can select the dye for the corresponding standard curve of a series of diluted standard samples from **Dye** drop-down list;
- User can select the number of standard samples from **Replicate** drop-down list;
- User can select the auto concentration setting direction from **Direction** drop-down list: **Horizontal** (from left to right) and **Vertical** (from up to down);
- User can click < **OK** > and the application software will conform to the auto concentration setting direction, calculate and display the concentration of all the selected standard samples.
- User can click < **Cancel** > and the application software will give up the current auto concentration setting and return back to the sample setting interface.

- **Replicate**: according to experiment requirements, user can classify the same samples into one replicate group in the **Replicate** group setting box, as shown in figure D-6.5.

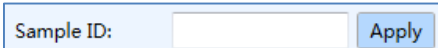
Figure D-6.5 Sample Setting interface - Replicate

- a. **Replicate No.:** user can select the corresponding sample wells in the sample setting area and select the replicate group number from the **Replicate No.** drop-down list, to classify the selected same samples into one replicate group.
- b. User can click < **Clear** > to cancel the replicate group number setting.
- c. The application software can automatically divide multiple replicate groups according to user' requirements. Please first select all wells of same samples in the sample setting area and click < **Auto Setting** >; then the auto setting dialog box will pop up automatically, as shown in figure D-6.6.


Figure D-6.6 Sample Setting interface - Auto Setting

- User can select the number of same samples in each replicate group from the **Replicate Size** drop-down list;
- User can select the starting value for replicate group number from the **Starting Replicate** drop-down list;
- User can select the direction for auto setting from **Direction** drop-down list: **Horizontal** (from Left to right) and **Vertical** (from up to down);


- Click **< OK >** and the application software will automatically divide the replicate group for all the selected samples;
- User can click **< Cancel >** and the application software will give up the current replicate setting and automatically return to the sample setting interface.


► **Sample ID :** 

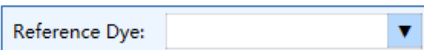
- User can select a well in the sample setting area and set the sample ID for the selected well in the **Sample ID** input box; click **< Apply >** to confirm the sample ID;
- User can also select multiple wells in the sample setting area, and let the application software to set the corresponding sample IDs. Please enter the starting ID number in the **Sample ID** input box and click **< Apply >**; the choose direction dialog box will pop-up, user could set the **Horizontal** or **Vertical** direction and then click **< OK >**, the application software will automatically set the sample IDs for the selected wells according to the direction setting.

► **Unique ID:** 

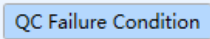
- User can select a well in the sample setting area and set the unique ID for the selected well in the **Unique ID** input box; click **< Apply >** to confirm the unique ID;
- User can also select multiple wells in the sample setting area, and let the application software to set the corresponding unique IDs. Please enter the starting ID number in the **Unique ID** input box and click **< Apply >**; the choose direction dialog box will pop-up, user could set the **Horizontal** or **Vertical** direction and then click **< OK >**, the application software will automatically set the unique ID for the selected wells according to the direction setting.

 **Warning:** The **Sample ID** and **Unique ID** should be at least double-digit, and last two digits must be numbers.

 **Reminding:** The **Sample ID** setting is primarily intended to facilitate operators to identify and differentiate samples. Samples can only be identified and confirmed according to the **Unique ID**.

► **Reference Dye:** 

- User can set the reference dye from the **Reference Dye** drop-down list.

► **QC Failure Condition Setting:** after completing the sample well setting, please click  icon and the QC failure condition setting dialog box will pop up automatically, as shown in figure D-6.7.

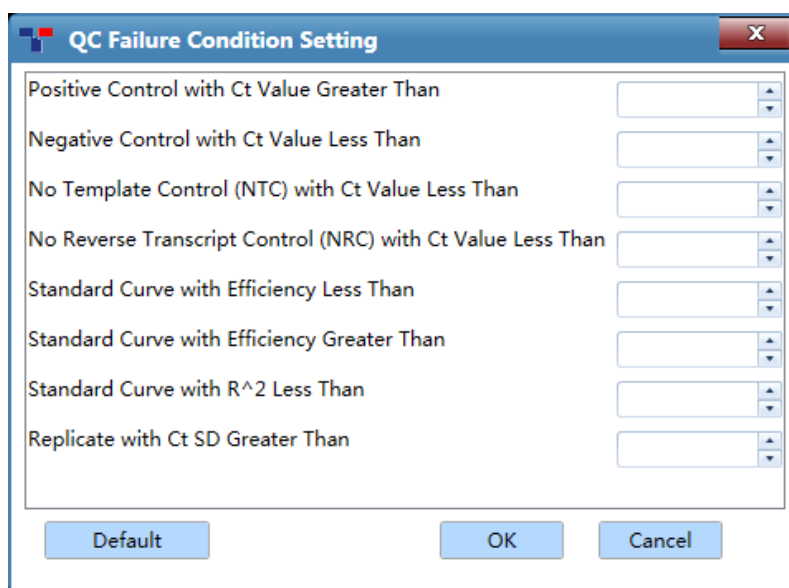


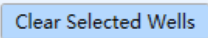


Figure D-6.7 Sample Setting interface - QC Failure Condition Setting

- User can manually enter or use  and  keys to set the QC failure conditions, and click **< OK >** to confirm the setting.
 - User can click **< Default >** to clear all QC failure condition settings.
- **Clear Selected Wells:** user can click  icon to clear the relevant property settings for all selected wells.
-

3.3.3 Run Monitoring

After finishing the experiment run setting and sample setting, user can click **Run Monitoring** tab to enter the run monitoring interface, in order to start running the experiment and monitor the experiment running process, as shown in figure D-7.

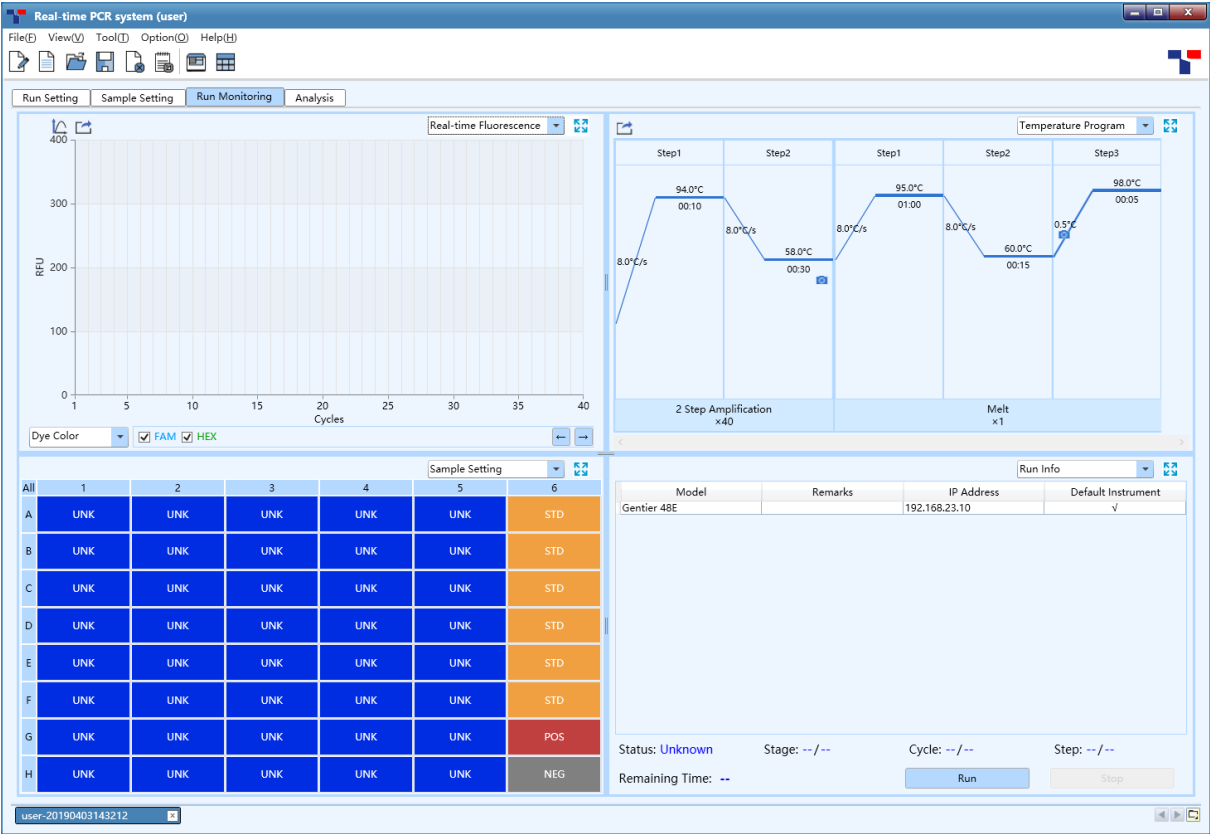



Figure D-7. Run Monitoring interface

Interface Introduction and Parameter Description

The run monitoring interface consists of six functional modules: **Real-time Fluorescence**, **Temperature Program**, **Sample Setting**, **Run Info**, **Sample Info** and **Heat Map**; after entering the run monitoring interface. This interface is divided into four areas and the application software displays four functional modules (**Real-time Fluorescence**, **Temperature Program**, **Sample Setting**, **Run Info**) by default.

- Reminding:** user can select the functional module displayed on the run monitoring interface from the drop-down list on the top right corner of each area.
 - Reminding:** user can click the  **Full Screen** > icon on the top right corner of each area to display the current functional module of the area in full screen.
 - Reminding:** user can click < **Run** > key in the **Run Info** functional module to run the current experiment and monitor the running state of current experiment on the run monitoring interface.
1. **Real-time Fluorescence:** displays the diagram of real-time fluorescence intensity against cycle number of the current running experiment, as shown in figure D-7.1.

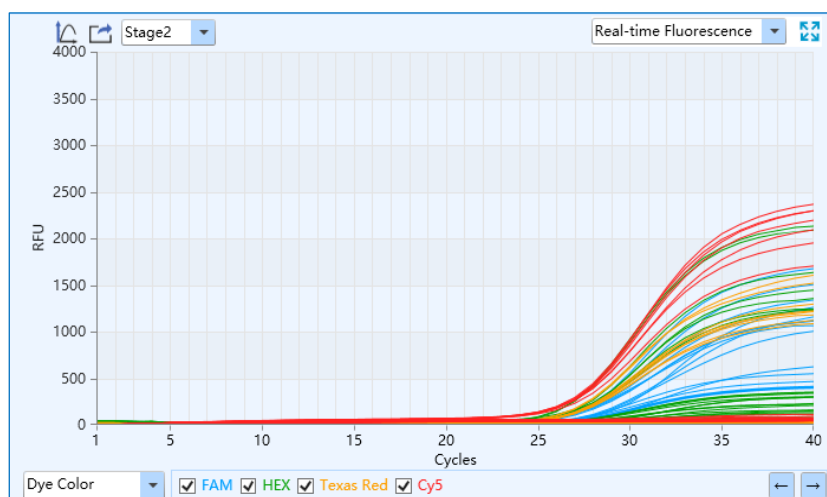






Figure D-7.1 Run Monitoring interface - Real-time Fluorescence function module

- User can set fluorescence curve display mode from the drop-down list on the bottom left of **Real-time Fluorescence** functional module. The real-time fluorescence curve can be displayed according to **Well Color**, **Dye Color**, **Sample Color** and **Gene Color**.
- <  **Y-axis Coordinate Adjustment** >: user can adjust the Y-axis coordinate of the real-time fluorescence monitoring diagram. Please click <  **Y-axis Coordinate Adjustment** > icon and the coordinate range setting box will pop up automatically, as shown in figure D-7.1a. User can select **Automatic** or **Manual** option in the coordinate range setting box to set the Y-axis coordinate. If the **Automatic** option is selected, the application software will automatically adjust the Y-axis coordinate according to the detected fluorescence value; if the **Manual** option is selected, user can manually enter or use  and  keys to set the maximum and minimum value of Y-axis coordinate range in the **Maximum** and **Minimum** input box and click < **OK** > to confirm.

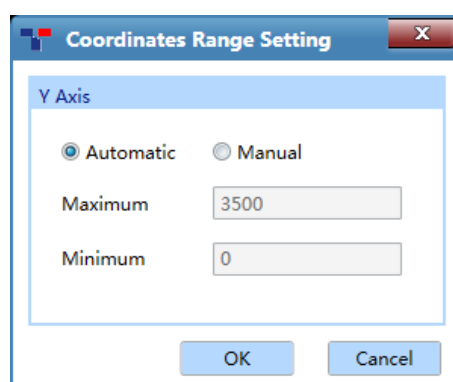


Figure D-7.1a Run Monitoring interface - Coordinate Range Setting

- <  **Export** >: user can click <  **Export** > icon to export the current real-time fluorescence monitoring diagram.

2. **Temperature Program**: display the temperature program of the current running experiment, as shown in figure D-7.2

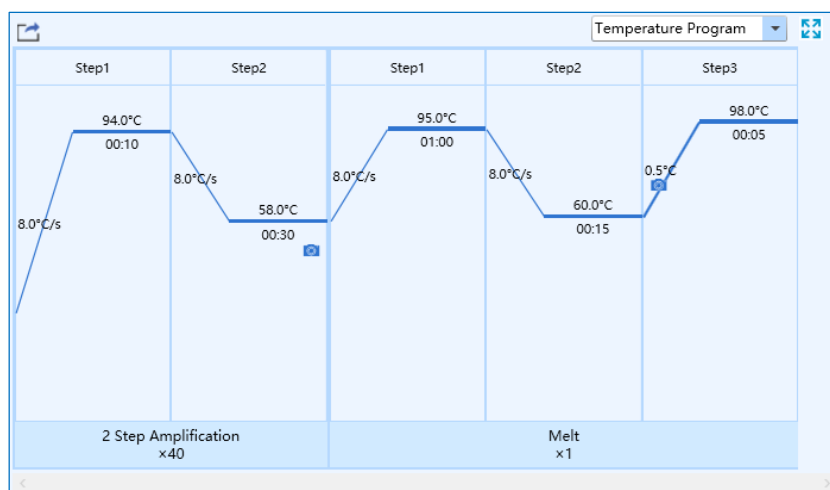



Figure D-7.2 Run Monitoring interface - Temperature Program function module

- **Parameter introduction**: same with the parameters introduced in [D-3.3.1 Run Setting - Temperature program area](#).
 - **Function key introduction**: user can click <  **Export** > icon to export the temperature program monitoring diagram of the current experiment.
3. **Sample Setting**: display the sample well setting of the current running experiment, as shown in figure D-7.3.

Sample Setting						
All	1	2	3	4	5	6
A	UNK	UNK	UNK	UNK	UNK	STD
B	UNK	UNK	UNK	UNK	UNK	STD
C	UNK	UNK	UNK	UNK	UNK	STD
D	UNK	UNK	UNK	UNK	UNK	STD
E	UNK	UNK	UNK	UNK	UNK	STD
F	UNK	UNK	UNK	UNK	UNK	STD
G	UNK	UNK	UNK	UNK	UNK	POS
H	UNK	UNK	UNK	UNK	UNK	NEG

Figure D-7.3 Run Monitoring interface - Sample Setting function module

- ▶ **Parameter introduction:** same with the parameters introduced in *D-3.3.2 Sample Setting - Sample Setting area*.
 - ▶ **Function key introduction:** user can select one or more sample wells in the sample setting function module to display the corresponding fluorescence curve and relevant data in the *Real-time Fluorescence* and *Sample Info* function modules.
4. **Run Info:** display the current instrument information and the real-time running status, such as the *Stage*, *Step*, *Cycle* and *Remaining Time* of of the current experiment, as shown in figure D-7.4.

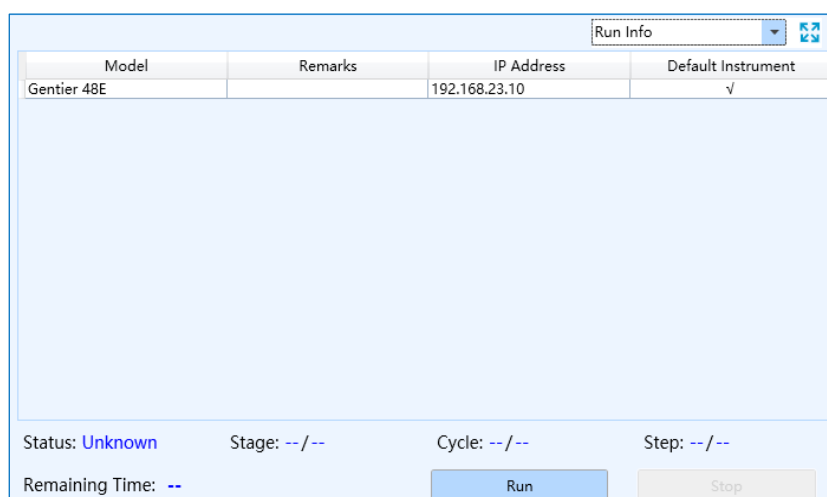


Figure D-7.4 Run Monitoring interface - Run Info function module

- ▶ **Function key introduction:**
 - User can click < **Run** > key to run the current experiment;
 - User can click < **Stop** > key to stop the current experiment;
5. **Sample Info:** display the detailed sample information of the current running experiment, as shown in figure D-7.5.

Sample Info

Well	Sample	Gene	Sample Type	Dye	Replicate
A1	test001	HBV	Unknown	FAM	1
A1	test001	HBV	Unknown	HEX	1
A1	test001	HBV	Unknown	Texas Red	1
A1	test001	HBV	Unknown	Cy5	1
A3	test001	HBV	Unknown	FAM	2
A3	test001	HBV	Unknown	HEX	2
A3	test001	HBV	Unknown	Texas Red	2
A3	test001	HBV	Unknown	Cy5	2
A5	test001	HBV	Unknown	FAM	3
A5	test001	HBV	Unknown	HEX	3
A5	test001	HBV	Unknown	Texas Red	3

Figure D-7.5 Run Monitoring interface - Sample Info function module

6. **Heat Map:** display the real-time fluorescence heat map of the current running experiment, as shown in figure D-7.6.

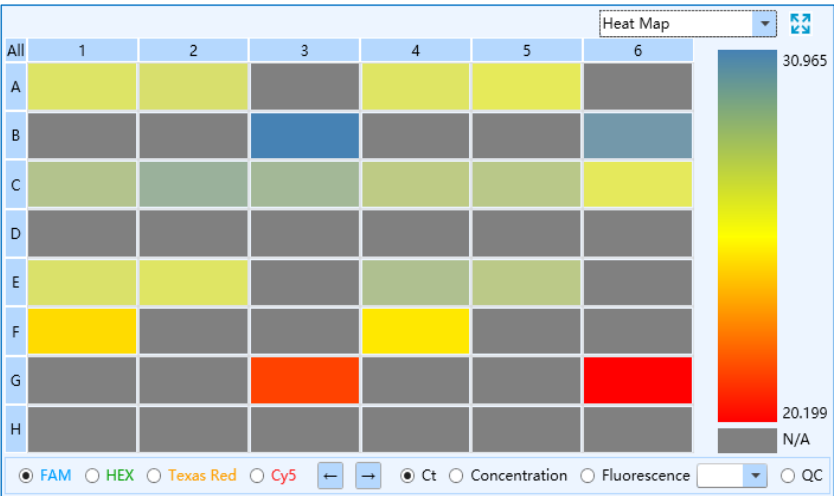



Figure D-7.6 Run Monitoring interface - Heat Map function module

3.3.4 Analysis

After experiment running, user can click the **Analysis** tab to enter the analysis interface. According to the experiment settings and requirements, user shall first select the suitable data analysis method for the current experiment on this interface.

- User can click <  **New Analysis** > icon in Tool Bar and the application software will automatically pop up the new analysis window, as shown in figure D-8.

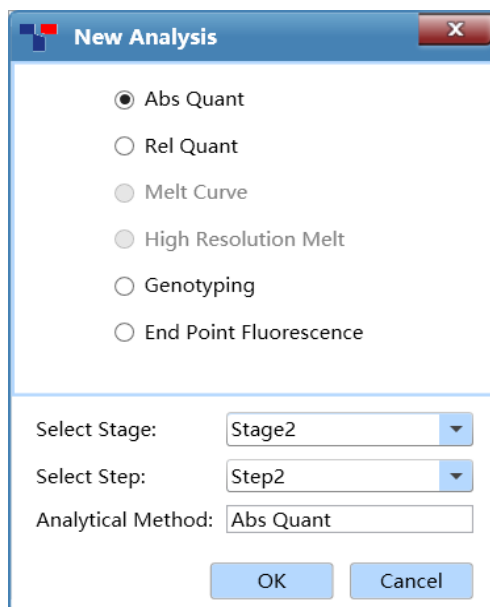


Figure D-8. New Analysis window


- The application software provides user with six analysis methods: ***Abs Quant***, ***Rel Quant***, ***Melting Curve***, ***High Resolution Melting***, ***Genotyping*** and ***End Point Fluorescence***.
- ***Select Stage***: user can choose the stages that need to be analyzed.
 - ***Select Step***: user can choose the step that need to be analyzed.
 - ***Analysis Method***: shown the current analysis method for the current step/stage.
 - User can choose the relevant data analysis method and click < ***OK*** > to confirm.

3.3.4.1 Abs Quant

The absolute quantification experiment is intended to quantify the amount of an interest nucleic acid. Samples with unknown initial nucleic acid quantities are amplified as well as a dilution series of gene-specific standard samples with known concentrations. The measured Ct values of the standard samples are plotted against their known concentrations to obtain a regression line named standard curve. The initial nucleic acid quantities of the samples can be obtained by plotting their Ct values on the standard curve.

The relative quantification analysis interface consists of seven functional modules: **Amplification Curve**, **Standard Curve**, **Sample Setting**, **Result Table**, **Raw Curve**, **Raw Fluorescence** and **Heat Map**. The relative quantification analysis interface is divided into four areas. When user enter the analysis interface, the application software displays four functional modules (**Amplification Curve**, **Standard Curve**, **Sample Setting**, **Result Table**) by default, as shown in figure D-9.

Reminding: user can select the functional module displayed on the analysis interface from the drop-down list on the top right corner of each area.

Reminding: user can click the  **Full Screen** > icon on the top right corner of each area to display the current functional module in full screen.

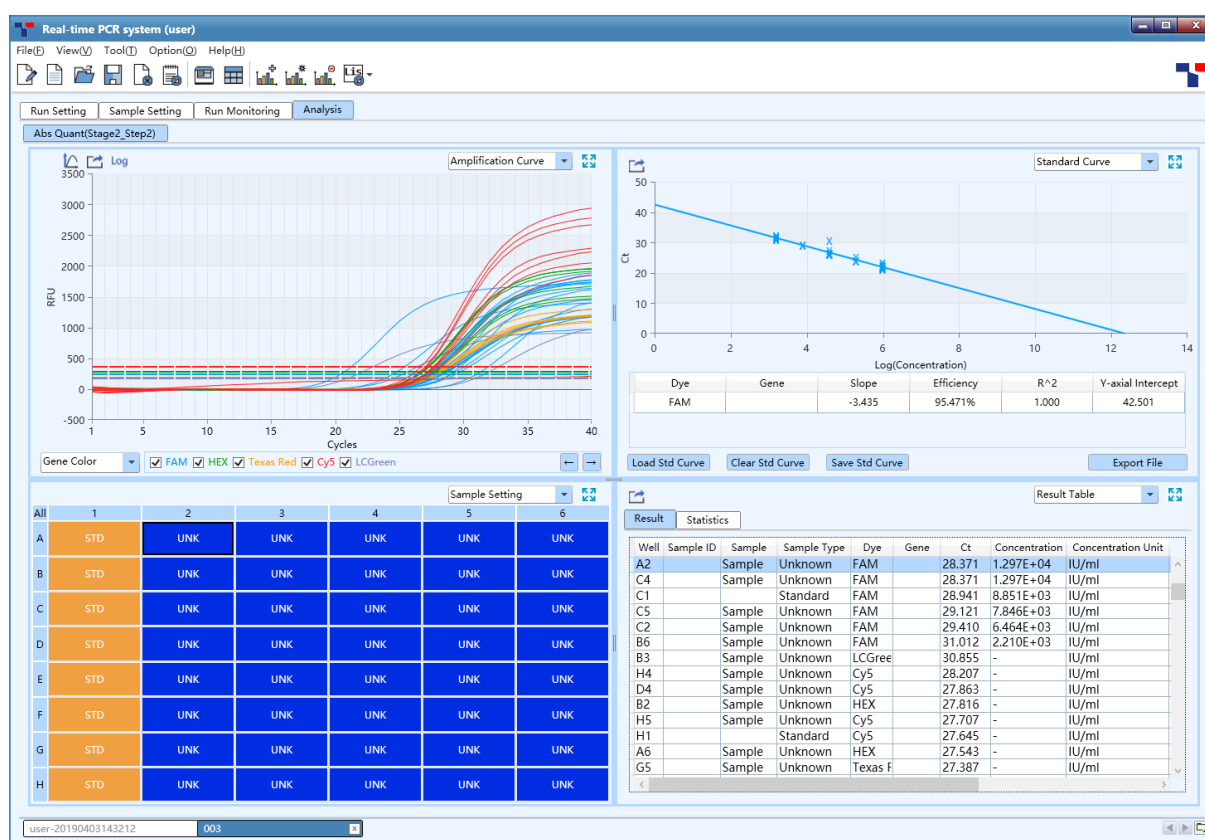


Figure D-9. Absolute quantification analysis interface

Interface Introduction and Parameter Description

- I. **Amplification Curve:** display the diagram of fluorescence intensity against cycle number of the current experiment, as shown in figure D-9.1.

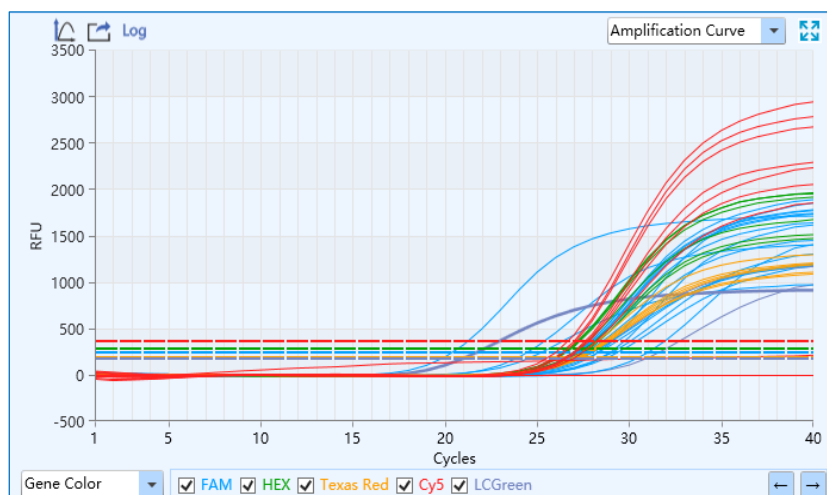






Figure D-9.1 Abs Quant analysis interface - Amplification Curve functional module

► **Parameter introduction:**

- X axis represents the cycle number of the current experiment;
- Y axis represents the Relative Fluorescence Unit (RFU);

► **Function key introduction:**

- User can select the amplification curve display mode from the drop-down list on the bottom left of **Amplification Curve** functional module. The amplification curve can be displayed according to **Well Color**, **Dye Color**, **Sample Color** and **Gene Color**.
- <  **Y-axis Coordinate Adjustment** >: user can adjust the Y-axis coordinate of the amplification curve diagram. Please click <  **Y-axis Coordinate Adjustment** > icon and the coordinate range setting box will pop up automatically, as shown in figure D-9.1a. User can select **Automatic** or **Manual** option in the coordinate range setting box to set the Y-axis coordinate. If the **Automatic** option is selected, the application software will automatically adjust the Y-axis coordinate according to the detected fluorescence value; if the **Manual** option is selected, user can manually enter or use  and  keys to set the maximum and minimum value of Y-axis coordinate range in the **Maximum** and **Minimum** input box and click < **OK** > to confirm.

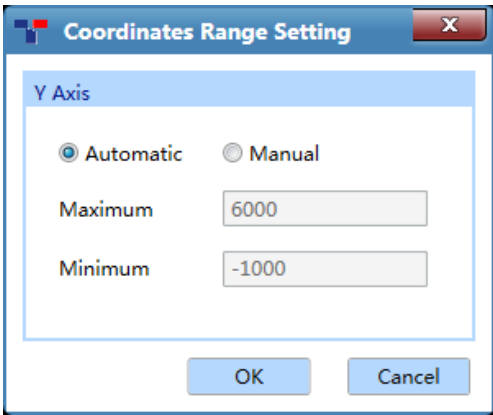






Figure D- 9.1a Abs Quant analysis interface - Coordinate Range Setting

- ▶ <  **Export** >: user can click <  **Export** > icon to export the current amplification curve diagram.
- ▶ <  **Log View** >: user can click <  **Log View** > icon to view the log image of the amplification curve diagram.
- ❗ **Reminding:** user can hover the mouse over a certain amplification curve, the curve will be highlighted and its corresponding channel and well coordinate will be displayed on the right of cursor.

II. **Standard Curve:** displays the threshold Ct value against the initial nucleic acid quantity of each standard sample, as shown in figure D-9.2. For absolute quantification, the standard curve is used to assign initial nucleic acid quantities to unknown samples.

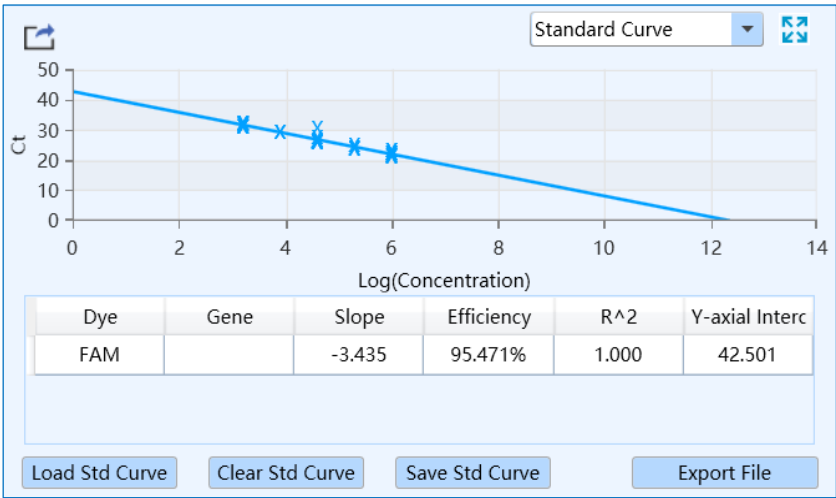


Figure D-9.2 Abs Quant analysis interface - Standard Curve functional module

► **Parameter introduction:**

- X axis represents the log concentration of samples;
- Y axis represents the threshold Ct value;
- **Dye**: display the relevant dye of standard curve;
- **Target Gene**: display the target gene of standard curve;
- **Slope**: display the slope of standard curve;
- **Efficiency**: display the amplification efficiency of standard curve;
- **R²**: display the linear regression coefficient square value of standard curve;
- **Y-axis intercept**: display the Y-axis intercept of standard curve;

► **Function key introduction:**

- **< Load Std Curve >**: click **< Load Std Curve >** and the application software will automatically pop up the standard curve window, which consists of **Import Standard Curve**, **Select Standard Curve** and **Standard Curve Preview** three parts, as shown in figure D-9.2a.

The image shows a software window titled "Standard Curve" with a close button (X) in the top right corner. The window is divided into three main sections:

- Import Standard Curve**: Contains a table with two columns, "No." and "Name". The first row has "1" and "STD1". Below the table are "Import" and "Delete" buttons.
- Select Standard Curve**: Contains a table with three columns: "Dye", "Gene", and "Standard Curve". The "Dye" column lists FAM, HEX, Texas Red, Cy5, and LCGreen. The "Gene" and "Standard Curve" columns are empty. Each row has a dropdown arrow in the "Standard Curve" column.
- Standard Curve Preview**: Contains a graph with a y-axis ranging from 38.8 to 40.6 and an x-axis ranging from 1 to 11. The graph area is currently empty.

At the bottom of the window are "OK" and "Cancel" buttons.

Figure D- 9.2a Abs Quant analysis interface - Standard Curve window

- a. **Import Standard Curve:** user can click < **Import** > key and select a saved standard curve file, then import it to the standard curve list. The corresponding standard curve diagram will be displayed in the **Standard Curve Preview** area below.
- b. **Select Standard Curve:** user can call one or more standard curve from the selected standard curve file according to the current experiment requirements. Please select a standard curve which corresponds to the current experiment setting from the **Standard Curve** drop-down list, and click < **OK** > at the bottom of Standard Curve window to load the selected standard curve to the standard curve functional module.
- c. **Standard Curve Preview:** display the preview of selected standard curve to be loaded.
- < **Clear Std Curve** >: user can click < **Clear Std Curve** > to clear the standard curve displayed on the Standard Curve functional module.
- < **Save Std Curve** >: user can click < **Save Std Curve** > and enter a name in the enter name dialog box for the standard curve file to save; click < **OK** > to confirm, as shown in figure D-9.2b.

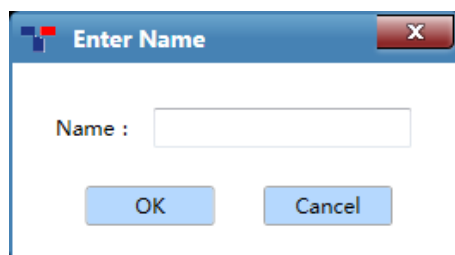


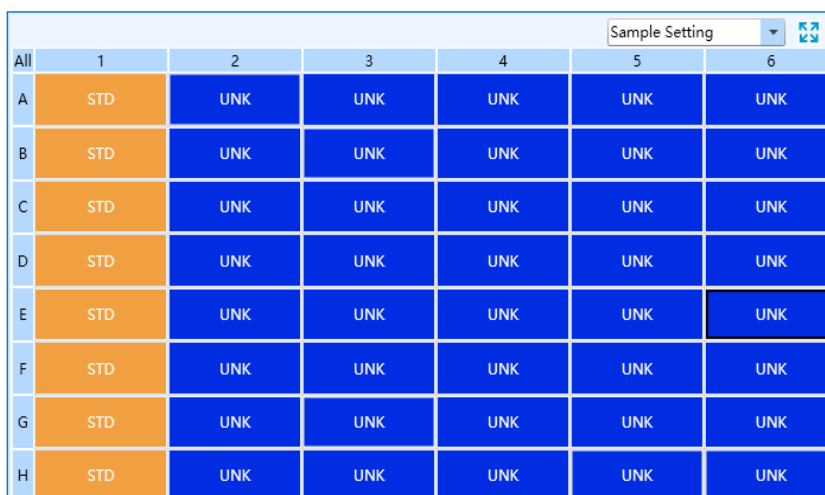


Figure D- 9.2b Abs Quant analysis interface - Enter Name dialog box

- < **Export File** >: user can click < **Export File** > to export the current standard curve file to a designated save path.
 - <  **Export** >: user can click <  **Export** > icon to export the current standard curve diagram.
- III. Sample Setting:** display the well setting of the current experiment, as shown in figure D-9.3.
- **Parameter introduction:** same with the parameters and operations introduced in [D-3.3.2 Sample Setting - Sample Setting area](#).
 - **Function key introduction:** user can click the relevant well in the sample setting area to display the corresponding amplification curve and data in the **Amplification Curve** and **Sample Info** functional modules.

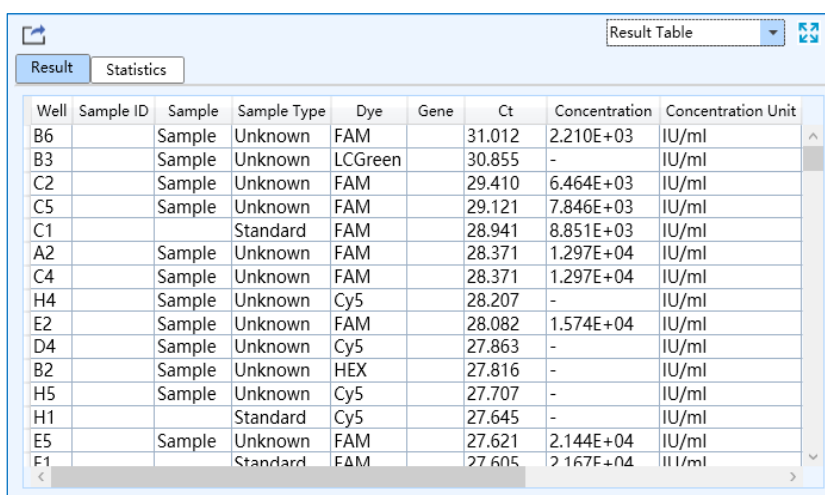


	1	2	3	4	5	6
A	STD	UNK	UNK	UNK	UNK	UNK
B	STD	UNK	UNK	UNK	UNK	UNK
C	STD	UNK	UNK	UNK	UNK	UNK
D	STD	UNK	UNK	UNK	UNK	UNK
E	STD	UNK	UNK	UNK	UNK	UNK
F	STD	UNK	UNK	UNK	UNK	UNK
G	STD	UNK	UNK	UNK	UNK	UNK
H	STD	UNK	UNK	UNK	UNK	UNK

Figure D-9.3 Abs Quant analysis interface - Sample Setting functional module

IV. Result Table: display the sample details and result data of the current experiment. The result table functional module consists of **Result** and **Statistics** sub-tabs.

- The the **Result** sub-tab is selected by default on the result table functional module, which display the sample details and results of the current experiment, as shown in figure D-9.4a.

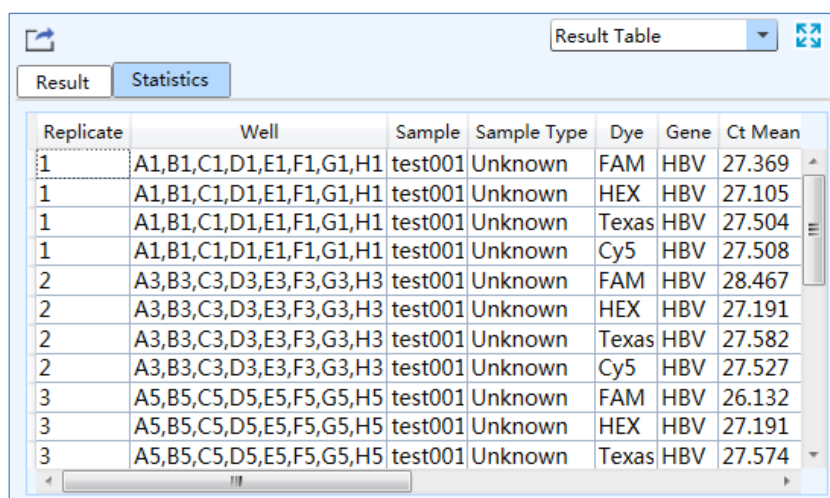


Well	Sample ID	Sample	Sample Type	Dye	Gene	Ct	Concentration	Concentration Unit
B6		Sample	Unknown	FAM		31.012	2.210E+03	IU/ml
B3		Sample	Unknown	LCGreen		30.855	-	IU/ml
C2		Sample	Unknown	FAM		29.410	6.464E+03	IU/ml
C5		Sample	Unknown	FAM		29.121	7.846E+03	IU/ml
C1		Standard	Unknown	FAM		28.941	8.851E+03	IU/ml
A2		Sample	Unknown	FAM		28.371	1.297E+04	IU/ml
C4		Sample	Unknown	FAM		28.371	1.297E+04	IU/ml
H4		Sample	Unknown	Cy5		28.207	-	IU/ml
E2		Sample	Unknown	FAM		28.082	1.574E+04	IU/ml
D4		Sample	Unknown	Cy5		27.863	-	IU/ml
B2		Sample	Unknown	HEX		27.816	-	IU/ml
H5		Sample	Unknown	Cy5		27.707	-	IU/ml
H1		Standard	Unknown	Cy5		27.645	-	IU/ml
E5		Sample	Unknown	FAM		27.621	2.144E+04	IU/ml
F1		Standard	Unknown	FAM		27.605	2.167E+04	IU/ml

Figure D-9.4a Result Table functional module – Result sub-tab

- User can double click the title of each column to sort all sample results according to the content in this column.
- User can click the title of each column and drag the column to the left or right, in order to adjust the sequence of sample results.

- User can click **Statistics** sub-tab to view the statistical results of the current experiment, as shown in figure D-9.4b.



Replicate	Well	Sample	Sample Type	Dye	Gene	Ct Mean
1	A1,B1,C1,D1,E1,F1,G1,H1	test001	Unknown	FAM	HBV	27.369
1	A1,B1,C1,D1,E1,F1,G1,H1	test001	Unknown	HEX	HBV	27.105
1	A1,B1,C1,D1,E1,F1,G1,H1	test001	Unknown	Texas	HBV	27.504
1	A1,B1,C1,D1,E1,F1,G1,H1	test001	Unknown	Cy5	HBV	27.508
2	A3,B3,C3,D3,E3,F3,G3,H3	test001	Unknown	FAM	HBV	28.467
2	A3,B3,C3,D3,E3,F3,G3,H3	test001	Unknown	HEX	HBV	27.191
2	A3,B3,C3,D3,E3,F3,G3,H3	test001	Unknown	Texas	HBV	27.582
2	A3,B3,C3,D3,E3,F3,G3,H3	test001	Unknown	Cy5	HBV	27.527
3	A5,B5,C5,D5,E5,F5,G5,H5	test001	Unknown	FAM	HBV	26.132
3	A5,B5,C5,D5,E5,F5,G5,H5	test001	Unknown	HEX	HBV	27.191
3	A5,B5,C5,D5,E5,F5,G5,H5	test001	Unknown	Texas	HBV	27.574

Figure D-9.4b Result Table functional module– Statistics sub-tab

- V. **Raw Curve:** display the diagram of raw amplification curve without subtract baseline, as shown in figure D-9.5. The parameters and function keys of raw curve functional module are same with those of amplification curve functional module.

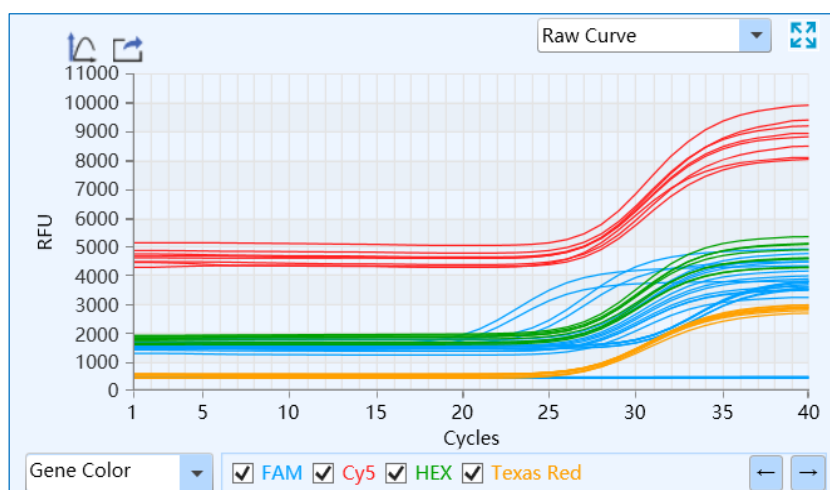


Figure D-9.5 Abs Quant analysis interface - Raw Curve functional module

- VI. **Raw Fluorescence:** display the raw fluorescence data map of the current experiment, as shown in figure D-9.6.

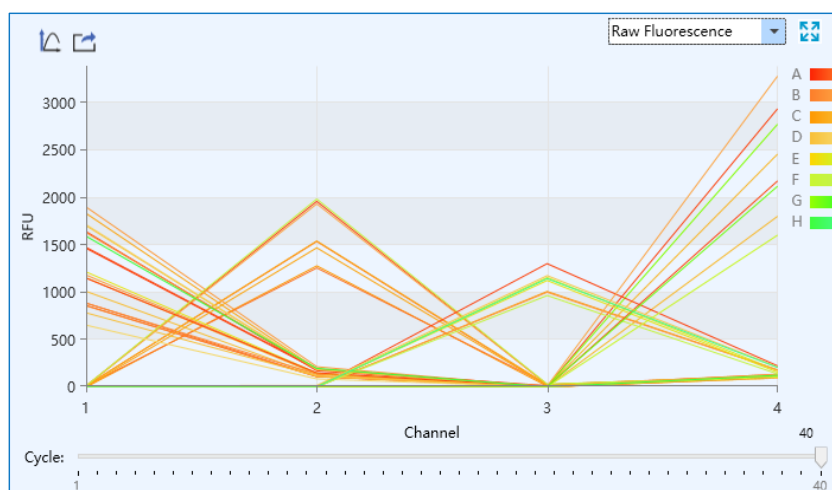




Figure D-9.6 Abs Quant analysis interface - Raw Fluorescence functional module

► **Parameter introduction:**

- X axis shows channel 1-4 (Gentier 48E/48S) or channel 1-2 (Gentier 48R).
- Y axis shows the relative fluorescence unit ((RFU).
- Different colors represent A-H rows of sample wells.

► **Function key introduction:**

- User can slide the **Cycle** slider at the bottom of raw fluorescence functional module to view the raw fluorescence value of different wells at different cycles.
- <  **Y-axis Coordinate Adjustment** > and <  **Export** > keys are same with those of amplification curve functional module.


VII. Heat Map: display the **Ct**, **Concentration**, **Fluorescence** heat map and QC diagram of the current experiment, as shown in figure D-9.7.



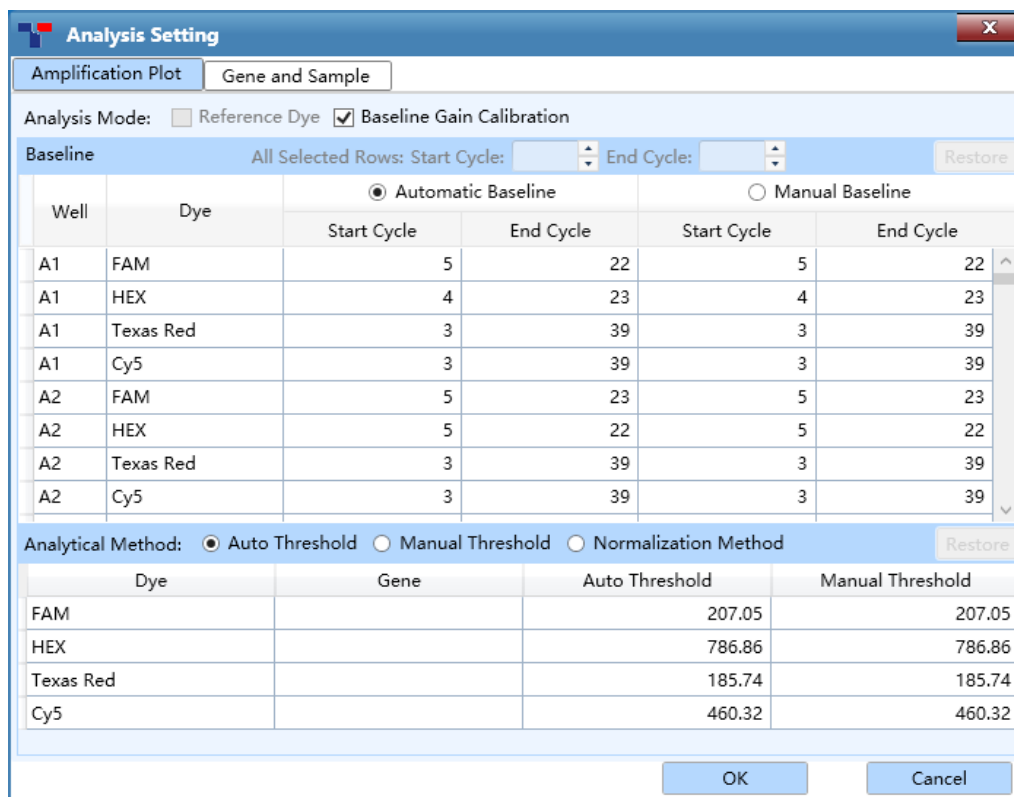
Figure D-9.7 Abs Quant analysis interface – Heat Map functional module

- **Ct**: user can check **Ct** option and click any dye to view the relevant Ct heat map.
- **Concentration**: user can select **Concentration** option and click any dye to view the relevant concentration heat map.
- **Fluorescence**: user can select **Fluorescence** option and select the cycle number from drop-down list; then click any dye to view the the relevant fluorescence heat map at certain cycle.
- **QC**: user can select **QC** option, and if a certain sample or reference conforms to the QC failure condition set on the Sample Setting interface, then the relevant sample well will be displayed as **N/A** on QC diagram.

Abs Quant Analysis Setting

User can click the  **Analysis Setting** > icon in the Tool Bar and the application software will automatically pop up the analysis setting window, which is composed of **Amplification Plot** and **Gene and Sample** two tabs. User can set corresponding parameters for the current experimental data analysis method in the analysis setting window.

- **Amplification Plot** tab: user can set the **Analysis Mode** and **Analysis Method** of the current experiment within the **Amplification Plot** tab, as shown in figure D-10.1.



Analysis Setting

Amplification Plot | Gene and Sample

Analysis Mode: ☐ Reference Dye ☒ Baseline Gain Calibration

Baseline All Selected Rows: Start Cycle: End Cycle: Restore





Well	Dye	Automatic Baseline		Manual Baseline	
		Start Cycle	End Cycle	Start Cycle	End Cycle
A1	FAM	5	22	5	22
A1	HEX	4	23	4	23
A1	Texas Red	3	39	3	39
A1	Cy5	3	39	3	39
A2	FAM	5	23	5	23
A2	HEX	5	22	5	22
A2	Texas Red	3	39	3	39
A2	Cy5	3	39	3	39

Analytical Method: ☒ Auto Threshold ☐ Manual Threshold ☐ Normalization Method Restore

Dye	Gene	Auto Threshold	Manual Threshold
FAM		207.05	207.05
HEX		786.86	786.86
Texas Red		185.74	185.74
Cy5		460.32	460.32

OK Cancel

Figure D-10.1 Abs Quant Analysis Setting interface - Amplification Plot tab

- **Analysis Mode:** user can check the analysis mode check box of **Reference Dye** and **Baseline Gain Calibration** according to the experiment requirements, and set the corresponding parameters in the **Baseline** list.
 - a. **Baseline Gain Calibration:** in general conditions, the application software select the **Baseline Gain Calibration** analysis mode by default, which provides user with **Automatic Baseline** and **Manual Baseline** two baseline setting methods.
 - User can click the **Automatic Baseline** option, the application software will automatically set the baseline for amplification curve, and shown the **Start Cycle** and **End Cycle** of the automatic baseline setting in the columns of the **Baseline** list.
 - User can also click the **Manual Baseline** option, select one well in the **Baseline** list, use  and  keys to manually set the **Start Cycle** and **End Cycle** of the baseline setting for this sample; or choose multiple wells and set the **Start Cycle** and **End Cycle** for **All Selected Rows** on top of the **Baseline** list.
 - User can click **< Restore >** on the top right corner of **Baseline** list to restore the **Start Cycle** and **End Cycle** of the automatic baseline setting.
 - b. **Reference Dye:** If user had set the reference dye in the Sample Setting tab according to the experiment requirements, user can check the analysis mode check box of **Reference Dye** to analysis experiment data.
- **Analytical Method:** the application software provides user with two Ct value analytical methods: **Auto Threshold**, **Manual Threshold** and **Normalization Method**.
 - a. User can click **Auto Threshold** option, the application software will automatically set and show the threshold value for all dyes in its column.
 - b. User can also click **Manual Threshold** option, select one fluorescein and use  and  keys to set the manual threshold value in its column.
 - c. User can click **< Restore >** on the top right corner of threshold list to restore the **Auto Threshold** settings.
 - d. User can click **Normalization Method** option and the application software will automatically calculate the Ct value according to the normalization value of amplification curve.

- **Gene and Sample** tab: user can set the dye, gene and sample required for the analysis of the current experiment within the gene and sample tab, as shown in figure D-10.2.

Analysis Setting

Amplification Plot Gene and Sample

Dye and Gene

No.	Dye	Gene	Remove
1	FAM		<input type="checkbox"/>
2	Cy5		<input type="checkbox"/>
3	HEX		<input type="checkbox"/>
4	Texas Red		<input type="checkbox"/>

Sample

No.	Sample	Remove
1	Sample1	<input type="checkbox"/>
2	Sample2	<input type="checkbox"/>
3		<input type="checkbox"/>

OK Cancel

Figure D-10.2 Abs Quant Analysis Setting interface – Gene and Sample tab

- **Dye and Gene:** display the name of target gene and its marking dye.
 - User can check a certain check box in the **Remove** column and click < **OK** > to remove the experiment data of the selected target gene and its marking dye.
 - Click < **Cancel** > to cancel operation and return to the absolute quantification analysis interface.
- **Sample:** display all the sample name of sample setting.
 - User can check a certain check box in the **Remove** column and click < **OK** > to remove the experiment data of the selected sample name.
 - Click < **Cancel** > to cancel operation and return to the absolute quantification analysis interface.

3.3.4.2 Rel Quant

The relative quantification experiment aims to compare the expression level of two or more genes within a same individual. Usually, endogenous housekeeping genes with constant expression level are taken as the internal reference genes. The relative quantification is realized by measuring the changes of the target genes expression level related to the internal reference genes.

The relative quantification analysis interface consists of eight functional modules: **Amplification Curve**, **Bar Chart**, **Sample Setting**, **Result Table**, **Standard Curve**, **Raw Curve**, **Raw Fluorescence** and **Heat Map**. This interface is divided into four areas and the application software displays four functional modules (**Amplification Curve**, **Bar Chart**, **Sample Setting**, **Result Table**) by default, as shown in figure D-11.

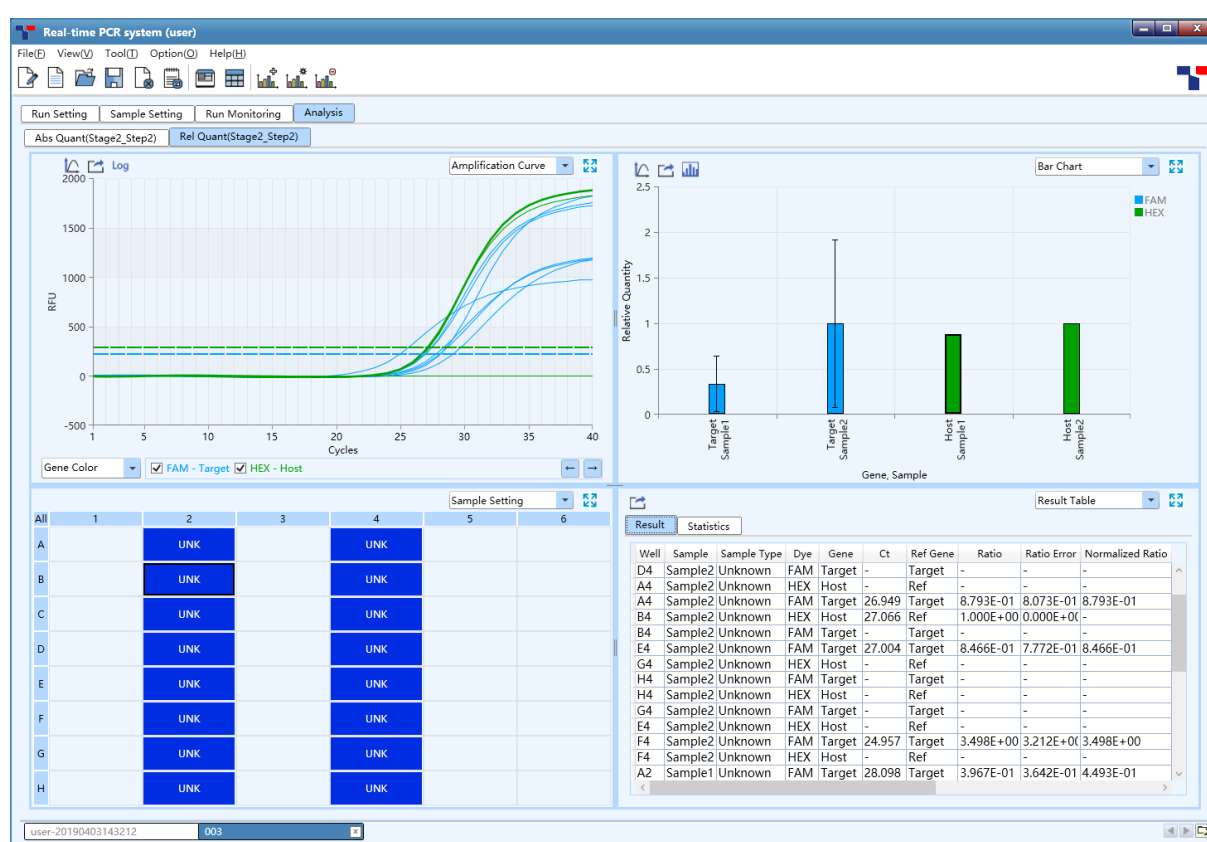


Figure D-11. Relative quantification analysis interface

Interface Introduction and Parameter Description

Reminding: the parameter descriptions of **Amplification Curve**, **Sample Setting**, **Result Table**, **Standard Curve**, **Raw Curve**, **Raw Fluorescence** and **Heat Map** functional modules are same with those introduced in **D-3.3.4.1 Abs Quant**.

- I. **Bar Chart:** display the relative ratio relationship between target gene & sample and reference gene & sample, as shown in figure D-11.1.

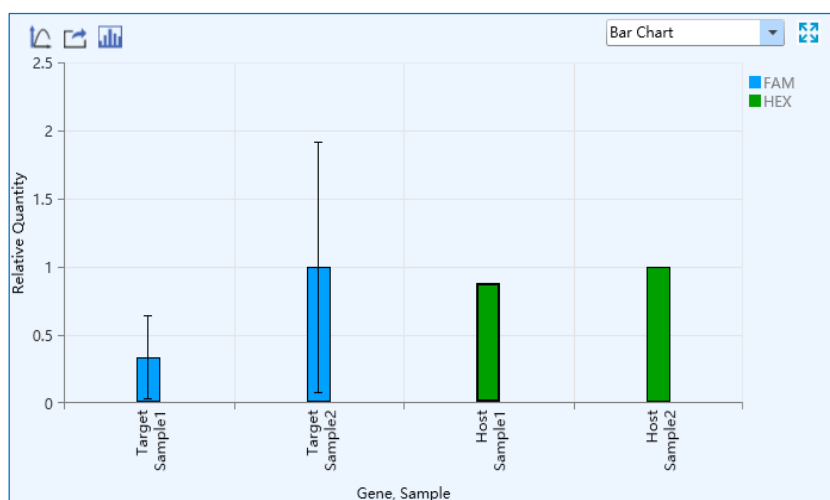






Figure D-11.1 Rel Quant analysis interface - Bar Chart function module

► **Parameter introduction:**

- X axis shows the gene and sample for comparison;
- Y axis shows the **Relative Quantity** ratio of the compared gene and sample;

❗ **Reminding:** the application software set the relative quantity ratio of the reference sample to 1 by default.

► **Function key introduction:**

- The introduction of <  **Y-axis Coordinate Adjustment** > and <  **Export** > keys are same with those keys introduced in [D-3.3.4.1 Abs Quant - Amplification Curve](#).
- <  **Display Setting** >: user can click <  **Display Setting** > icon and the application software will automatically pop up the display setting window, as shown in figure D-11.2.

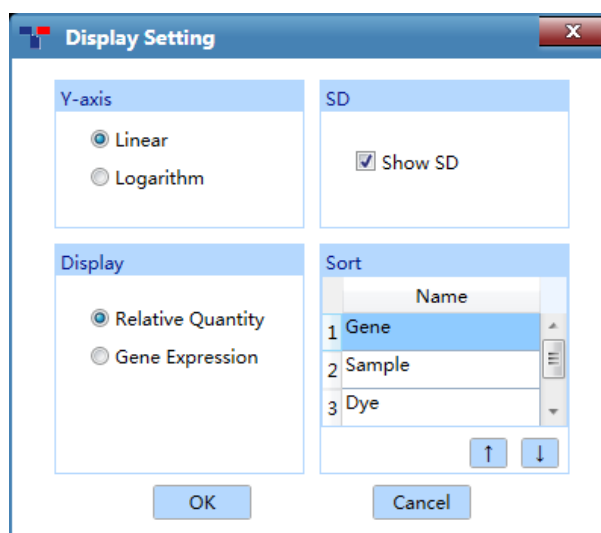




Figure D-11.2 Rel Quant interface - Display Setting window

- a. **Y axis:** If user select **Linear** option, the **Bar Chart** functional module will display the linear coordinates of Y axis; if user select **Logarithm** option, the **Bar Chart** functional module will display the logarithm coordinates of Y axis.
- b. **SD:** user can check **Show SD** check box to display the SD value on **Bar Chart** functional module.
- c. **Display:** user can set the display mode of **Bar Chart** functional module. If user select **Relative Quantity** option, the **Bar Chart** functional module will display the relative quantity ratio of the target gene and reference gene; if user select **Gene Expression** option, the **Bar Chart** functional module will display the relative quantity ratio of target gene & sample and reference gene & sample.
- d. **Sort:** User can use  and  keys to set the display sort order of **Bar Chart** functional module, including **Sample**, **Dye** or **Gene**.

Rel Quant Analysis Setting

After entering the relative quantification analysis interface, the application software will automatically pop up the analysis setting window, which consists of **Gene and Sample**, **Amplification Plot** two tabs. User can set the relevant parameters for current relative quantification analytical method in the analysis setting window.

- I. **Amplification Plot** tab: the introduction of relevant parameters and function keys is same with those introduced in [D-3.3.4.1 Abs Quant - Abs Quant Analysis Setting](#).

- II. **Gene and Sample** tab: user can set the dye, gene, sample and reference required for the analysis of the current experiment within the gene and sample tab, as shown in figure D-12.

Analysis Setting

Amplification Plot Gene and Sample

Dye and Gene



No.	Dye	Gene	Ref Gene	Remove	Efficiency
1	FAM	HPV	<input type="checkbox"/>	<input type="checkbox"/>	100.000 %
2	Cy5	Host	<input checked="" type="checkbox"/>	<input type="checkbox"/>	100.000 %

Sample

No.	Sample	Ref Sample	Remove
1	Sample1	<input type="checkbox"/>	<input type="checkbox"/>
2	Sample2	<input checked="" type="checkbox"/>	<input type="checkbox"/>

OK Cancel

Figure D-12 Rel Quant Analysis Setting interface - Gene and Sample tab

- **Dye and Gene:** display the name of target gene and its marking dye.
 - a. User can check a certain check box in **Ref Gene** column and click **< OK >** to set the reference gene of the current experiment.
 - b. User can check a certain check box in **Remove** column and click **< OK >** to remove the experiment data of corresponding dye labeled genes from the Analysis interface.
 - c. User can click a certain **Efficiency** column, use  and  keys to set the amplification efficiency of the corresponding dye labeled gene.
- ❗ **Reminding:** if a standard curve has obtained for the current experiment, the application software will automatically load the amplification efficiency of the standard curve.
- **Sample:** display all the sample name of sample setting.
 - a. User can check a certain check box in **Ref Sample** column and click **< OK >** to set the reference sample of the current experiment.

- b. User can check a certain check box in **Remove** column and click < **OK** > to remove the experiment data of samples with corresponding sample name from the relative quantification analysis interface.

3.3.4.3 Melting Curve

For intercalating dyes (such as SYBR Green) and non-cleavable hybridization probes, the fluorescence intensity is proportional to the amount of double-strand DNA (dsDNA). However, these dsDNA including specific and non-specific PCR products, which means that the presence of primer dimers and other non-specific products can affect the quality of real-time PCR data.

Melting curve show that the melting degree of dsDNA varies with the rising temperature. Different dsDNA has different melting temperature T_m (the temperature at which, 50% of the DNA has dissociated from double-stranded to single-stranded) owing to different base sequence, fragment length and GC content. Therefore, the melting curve is usually used to analyze whether there are non-specific PCR products in PCR products.

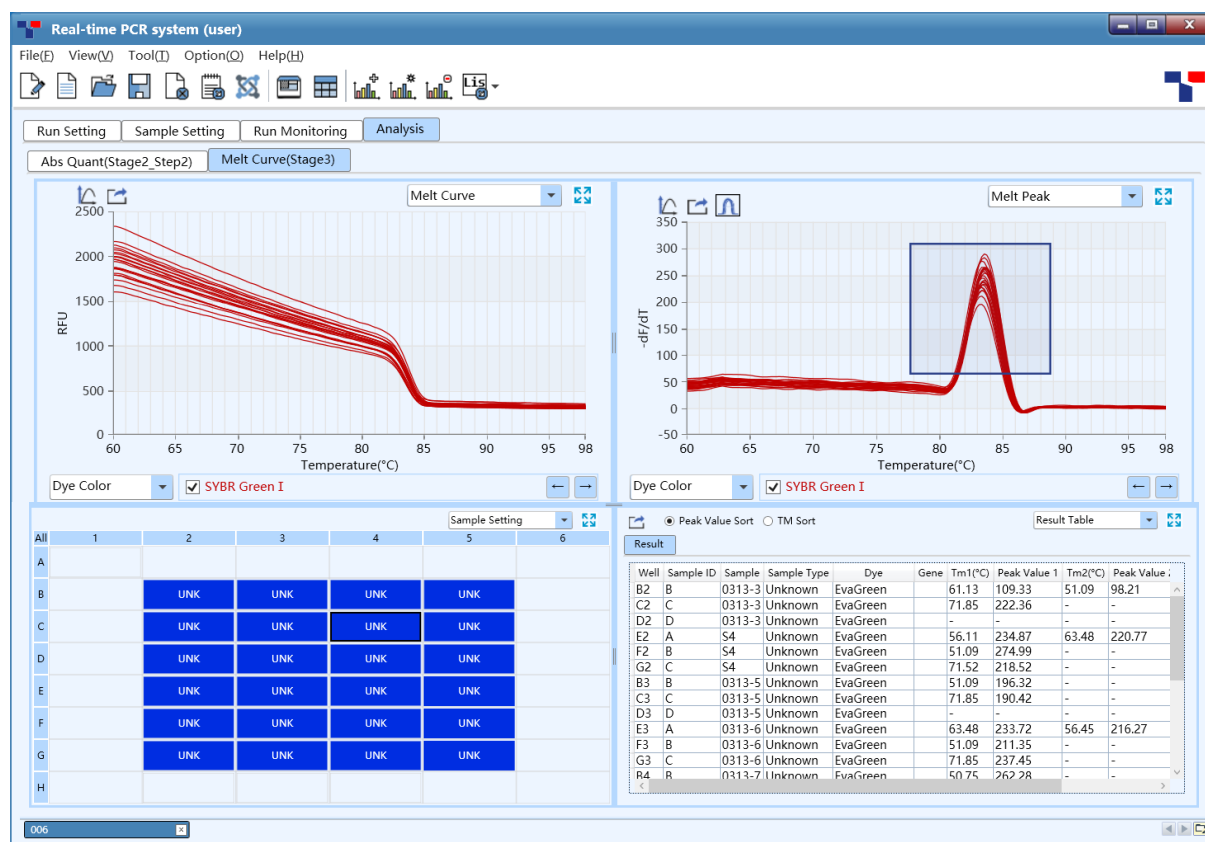


Figure D-13. Melting Curve analysis interface

The melting curve analysis interface consists of five functional modules: **Melting Curve**, **Melting Peak**, **Sample Setting**, **Result Table** and **Heat Map**. The melting curve analysis interface is divided into four areas, the application software displays four functional modules (**Melting Curve**, **Melting Peak**, **Sample Setting**, **Result Table**) by default, as shown in figure D-13.

Interface Introduction and Parameter Description

Reminding: the parameter descriptions of **Sample Setting**, **Result Table** and **Heat Map** functional modules are same with those introduced in **D-3.3.4.1 Abs Quant**.

I. **Melting Curve:** display the fluorescence intensity against the temperature, as shown in figure D-13.1.

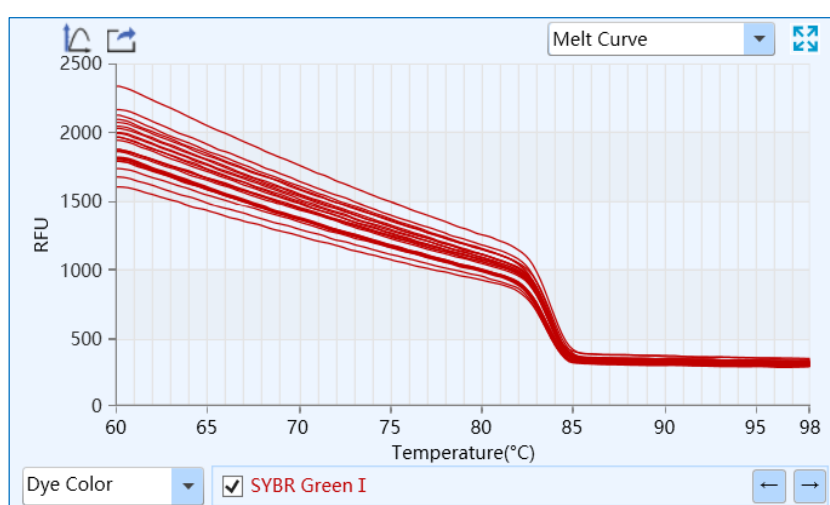




Figure D-13.1 Melting Curve analysis interface - Melting Curve functional module

► Parameter introduction:

- X axis represents the temperature value;
- Y axis represents the Relative Fluorescence Unit (RFU);

► Function key introduction:

- User can select the melting curve display mode from the drop-down list on the bottom left of **Melting Curve** functional module. The melting curve can be displayed according to **Well Color**, **Dye Color**, **Sample Color** and **Gene Color**.
- The introduction of  **Y-axis Coordinate Adjustment** > and  **Export** > keys are same with those keys introduced in **D-3.3.4.1 Abs Quant - Amplification Curve**.

- ❗ **Reminding:** user can hover the mouse over a certain melting curve, the curve will be highlighted and its corresponding channel and well coordinate will be displayed on the right of cursor.
- ❗ **Reminding:** user can drag and drop the left mouse button to zoom in the local melting curve diagram, or click the right mouse button to zoom out.
- II. **Melting Peak:** display the first negative derivative of the fluorescence intensity ($-dF/dT$) respect to the corresponding temperature, as shown in the figure 13.2. If there is only a single melting peak, it means that no non-specific PCR products are detected in the experiment.

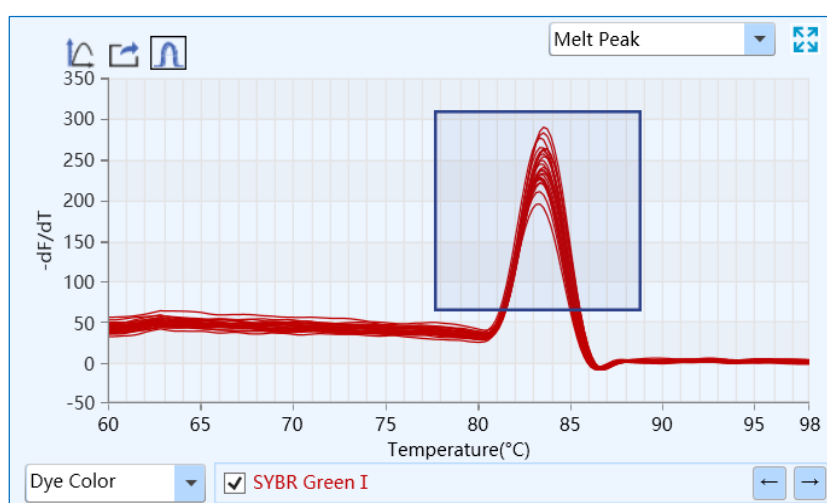







Figure D-13.2 Melting Curve analysis interface - Melting Peak functional module

► **Parameter introduction:**

- X axis represents the temperature value;
- Y axis represents the first negative derivative of the fluorescence intensity against the corresponding temperature;
- User can select the melting peak display mode from the drop-down list on the bottom left of **Melting Peak** functional module. The melting curve can be displayed according to **Well Color**, **Dye Color**, **Sample Color** and **Gene Color**.
- The introduction of <  **Y-axis Coordinate Adjustment** > and <  **Export** > keys are same with those keys introduced in [D-3.3.4.1 Abs Quant - Amplification Curve](#).
- <  **Peak Area** >: user can click <  **Peak Area** > icon to define the melting peak area; each side of the rectangle delimits the temperature and fluorescence range of the peak area.

Melting Curve Analysis Setting

User can click the <  **Analysis Setting** > icon in Tool Bar and the application software will automatically pop up the melting curve analysis setting window, which consists of **Dye and Gene**, **Sample** and **Peak Area** three tabs. User can set the relevant parameters for melting curve analytical method in the melting curve analysis setting window.

I. **Melt Curve** tab: display the parameter range of each peak area, as shown in figure D-13.3.

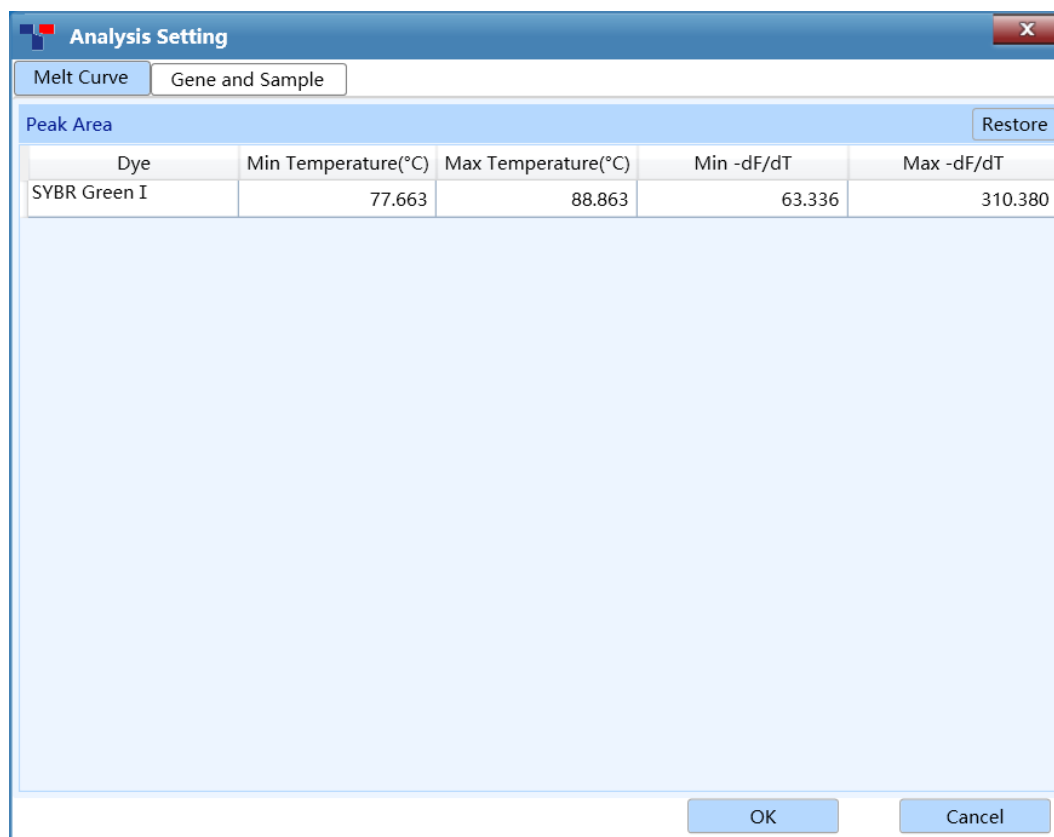






Figure D-13.3 Melting Curve Analysis Setting interface - Melt Curve tab

- **Dye**: display the settings of **Dye** and its marked **Gene** of the current experiment, as shown in figure D-13.3.
- **Peak Area**: display the melting peak area appearing with temperature rise.
- User can use  and  keys to set the temperature range of melting peak area in **Min Temperature** and **Max Temperature** columns.
- User can use  and  keys to set the fluorescence range of melting area in **Min -df/dT** and **Max -df/dT** columns.

II. **Gene and Sample** tab: the introduction of relevant parameters and function keys is same with those introduced in **D-3.3.4.1 Abs Quant - Abs Quant Analysis Setting**.

3.3.4.4 High Resolution Melting

The melting temperature T_m of dsDNA depends on its base sequence, fragment length and GC content. In theory, any base change can cause difference in melting temperature; however, the difference caused by a base change is tiny, usually less than 1 degree. As the resolution of melting curve is not enough, it is required to read the fluorescence signal several times within one degree of temperature change to accurately detect the difference of melting temperature. Therefore, high resolution melting curve is often used for single nucleotide polymorphism (SNP) analysis, mutation scanning, methylation research and genotyping, etc.

The high resolution melting analysis interface consists of seven functional modules: **Melting Curve**, **Normalized Melting Curve**, **Difference Plot**, **Result Table** and **Sample Setting**, **Normalized Peak Melting** and **Heat Map**. The high resolution melting analysis interface is divided into four areas, the application software displays four functional modules (**Melting Curve**, **Normalized Melting Curve**, **Difference Plot** and **Result Table**) by default, as shown in figure D-14.

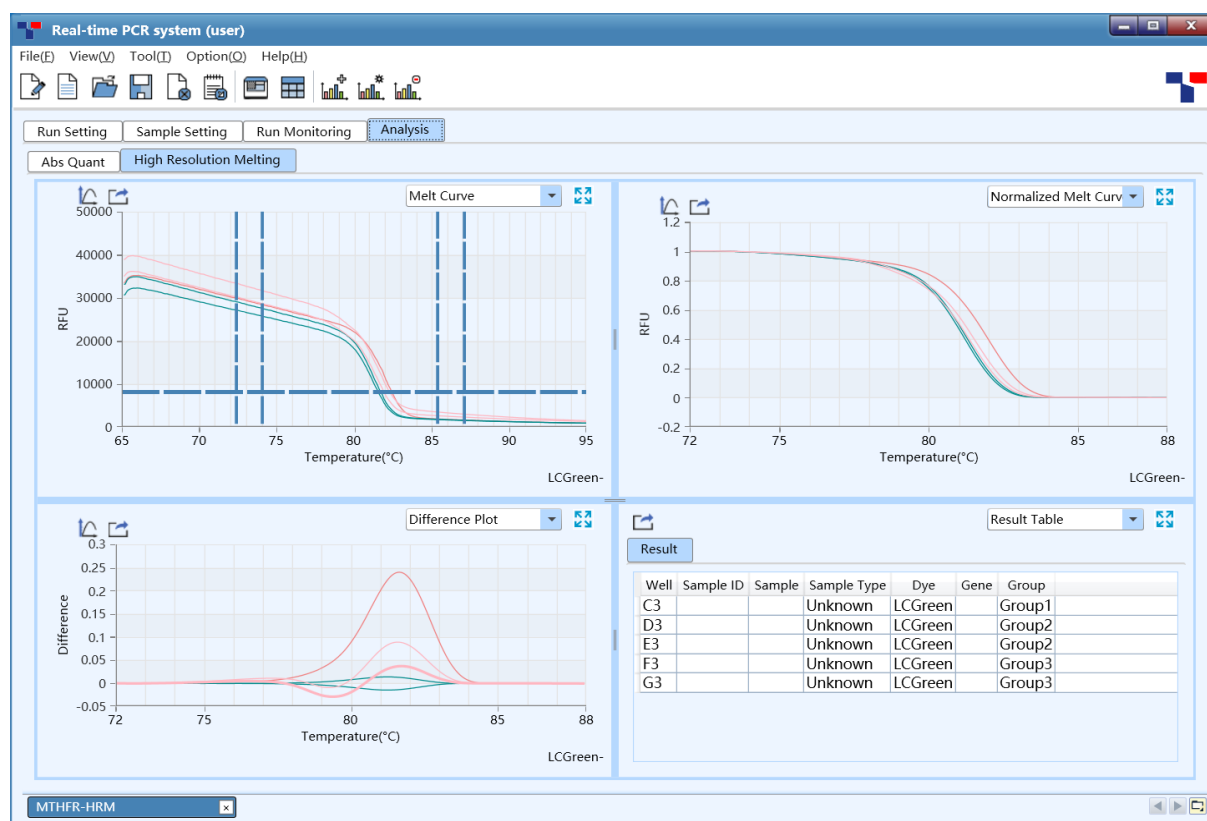


Figure D-14. High Resolution Melting analysis interface

Interface Introduction and Parameter Description

Reminding: the parameter descriptions of *Sample Setting*, *Result Table* and *Heat Map* functional modules are same with those introduced in [D-3.3.4.1 Abs Quant](#).

I. **Melting Curve:** display the fluorescence intensity against the temperature, as shown in figure D-14.1.

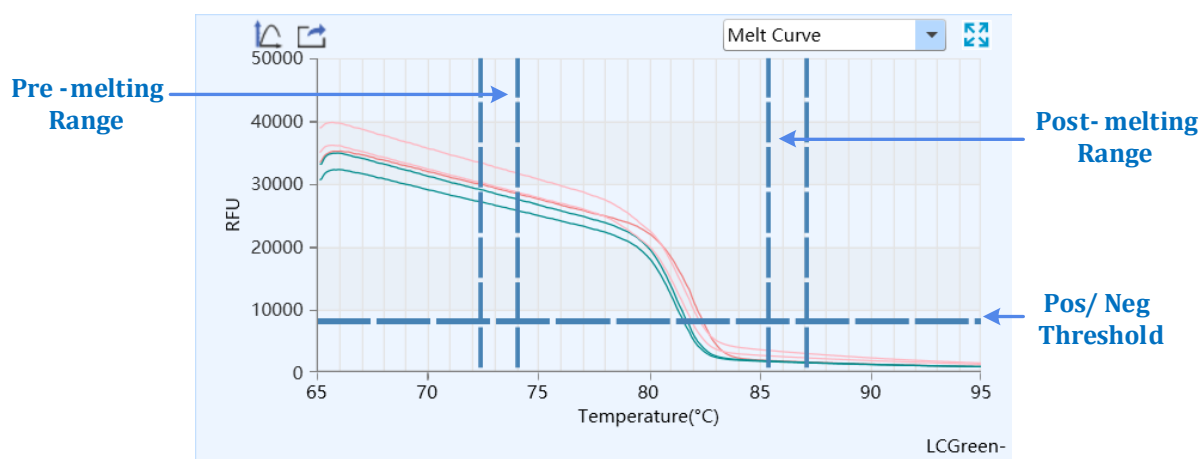




Figure D-14.1 HRM analysis interface - Melting Curve functional module

► Parameter introduction:

- X axis represents the temperature;
- Y axis represents the Relative Fluorescence Unit (RFU);
- By default, the application software will automatically determine normalized area of melting curve. The two vertical sliders on the left are the **Pre-melting Range** threshold lines, which are used to specify the pre-melting temperature range; while two vertical sliders on the right are the **Post-melting Range** threshold lines, which are used to specify the post-melting temperature range; the single line at the bottom is the **Pos/Neg Threshold** line, which is used to determine the positivity or negativity of sample.

► Function key introduction:

- The introduction of  **Y-axis Coordinate Adjustment** > and  **Export** > keys are same with those keys introduced in [D-3.3.4.1 Abs Quant - Amplification Curve](#).
- User can manually drag the vertical sliders or enter relevant parameter in the high resolution melting analysis setting window to specify the normalized area of melting curve.

Reminding: if the normalized area is manually changed, the relevant curve shape on **Normalized Melting Peak, Difference Plot, Normalized Melting Curve** functional modules and the group calling result will change accordingly.

II. **Normalized Melting Curve:** display the normalized melting curve according to the normalization method and fluorescence normalization settings. For more details, please refer to [D-3.3.4.4 High Resolution Melting - High Resolution Melting Analysis Setting](#).

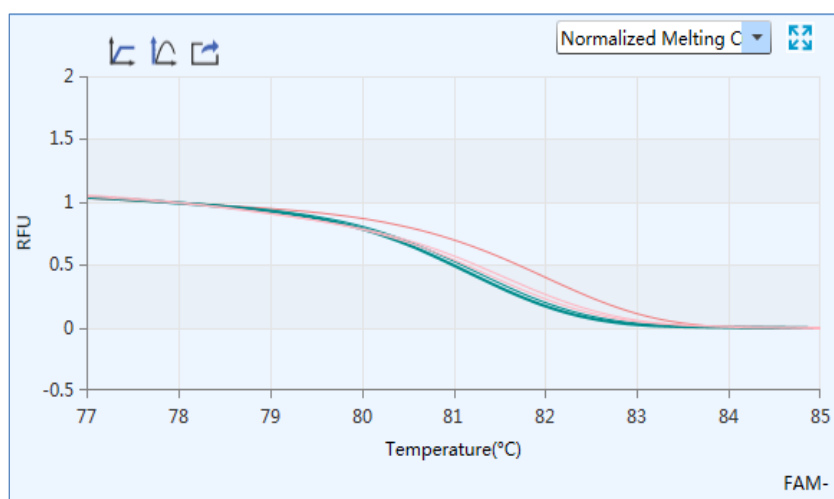


Figure D-14.2 HRM analysis interface - Normalized Melting Curve functional module

III. **Difference Plot:** display the curve after subtracting the baseline group curve, or after normalization and optionally, or after temperature shift. The appearance of the curve on **Difference Plot** functional module depends on the selected baseline group. For more details, please refer to [D-3.3.4.4 High Resolution Melting- High Resolution Melting Analysis Setting](#).

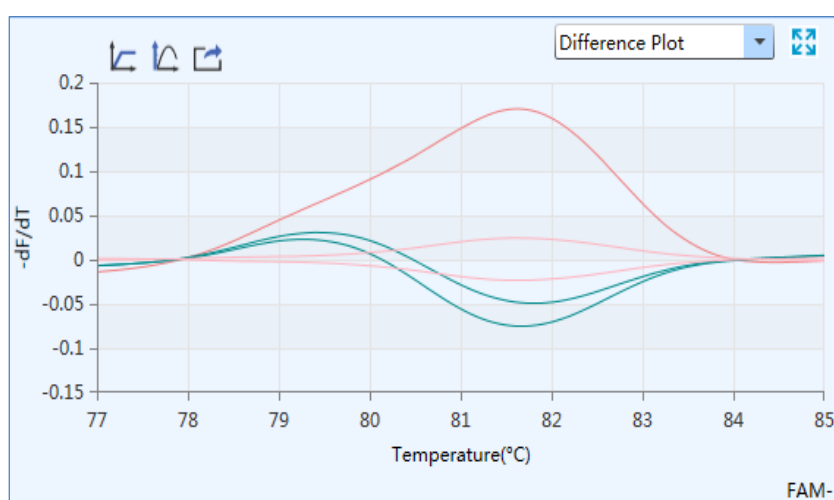


Figure D-14.3 HRM analysis interface - Difference Plot functional module

- IV. **Normalized Melting Peak**: display the first negative derivative of the normalized melting curve, as shown in figure 14.4. The melting temperature range of each sample appears as a peak after normalization, and optionally, temperature shift.

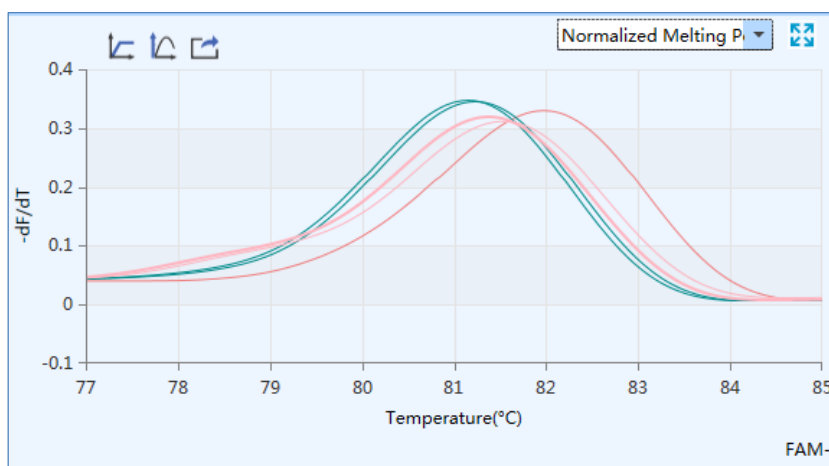



Figure D-14.4 HRM analysis interface - Normalized Peak Melting functional module

High Resolution Melting Analysis Setting

User can click the  **Analysis Setting** > icon in Tool Bar and the application software will automatically pop up the high resolution melting analysis setting window, which consists of **Dye and Gene, Sample** and **High Resolution Melting** three tabs. User can set the relevant parameters for current analytical method in the high resolution melting analysis setting window.

- I. **High Resolution Melting** tab: user can set the **Normalization Method, Fluorescence Normalization, Sensitivity** and **Other** parameters for the current analytical method as shown in figure D-15.

Analysis Setting

High Resolution Melt | Gene and Sample

Normalization Method ☐ Ratio Method ☒ Exponential Method

Fluorescence Normalization

Pre-Melt Range from: 72.395 Default 77.341 to: 74.089 Default 78.328

Post-Melt Range from: 85.435 Default 83.755 to: 87.129 Default 84.742

Sensitivity

Delta Tm Discrimination: 0.50 Default 0.50

Curve Shape Discrimination: 0.50 Default 0.50

Other

Pos/Neg Threshold: 8,015.60 Default 7611.22





☐ Temperature Compensation 0.0665 Default 0.0560

Baseline Group: Group2

OK Cancel

Figure D- 15. High Resolution Melting Analysis Setting interface - High Resolution Melting tab


- **Normalization Method:** the application software provides two normalization methods: **Ratio Method** and **Exponential Method**. The **Ratio Method** option is selected by default.
- **Fluorescence Normalization:** used to set the temperature range for the fluorescence normalization of the melting curve.
 - a. **Pre-Melt Range:** user can use and keys to set the pre-melting temperature range of the normalization area of the melting curve.
 - b. **Post-Melt Range:** user can use and keys to set the post-melting temperature range of the normalization area of the melting curve.
- **Sensitivity:** used to set the discrimination sensitivity of melting curve.
- ❗ **Reminding:** by default, the application software will automatically set the corresponding parameters for the fluorescence normalization of melting curve; user can also drag the relevant threshold lines (vertical sliders) on the **Melting Curve** functional module to determine the normalized area of melting curve.

- a. **Delta T_m Discrimination**: the default Delta T_m discrimination sensitivity provided by the amplification software is 0.5; reduce this value, the application software will grouping the melting curve at a higher temperature resolution.
- b. **Curve Shape Discrimination**: the default curve shape discrimination sensitivity provided by the amplification software of is 0.5; reduce this value, the application software will grouping the melting curve at a higher shape resolution.
- **Other:**
 - a. **Pos/Neg Threshold**: user can use  and  keys to set the threshold value for the application software to determine the positivity or negativity of samples
 - b. **Temperature Compensation**: user can check **Temperature Compensation** check box and the application software will automatically calculate the temperature compensation value between sample block wells. User can use  and  keys to change the temperature compensation value.
 - c. **Baseline Group**: user can set the baseline group gene for the **Difference Plot** functional module.
- II. **Gene and Sample** tab: the introduction of relevant parameters and function keys is same with those introduced in [D-3.3.4.1 Abs Quant - Abs Quant Analysis Setting](#).

3.3.4.5 Genotyping

The Genotyping analysis adopts two sequence-specific probes labeled with different dyes to identify the wild type and mutation alleles, respectively. After running the experiment program, the application software will automatically detect the end point fluorescence and distinguish different genotypes based on the distribution of two dyes on the scatter plot.

The Genotyping analysis interface consists of seven functional modules: **Amplification Curve**, **Scatter Plot**, **Sample Setting**, **Result Table**, **Raw Curve**, **Raw Fluorescence** and **Heat Map**. The Genotyping analysis interface is divided into four areas, the application software displays four functional modules (**Amplification Curve**, **Scatter Plot**, **Sample Setting**, **Result Table**) by default, as shown in figure D-16.

 **Reminding**: the parameter descriptions of **Amplification Curve**, **Sample Setting**, **Raw Curve**, **Raw Fluorescence** and **Heat Map** functional modules are same with those introduced in [D-3.3.4.1 Abs Quant](#).

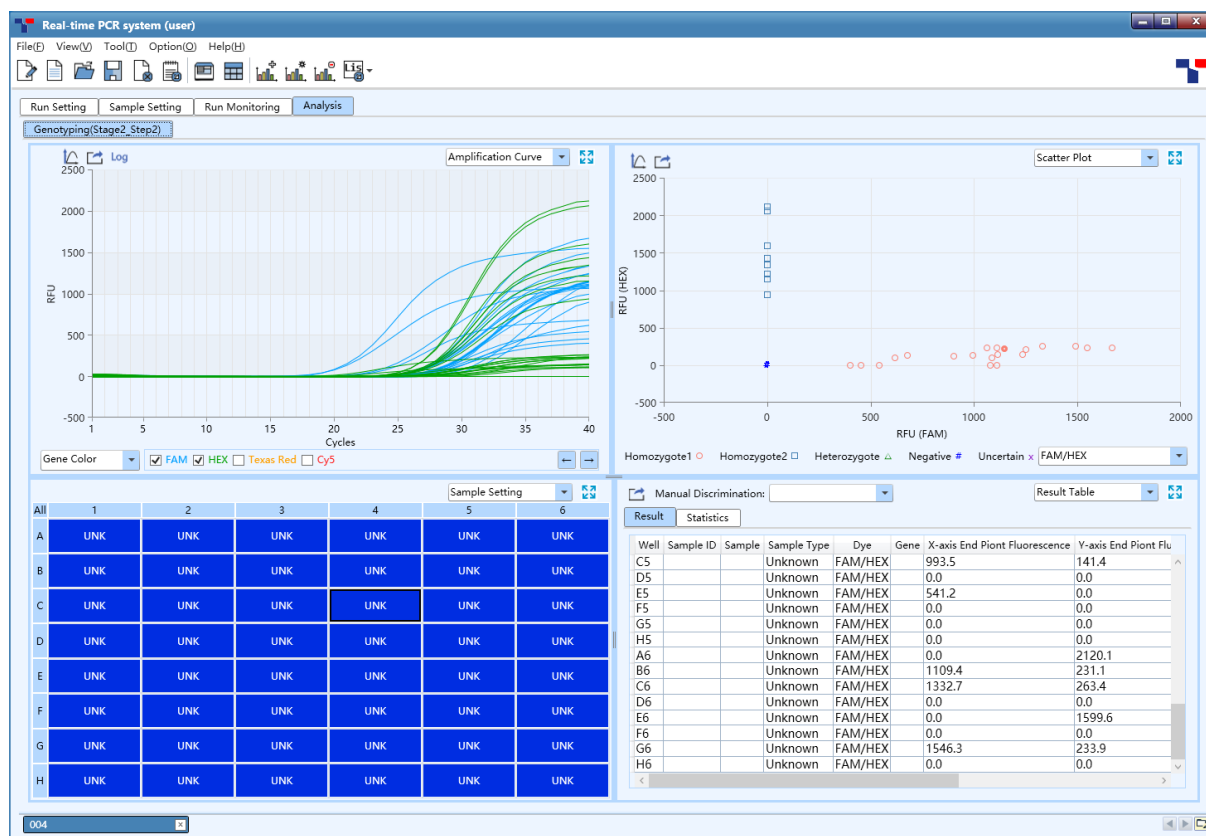


Figure D-16 Genotyping analysis interface

Interface Introduction and Parameter Description

- I. **Scatter Plot:** display the end point fluorescence distribution of two dyes, as shown in figure 16.1.

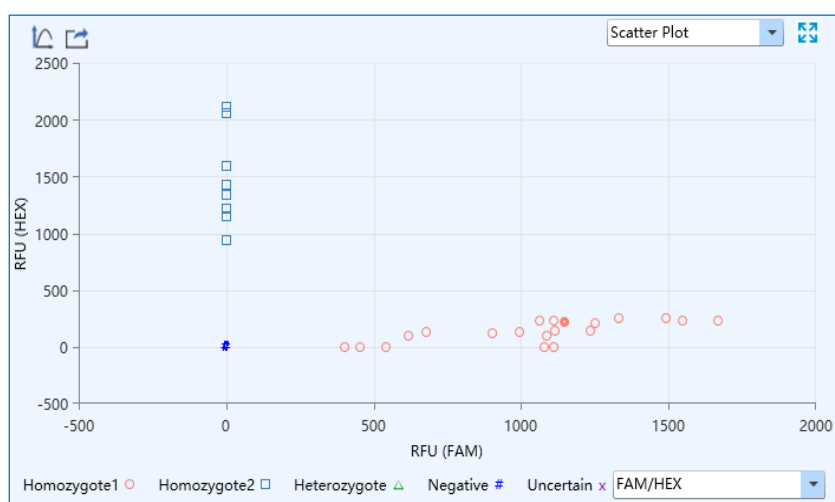




Figure D-16.1 Genotyping analysis interface - Scatter Plot functional module

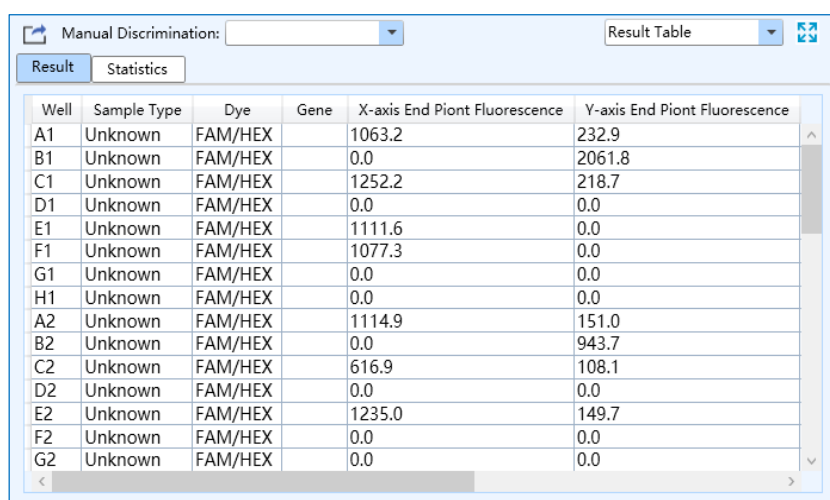
► **Parameter introduction:**

- X axis and Y axis represent the Relative Fluorescence Unit (RFU) of different dyes respectively;
- Each point represents a sample; different icons represent different genotypes or interpretation results.

► **Function key introduction:**

- The introduction of  **Y-axis Coordinate Adjustment** > and  **Export** > keys are same with those keys introduced in [D-3.3.4.1 Abs Quant - Amplification Curve](#).

II. **Result Table:** display the sample details of and result data of the current experiment, as shown in figure 16.2.




Well	Sample Type	Dye	Gene	X-axis End Point Fluorescence	Y-axis End Point Fluorescence
A1	Unknown	FAM/HEX		1063.2	232.9
B1	Unknown	FAM/HEX		0.0	2061.8
C1	Unknown	FAM/HEX		1252.2	218.7
D1	Unknown	FAM/HEX		0.0	0.0
E1	Unknown	FAM/HEX		1111.6	0.0
F1	Unknown	FAM/HEX		1077.3	0.0
G1	Unknown	FAM/HEX		0.0	0.0
H1	Unknown	FAM/HEX		0.0	0.0
A2	Unknown	FAM/HEX		1114.9	151.0
B2	Unknown	FAM/HEX		0.0	943.7
C2	Unknown	FAM/HEX		616.9	108.1
D2	Unknown	FAM/HEX		0.0	0.0
E2	Unknown	FAM/HEX		1235.0	149.7
F2	Unknown	FAM/HEX		0.0	0.0
G2	Unknown	FAM/HEX		0.0	0.0

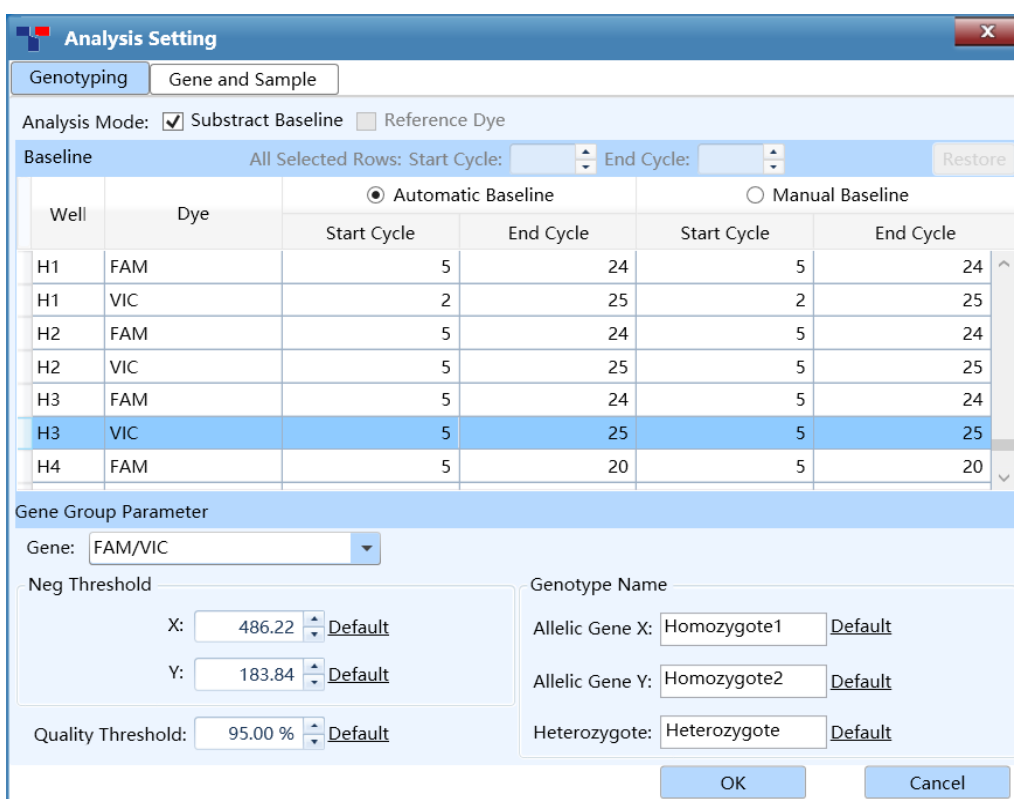
Figure D-16.2 Genotyping analysis interface - Result Table functional module

- User can double click the title of each column to sort all sample results according to the content in this column.
- User can click the title of each column and drag the column to the left or right to adjust the sequence of sample results.
- By default, the application software will automatically discriminate the different genotypes. But user can also select any sample in the **Result Table** list and set the genotype for the selected sample from the **Manual Discrimination** drop-down list.

Genotyping Analysis Setting

User can click the <  **Analysis Setting** > icon in the Tool Bar and the application software will automatically pop up the genotyping analysis setting window, which consists of **Dye and Gene**, **Sample** and **Genotyping** three tabs. User could set the relevant parameters for the current analytical method in the genotyping analysis setting window.

- I. **Genotyping** tab: user can set **Analysis Mode**, **Reference Dye**, **Baseline** and **Gene Group Parameter** in **Genotyping** tab, as shown in figure D-17.



Analysis Setting

Genotyping | Gene and Sample

Analysis Mode: ☒ Subtract Baseline ☐ Reference Dye

Baseline: All Selected Rows: Start Cycle: End Cycle: Restore

Well	Dye	Automatic Baseline		Manual Baseline	
		Start Cycle	End Cycle	Start Cycle	End Cycle
H1	FAM	5	24	5	24
H1	VIC	2	25	2	25
H2	FAM	5	24	5	24
H2	VIC	5	25	5	25
H3	FAM	5	24	5	24
H3	VIC	5	25	5	25
H4	FAM	5	20	5	20

Gene Group Parameter

Gene: FAM/VIC

Neg Threshold

X: 486.22 Default

Y: 183.84 Default

Quality Threshold: 95.00 % Default

Genotype Name



Allelic Gene X: Homozygote1 Default



Allelic Gene Y: Homozygote2 Default

Heterozygote: Heterozygote Default

OK Cancel

Figure D-17. Genotyping Analysis Setting interface - Genotyping tab

- **Analysis Mode:** same with the introduction of **D-3.3.4.1 Abs Quant - Abs Quant Analysis Setting - Amplification Curve tab**.
- **Gene Group Parameter:**
 - a. **Gene:** display the detective dyes of different genotypes;
 - b. **Neg Threshold:** user can use  and  keys to set the corresponding negative threshold for the X axis and Y axis of scatter plot.

c. **Quality Threshold:** user can use  and  keys to set the quality threshold of application software algorithm. Increase the quality threshold value can improve the credibility of grouping result.

d. **Genotype Name:** user can change the grouping name of allelic gene and heterozygote gene.

II. **Gene and Sample** tab: the introduction of relevant parameters and function keys is same with those introduced in **D-3.3.4.1 Abs Quant - Abs Quant Analysis Setting**.

3.3.4.6 End Point Fluorescence

The end point fluorescence analysis displays the final detection result based on the fluorescence intensities measured at the end point of amplification plateau phase.

The end point fluorescence analysis interface consists of seven functional modules: **Amplification Curve**, **End Point Fluorescence Scatter Plot**, **Sample Setting**, **Result Table**, **Raw Curve**, **Raw Fluorescence** and **Heat Map**. The end point fluorescence analysis interface is divided into four areas, the application software displays four functional modules (**Amplification Curve**, **End Point Fluorescence Scatter Plot**, **Sample Setting**, **Result Table**) by default, as shown in figure D-18.

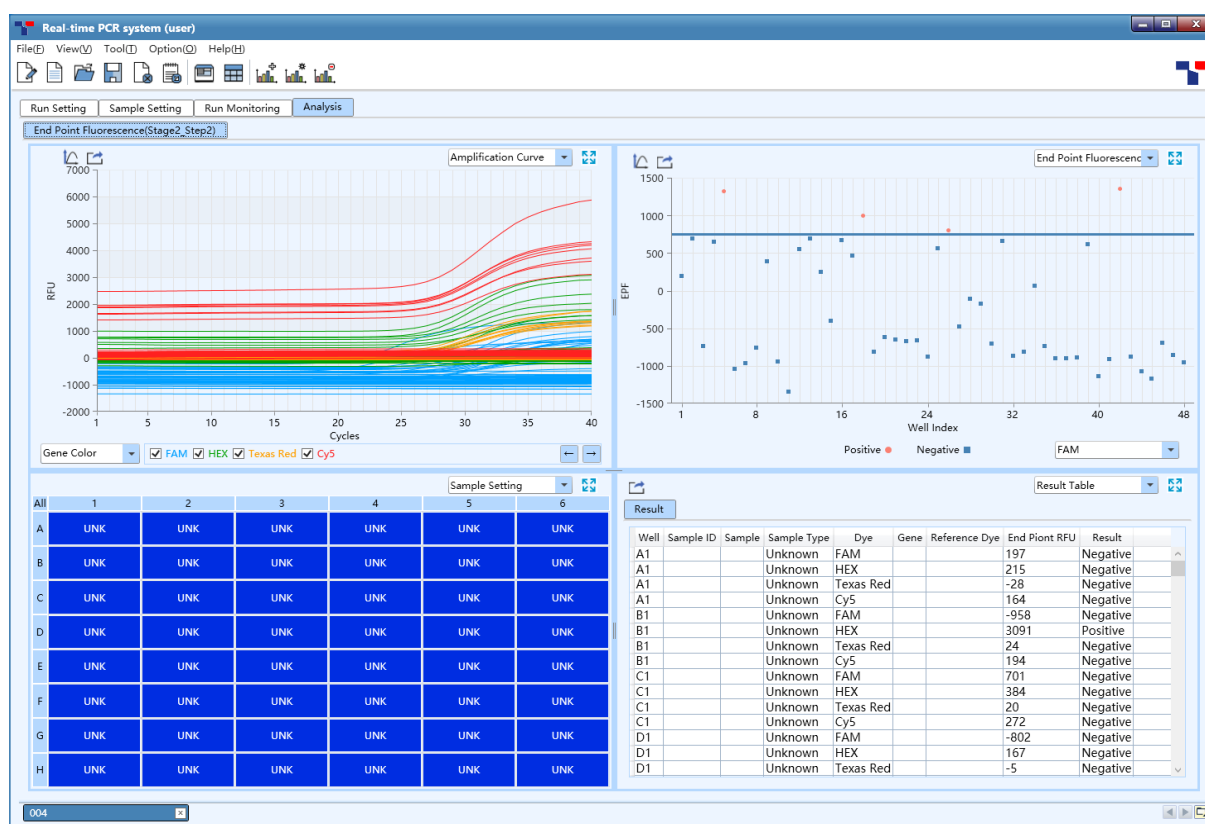


Figure D-18. End Point Fluorescence analysis interface

Reminding: the parameter descriptions of *Amplification Curve*, *Sample Setting*, *Result Table*, *Raw Curve*, *Raw Fluorescence* and *Heat Map* functional modules are same with those introduced in [D-3.3.4.1 Abs Quant](#).

Interface Introduction and Parameter Description

- I. **End Point Fluorescence Scatter Plot:** display the scatter plot of fluorescence intensities measured at end the point of amplification plateau phase, as shown in figure 18.1.

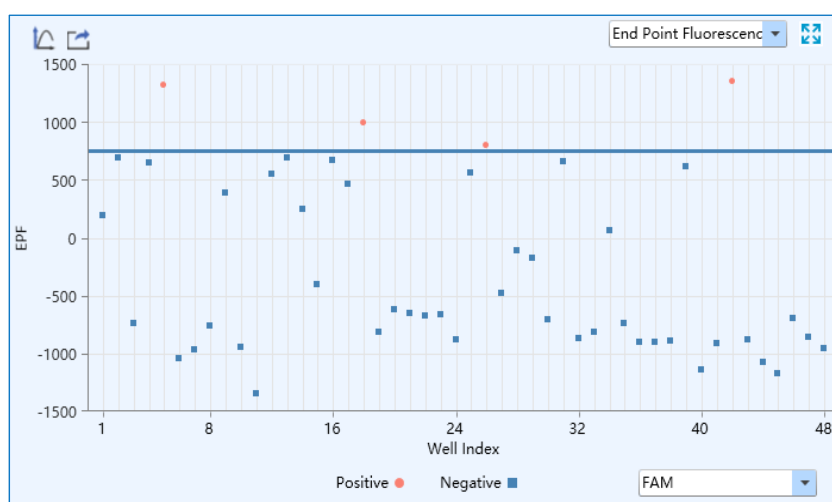




Figure D-18.1 End Point Fluorescence analysis interface - End Point Fluorescence Scatter Plot


► Parameter introduction:

- X axis represents the sample well index of the current experiment samples;
- Y axis represents the End Point Fluorescence (EPF) intensity;
- Each point represents a sample; different icons represent the negative or positive interpretation results.

► Function key introduction:

- The introduction of <  **Y-axis Coordinate Adjustment** > and <  **Export** > keys are same with those keys introduced in [D-3.3.4.1 Abs Quant - Amplification Curve](#).

End Point Fluorescence Analysis Setting

User can click the  **Analysis Setting** > icon in Tool Bar and the application software will automatically pop up the end point fluorescence analysis setting window, which consists of **End Point Fluorescence** and **Gene and Sample** two tabs. User can set the relevant parameters for the current analytical method in the end point fluorescence analysis setting window.

I. **End Point Fluorescence** tab: as shown in figure D-19.

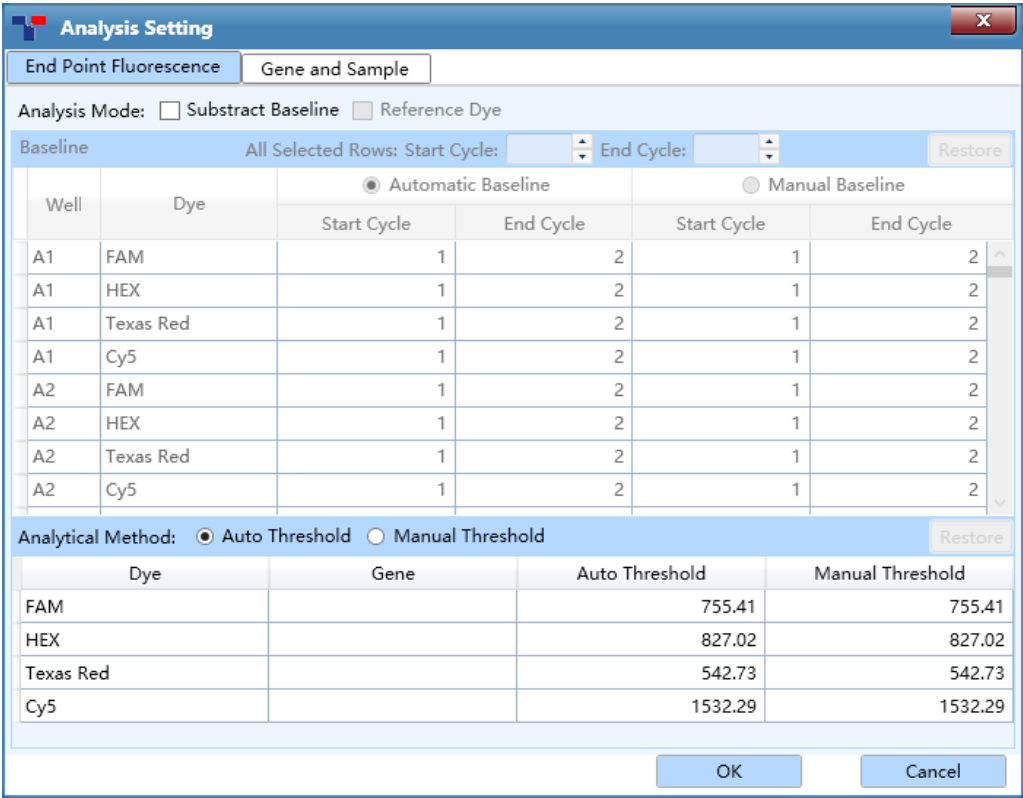


Figure D-19. End Point Fluorescence Analysis Setting interface - End Point Fluorescence tab

- **Analysis Mode:** same with the introduction of [D-3.3.4.5 Genotyping - Genotyping Analysis Setting - Genotyping tab](#).
 - **Analytical Method:** same with the introduction of [D-3.3.4.5 Genotyping - Genotyping Analysis Setting - Genotyping tab](#).
- II. **Gene and Sample** tab: the introduction of relevant parameters and function keys is same with those introduced in [D-3.3.4.1 Abs Quant - Abs Quant Analysis Setting](#).

E. Instrument Software Operation

The Gentier instrument can run independently from the control computer. User can use the touch screen or external mouse to operate the instrument system software, so as to realize the experiment program setting and other relevant operations.

1. Boot Screen & Main Menu Interface

Switch on the power switch of the Gentier instrument. The instrument system will automatically conduct self-inspection, the touch screen will light up and display instrument software boot screen.

After the self-inspection, the instrument software will enter the main interface, which is consisted of status bar, operation area and main function keys, as shown in figure E-1.

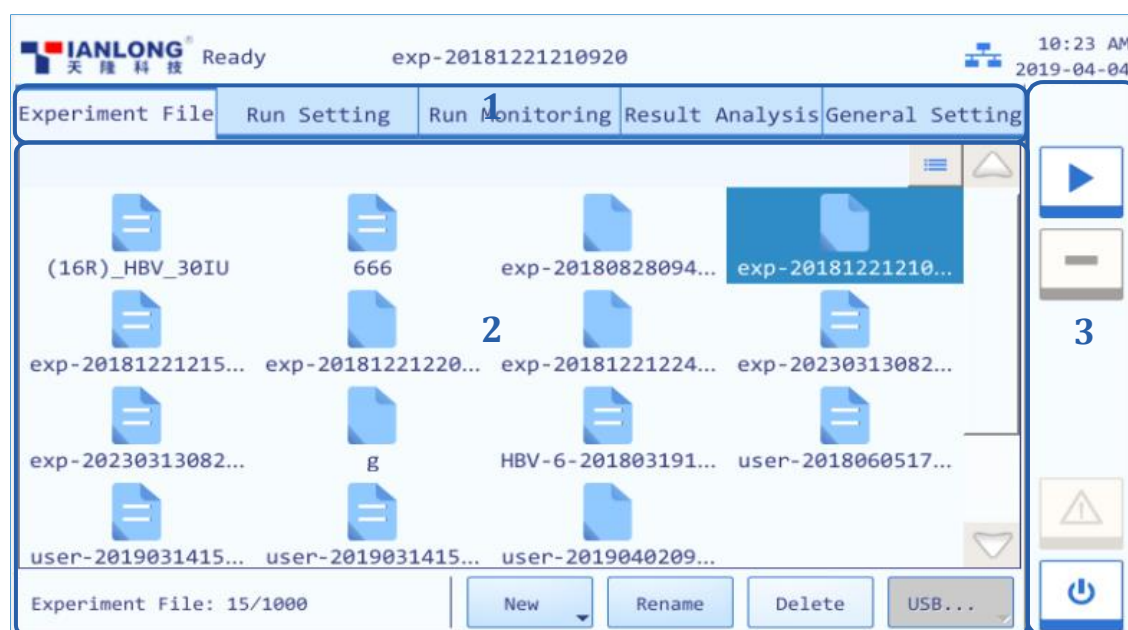



















Figure E-1. Instrument software main interface

Interface Introduction & Parameter Descriptions

1. **Status Bar:** display the system status, current file name, instrument status and system time.
 - ▶ **System Status:** display the current instrument system status.
 - **Initializing:** the instrument system is in the process of initialization;

- **Ready:** the instrument system initialization is completed, ready for running;
 - **Running:** the experiment is running;
 - **Pause:** the current running experiment is paused;
 - **Error:** instrument hardware or software error, cannot execute any operation.
- **File Name:** display the current experiment file name and its file path.
- **Instrument Status:** display the current instrument status.
- <  **Not Connected** >: the instrument is not connected to the network;
 - <  **Connected** >: the instrument is connected to the network;
 - <  **USB** >: the instrument is connected with USB device;
 - <  **Hot Lid Used** >: the hot lid heating function is enabled;
 - <  **Hot Lid Unused** >: the hot lid heating function is not enabled;
 - <  **Top Lid Open** >: the top lid is open;
 - <  **Error** >: the instrument hardware or software error has occurred, cannot execute any operation.
- **System Time:** display the current system date and time.
2. **Operation Area:** includes **Experiment File**, **Run Setting**, **Run Monitoring**, **General Setting** four tabs, for more operation details, please refer to [E-2 Instrument Software Operation Area](#).
3. **Main Function Keys:**
- <  **Run Experiment** >: start running the current experiment. This function key is activated when the instrument system is under **Ready** or **Pause** status.
- ❗ **Reminding:** when the instrument system is under **Running** status, the <  **Run Experiment** > function key will be converted to <  **Pause Experiment** > function key, user can press this function key to pause the current running experiment.
- <  **Stop Experiment** >: stop the current running experiment. This function key is activated when the instrument system is under **Running** status.

- ▶ <  **Shutdown / Restart** >: shutdown or restart the instrument. This function key is inactivated when the instrument system is under **Running** status.
- ▶ <  /  /  **Warning** >: this function key is inactivated (grey icon) under normal system status, however it will turn yellow or red when a certain system error happened, press this key, the system will display the warning information.
 -  Yellow icon: the instrument can still execute the current operation;
 -  Red icon: the instrument cannot execute the current operation;

2. Instrument Software Operation Area

The operation area of the system software includes **Experiment File**, **Run Setting**, **Run Monitoring**, **General Setting** four tabs. User could perform relevant operations and set the corresponding contents according to the specific experimental requirements.

- ▶ **Experiment File**: display experiment files and provide the relevant experiment file operations;
- ▶ **Run Setting**: provide the experiment temperature and fluorescence setting operations;
- ▶ **Run Monitoring**: provide the real-time running state of the experiment and display the real-time fluorescence and temperature data of the current running experiment.
- ▶ **General Setting**: provide general setting operations of the current instrument system.

2.1 Experiment File

The main interface of instrument software displays the experiment file tab by default. This tab is consisted of three parts: experiment file display area, experiment file information window and experiment file action bar, as shown in figure E-2.

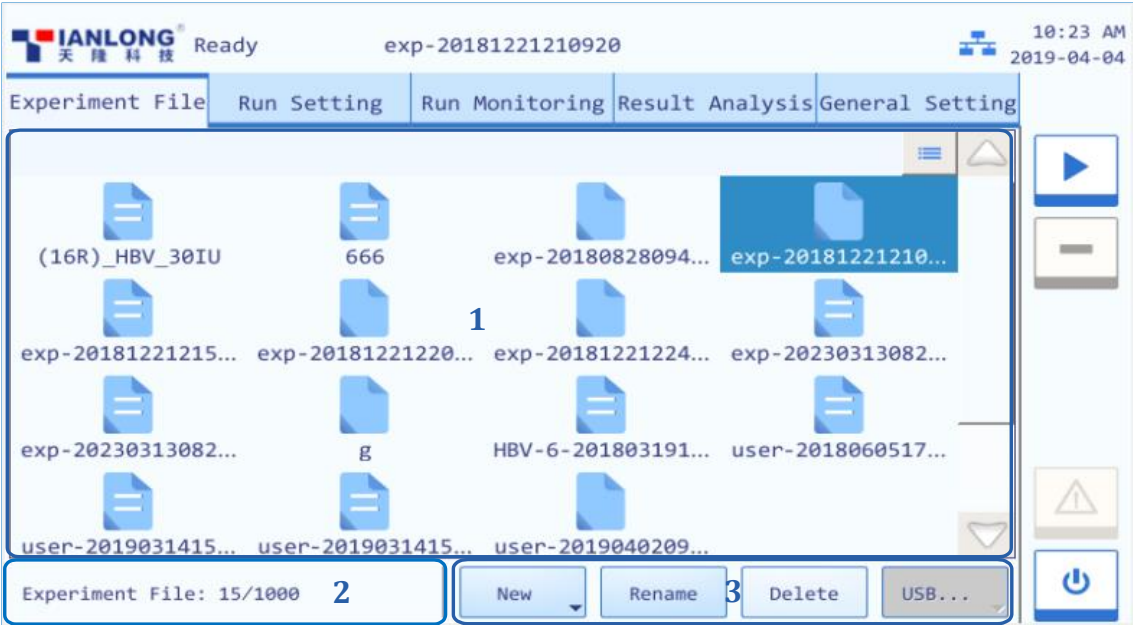


Figure E-2. Instrument software – Experiment File interface

Interface Introduction & Parameter Descriptions

- 1. **Experiment File Displaying Area:** display the pre-existed experiment files or file folders within the instrument system, as shown in figure E-2.1.

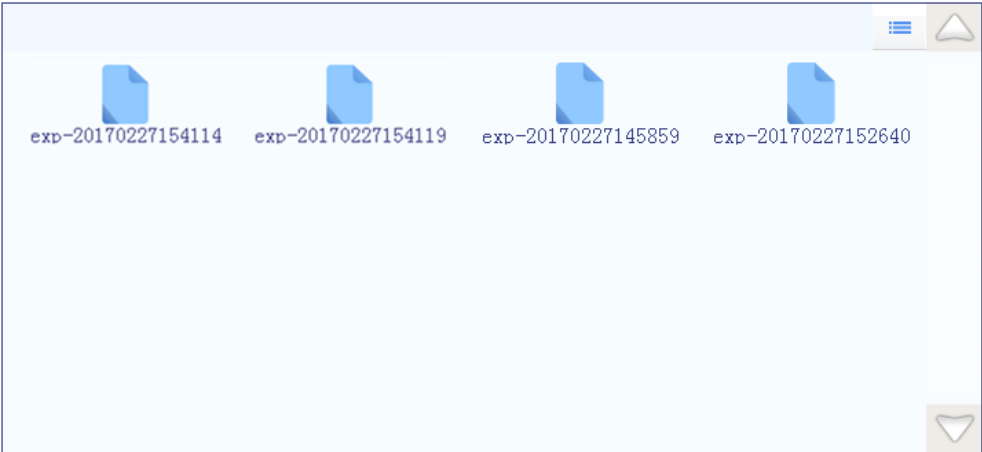






Figure E-2.1 Experiment File tab - Experiment file display area

- User can press the icon of a certain experiment file or file folder in the experiment file display area to open it.

- ▶ User can slide the slider or press  and  icons to view all the experiment files or file folders in the experiment file display area.
 - ▶ User can press the  icon on the top left corner of the experiment file display area and it will turn into  icon, and the display mode of the experiment file displaying area will change from icons to file details list.
2. **Experiment File Information Window:** display the pre-existed experiment quantity within the instrument system, as shown in figure E-2.2.




Experiment File: 2/1000

Figure E-2.2 Experiment File tab - Experiment file information window

3. **Experimental file action bar:** consists of < New >, < Rename >, < Delete > and < USB... > four keys, as shown in figure E-2.3.



Figure E-2.3 Experiment File tab - Experimental file action bar

- ▶ < New >: create a new experiment file or file folder, this key is inactivated when the instrument system is under **Running** status. Press the  icon, there are three options in the drop-down list.
 - **New Folder:** create a new experiment file folder.
 - **New Experiment:** create a new experiment file.
 - **New Experiment from Selected Experiment:** user could press and select a pre-existed experiment file in the experiment file displaying area, and create a new experiment based on the selected experiment settings.
- ❗ **Reminding:** the system will pop up a keyboard for user to name the new experiment file or file folder, as shown in figure E-2.4. User can press < Close > to exit the keyboard.
- ❗ **Reminding:** the instrument software will name the new experiment file or file folder with creation date and time by default.

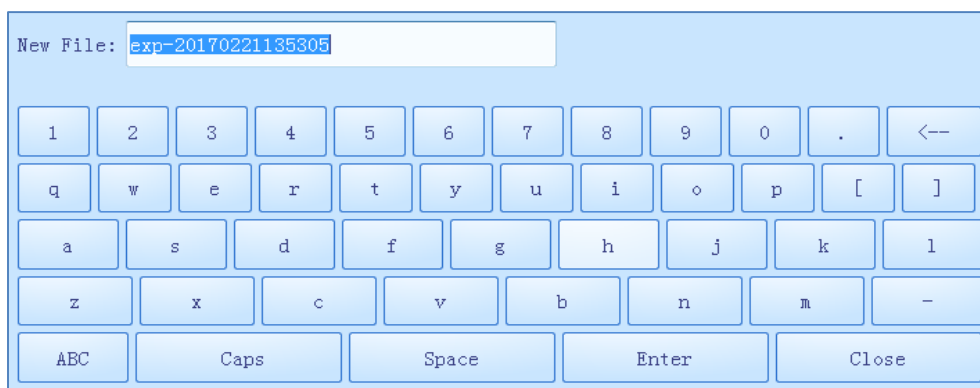






Figure E-2.4 Experiment File tab - keyboard

- ▶ **< Rename >**: rename the selected experiment file or file folder, this key is inactivated when the instrument system is under **Running** status.
 - ▶ **< Delete >**: delete the selected experiment file or file folder, this key is inactivated when the instrument system is under **Running** status.
 - ▶ **< USB... >**: execute file transmission between the instrument and USB device. Press the  icon, there are two options in the drop-down list.
 - **Export Experiment**: export the selected experiment file or file folder from the instrument system to the USB device.
 - **Import Experiment**: import the selected experiment file or file folder from the USB device to the instrument system.
-  **Reminding**: If there are more than one USB devices connected to the current instrument, after selecting the **Export Experiment** or the **Import Experiment** option, the instrument software will automatically pop up USB device list for user to choose the proper USB device.

2.2 Run Setting

The run setting tab consists of **Temperature Setting** and **Fluorescence Setting** two sub-tabs, as shown in figure E-3.

-  **Reminding**: the system will automatically enter the run setting interface after new experiment file creation;
-  **Reminding**: user could also select an experiment file, press **Run Setting** tab to enter the run setting interface to view or edit the experimental settings.

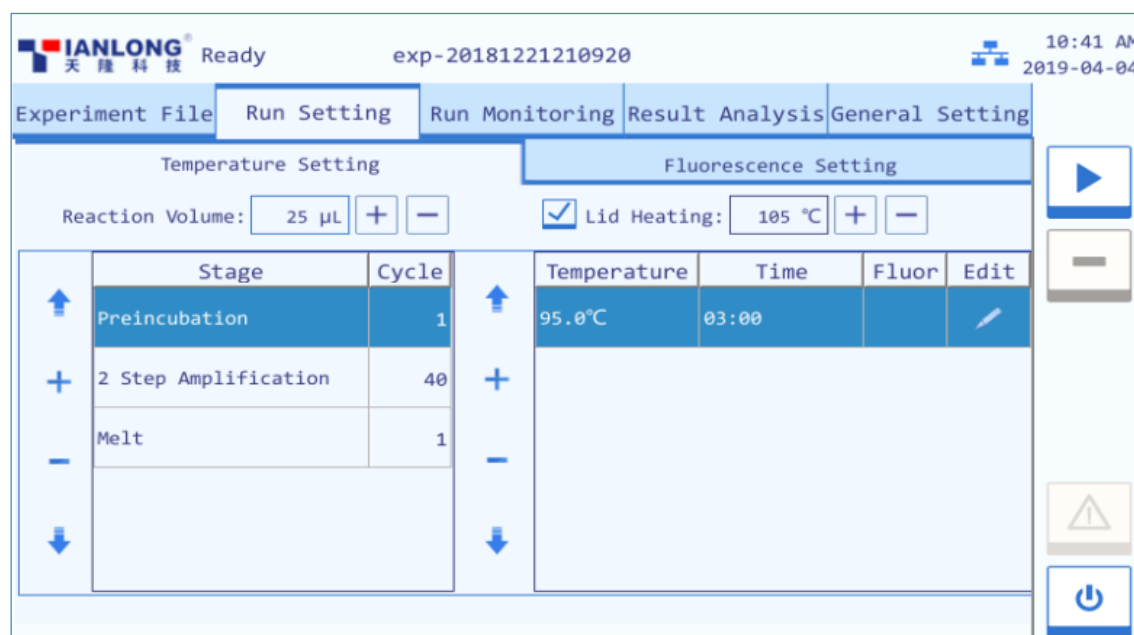



Figure E-3. Run Setting interface - Temperature Setting sub-tab

2.2.1 Temperature Setting



The **Temperature Setting** sub-tab is selected on the run setting interface by default, as shown in figure E-3. User can edit the relevant experiment settings and temperature program in the **Temperature Setting** sub-tab.

Interface Introduction & Parameter Descriptions

1. **Tube Type:** the instrument software provides three tube type options for user, including **Clear**, **White** and **Frosted**. User can press  icon to select.

2. **Reaction Volume:** user can press  or  icons to set the reaction volume.





 **Reminding:** the reaction volume setting range is 0µL~100µL.

3. **Hot Lid:** user could press the check box of **Hot Lid** to decide whether to utilize the hot lid heating function, and press  or  icons to edit the hot lid temperature.






 **Reminding:** the hot lid temperature setting range is 40°C ~110°C.

4. **Stage Setting Box:**

- **Stage:** display the stage type of the temperature program.
- **Cycle:** display the cycle number of the current stage.

- <  **Up** >: move up the selected stage.
- <  **Add** >: add a new stage.
- <  **Delete** >: delete the selected stage.
- <  **Down** >: move down the selected stage.

5. Step Setting Box:

- **Temperature**: display the target temperature of the current step.
- **Time**: display the temperature holding time of the current step.
- **Fluor**: display whether to read the fluorescence at the current temperature step;
- **Edit**: press  icon in **Edit** column to edit the corresponding step.
- <  **Up** >: move up the selected step.
- <  **Add** >: add a new step.
- <  **Delete** >: delete the selected step.
- <  **Down** >: move down the selected step.

Temperature Program Setting Operation

Step 1: Add stage


User can press <  **Add** > in the stage setting box and the instrument software will automatically pop up a stage type selection window, as shown in figure E-3.1



Figure E-3.1Temperature Setting sub-tab - Stage type selection window

- ▶ Seven predefined stage types are provided by the instrument software for user: **Preincubation, Reverse Transcription, 2 Step Amplification, 3 Step Amplification, Melting, Continuous Melting** and **Cooling**. User can also press **Custom Stage** to define the stage setting according to specific experiment requirements.
- ▶ User can press < **Add** > to add the selected stage to stage setting box.
- ▶ User can press < **Back** > to return back to the run setting interface.
- ❗ **Reminding:** at least one stage should be included in the temperature program.

Step 2: Edit stage

- ▶ The added stage will be automatically displayed in the stage setting box and user can select any stage to delete or change its sequence.
- ▶ User can select a certain stage and press its cycle number in the corresponding **Cycle** column, the cycle edit box will pop up as shown in figure E-3.2.

Figure E-3.2 Cycle edit box

- User could set the cycle number in the cycle edit box, press < **OK** > to confirm the cycle setting or press < **Cancel** > to cancel the setting.


❗ **Reminding:** the cycle setting range is 1~99.

Step 3: Add/delete a stage

- ▶ Select a stage in the stage setting box and the step setting box on the right will display the corresponding steps of the selected stage;
- ▶ User can add a new step or select a step to delete or change its order;

❗ **Reminding:** at least one step should be included in one stage.


Step 4: Set fluorescence reading step

- In the step setting box, user can press the **Fluor** coloum of a cetain step to set whether to read its fluorescence, the  icon will be displayed in the corresponding **Fluor** coloum of the fluorescence reading step.



Reminding: user can set only one fluorescence reading step in one stage.

Step 5: Edit step


User can select a certain step in the step setting box, and press the  icon in the corresponding **Edit** column, the instrument software will automatically pop up the step edit window, as shown in figure E-3.3.



The image shows a 'Step Setting' window with the following fields and options:

- Temperature:** 58.0 °C
- Time:** 00:30
- Ramp:** 6.1 °C/s
- Step Mode:**
 - ☒ Standard
 - ☐ Touchdown
 - ☐ Long
 - ☐ Gradient
- Back** button

Figure E-3.3 Temperature Setting sub-tab - Step Edit window - Standard step mode

Four **Step Mode** options are provided by the instrument software: **Standard**, **Touchdown**, **Gradient** and **Long**. User can select a certain step mode option by pressing the  icon in front of it, and the icon will turn blue.

1) The **Standard** step mode option is selected by default.

- User can press the input boxes of **Temperature**, **Time** and **Ramp** to edit the corresponding parameters for the current standard step, as shown in figure E-3.3a.
- **Temperature:** the temperature in °C, which is to be held for a defined time.
 - **Time:** the time for which the temperature is to be held.
 - **Ramp:** the rate of temperature change in °C per second.

Figure E-3.3a shows the Step Edit window for a standard step. It consists of three main sections: Temperature [°C], Time, and Ramp [°C/s]. Each section has a numeric keypad with digits 0-9, a decimal point, and a 'Cancel' button. The Temperature section is set to 94.0, the Time section is set to 00:10, and the Ramp section is set to 8.0. Each section also has an 'Ok' button.

Figure E-3.3a. Step Edit window - Standard step parameter setting

- ❗ **Reminding:** the temperature setting range is 0.0°C ~100.0°C.
- ❗ **Reminding:** the time setting range is 1s~60min.
- ❗ **Reminding:** the ramp setting range is 0.1°C/s~8°C/s.

2) If the **Touchdown** step mode option is selected, the step edit window will be shown as figure E-3.4.

Figure E-3.4 shows the Step Edit window for a Touchdown step. It includes fields for Initial Temp. (58.0 °C), Target Temp. (50.0 °C), Time (00:30), Delta Temp. (1.0 °C/Cycle), Ramp (6.1 °C/s), and Start Cycle (1). At the bottom, the Step Mode is set to Touchdown (selected), with options for Standard, Long, and Gradient. A 'Back' button is also present.

Figure E-3.4 Temperature Setting sub-tab - Step Edit window - Touchdown step mode

- The touchdown step mode allows the temperature program to change the annealing step temperature from the initial temperature to the target temperature as the cycling proceeds.
- User can press the input boxes of **Initial Temp.**, **Target Temp.**, **Delta Temp.** and **Start Cycle** to edit the corresponding parameters for the current touchdown step, as shown in E-3.4a.


The figure displays four separate windows for editing PCR step parameters in Touchdown mode. Each window features a title bar, a parameter label with units and a range, a numeric input field, a keypad with digits 0-9 and a decimal point, and 'Ok' and 'Cancel' buttons.


- Initial Temp. [°C] {35.0-100.0}**: The value 72.0 is entered.
- Temperature [°C] {35.0-100.0}**: The value 50.0 is entered.
- Delta Temp. [°C/Cycle] {0.1-5.0}**: The value 1.0 is entered.
- Start Cycle {1-40}**: The value 1 is entered.

Figure E-3.4a Step Edit window - Touchdown step parameter setting

- **Initial Temp.:** the initial value of annealing temperature change range.
- **Target Temp.:** the target value of annealing temperature change range.
- **Delta Temp.:** the temperature change (°C) per cycle.
- **Start Cycle:** the cycle number after which the temperature change is started.

 **Reminding:** the initial and target temperature setting range is 35.0°C ~100.0°C.

 **Reminding:** the delta temperature setting range is 0.1°C ~5.0°C.

 **Reminding:** the star cycle range is 1~ max cycle number of the current stage.

 **Reminding:** the other parameter settings are same with the standard step mode.

3) If the **Long** step mode option is selected, the step edit window will be shown as E-3.5.

Step Setting				
Temperature:	58.0	°C	Target Time:	02:00
Initial Time:	00:30		Delta Time:	00:05 /Cycle
Ramp:	6.1	°C/s	Start Cycle:	1
Step Mode: <input type="radio"/> Standard <input type="radio"/> Touchdown <input checked="" type="radio"/> Long <input type="radio"/> Gradient <button>Back</button>				

Figure E-3.5 Step Edit window - Long step mode

- The long step mode allows the temperature program to change the elongation step temperature holding time from the initial time to the target time as the cycling proceeds.
- User can press the input boxes of **Initial Time**, **Target Time**, **Delta Time** and **Start Cycle** to edit the corresponding parameters for the current long step, as shown in figure E-3.5a.

Initial Time {00:01-60:00} 00 : 30 <-- 1 2 3 4 5 6 7 8 9 0 Ok Cancel	Target Time {00:01-60:00} 02 : 00 <-- 1 2 3 4 5 6 7 8 9 0 Ok Cancel
Delta Time [/Cycle] {00:01-10:00} 00 : 05 <-- 1 2 3 4 5 6 7 8 9 0 Ok Cancel	Start Cycle {1-40} 1 <-- 1 2 3 4 5 6 7 8 9 0 Ok Cancel

Figure E-3.5a Step Edit window - Long step parameter setting

- **Initial Time:** the initial value of the elongation time change range.
- **Target Time:** the target value of the elongation time change range.
- **Delta Time:** the temperature change(°C) per cycle.
- **Start Cycle:** the cycle number after which the time change is started.
- ❗ **Reminding:** the initial and target time setting range is 1s~60min.
- ❗ **Reminding:** the delta time setting range is 1s~10min.
- ❗ **Reminding:** the star cycle range is 1~ max cycle number of the current stage.
- ❗ **Reminding:** the other parameter settings are same with the standard step mode.

4) If the **Gradient** step mode option is selected, the step edit window will be shown as figure E-3.6.

The screenshot shows the 'Step Setting' window. It has a title bar 'Step Setting'. Inside, there are four input fields: 'Time:' with value '00:30', 'Ramp:' with value '6.1 °C/s', 'Temp. Center:' with value '58.0 °C', and 'Temp. Offset:' with value '5.0 °C'. Below these fields is a 'Details' button. At the bottom, there is a 'Step Mode:' section with four radio buttons: 'Standard', 'Touchdown', 'Long', and 'Gradient'. The 'Gradient' radio button is selected. To the right of the radio buttons is a 'Back' button.

Figure E-3.6 Temperature Setting sub-tab - Step Edit window - Gradient step mode

- The gradient step mode allows the Gentier sample to adopt different temperatures. After setting the temperature center value and the temperature offset value for the current gradient step, the system will automatically calculate the gradient temperatures.
- User can press the input boxes of **Temp. Center** and **Temp. Offset** to edit the corresponding parameters for the current gradient step. User could also and press < **Details** > to view the specific gradient temperatures of each row, as shown in figure E-3.6a.

Figure E-3.6a Step Edit window - Gradient step parameter setting

- **Temp. Center:** the center value of the temperature grading range.

- **Temp. Offset:** the offset value of the temperature grading range.

❗ **Reminding:** the temperature center setting range is 35.5°C ~99.5°C.

❗ **Reminding:** the temperature offset setting range is 0.5°C ~20.0°C.

❗ **Reminding:** the other parameter settings are same with the standard step mode.

❗ **Reminding:** the Gentier 48S instrument model cannot support gradient PCR step mode.

5) If the current step is the last step of a melting stage, the step setting window of this step is as shown in figure E-3.7.

Figure E-3.7 Step Edit window - Melting step mode

► The melting stage allows the instrument system to read fluorescence signals after each temperature increment. User can press the input boxes of **Increment** to edit the temperature increment for the current melting step, as shown in figure E-3.7a.

Increment [°C] {0.1-5.0}

0.5 <--

1	2	3
4	5	6
7	8	9
0	.	

Ok Cancel

Figure E-3.7a Step Edit window - Melting step parameter setting

- **Increment:** the temperature change(°C) after which the system will read the fluorescence.

 **Reminding:** the temperature increment setting range is 0.1°C - 5.0°C.

- 6) If the current step is the last step of a continuous melting stage, the step setting window of this step is as shown in figure E-3.8.

Step Setting

Temperature: 97.0 °C Readings: 10 Readings/°C

Back

Figure E-3.8 Step Edit window - Continuous Melting step mode

- Continuous melting stage allows the instrument system to read fluorescence more frequently, user can press the input boxes of **Readings** to set the fluorescence reading times per °C for the current continuous melting step, as shown in figure E-3.8a.

- **Readings:** the fluorescence reading times per °C.

 **Reminding:** the reading frequency setting range is 2readings/°C - 15readings/°C.

Figure E-3.8a Step Edit window - Continuous Melting step parameter setting


2.2.2 Fluorescence Setting

The **Fluorescence Setting** sub-tab on the run setting interface is as shown in figure E-3.9.


Channel	Dye	Excitation/Emission
<input checked="" type="checkbox"/> 1	FAM	465/510
<input checked="" type="checkbox"/> 2	HEX	527/563
<input checked="" type="checkbox"/> 3	Texas Red	580/616
<input checked="" type="checkbox"/> 4	Cy5	632/664

Figure E-3.9. Run Setting Interface - Fluorescence Setting sub-tab

- User can set the fluorescence channel and dyes for the current experiment in the **Fluorescence Setting** sub-tab.
- User can check the corresponding check box in the **Channel** column to set the fluorescence channel for the current experiment (1-4 channels for Gentier 48E/48S and 1-2 channels for Gentier 48R);

- User can press the corresponding  icon in the **Dye** column and select the proper dye from the drop-down list;
- User can view the excitation and emission wave length of the corresponding channel in the **Excitation/Emission** column;

2.3 Run Monitoring

After finishing the experiment setting, user can press the <  **Run Experiment** > function key to start running the current experiment, the system will enter the run monitoring interface, which is consisted of three parts: run status bar, run monitoring option bar and run monitoring diagram, as shown in figure E-4.

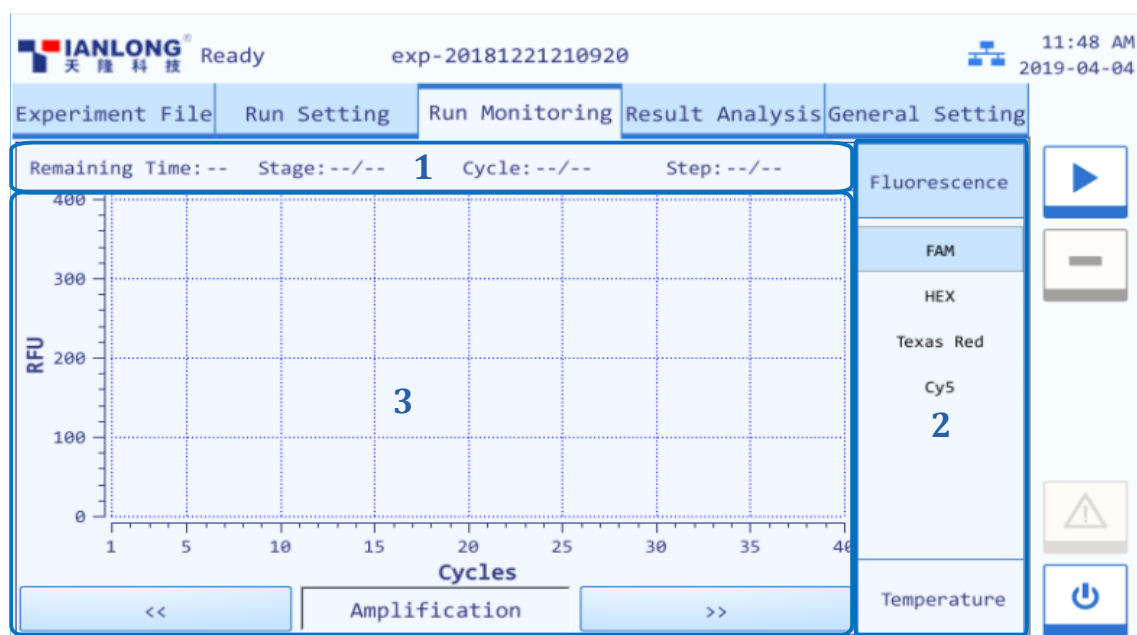




Figure E-4. System software – Run Monitoring interface

1. **Run Status Bar:** displays the real-time running status of the the current experiment, includes **Remaining Time**, **Stage**, **Cycle** and **Step**.
2. **Run Monitoring Option Bar:** user can select **Fluorescence** or **Temperature** option in the run monitoring option bar to monitor the corresponding content of the current experiment.

- ▶ User can press the **Fluorescence** option to monitor the real-time amplification curve, melting curve and fluorescence heat map of the current experiment.
- ▶ User can press the **Temperature** option to monitor real-time temperature program of the current experiment

3. Run Monitoring Diagram:

- ▶ If user select the **Fluorescence** option in the run monitoring option bar, user can press  or  keys below the run monitoring diagram to choose the monitor content, which can be the real-time **Amplification** curve, **Melting** curve or fluorescence **Heat Map** of the current experiment, as shown in figure E-4.1.
- a. The run monitoring diagram will display the real-time **Amplification** curve of the current experiment by default, as shown in figure E-4.1.

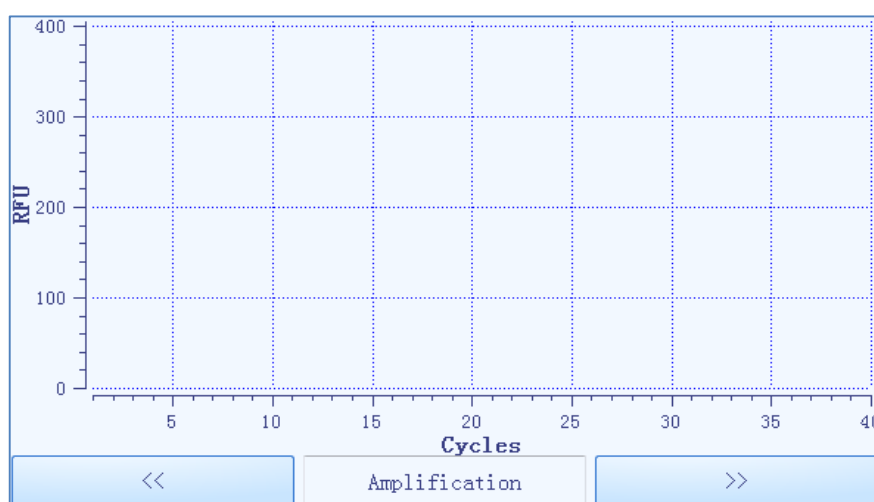



Figure E-4.1 Run Monitoring Diagram - Real-time Amplification Curve

- X axis represents the cycle number;
- Y axis represents the Relative Fluorescence Unit (RFU);
- User can press any dye under the **Fluorescence** option in the run monitoring option bar, and the run monitoring diagram will only display the corresponding real-time amplification curve;
- b. User can press  key below the run monitoring diagram to monitor the real-time **Melting** curve of the current experiment, as shown in figure E-4.2.

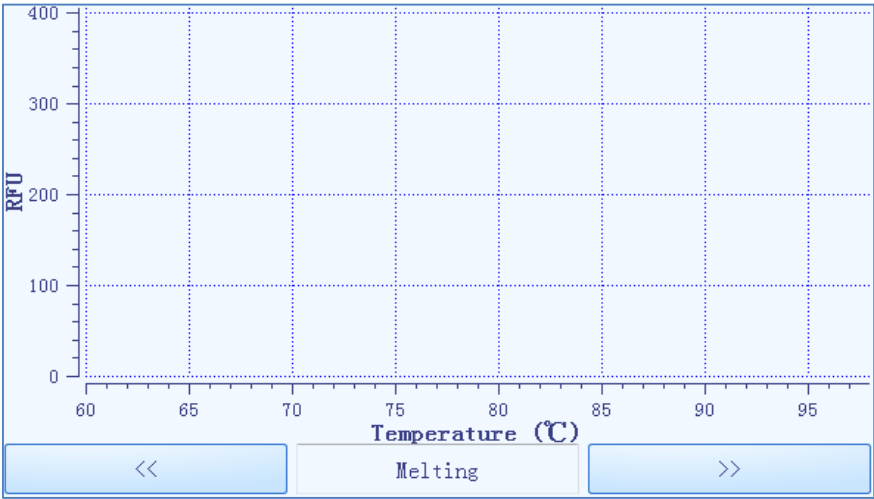


Figure E-4.2 Run Monitoring Diagram - Real-time Melting Curve

- X axis represents the temperature;
- Y axis represents the Relative Fluorescence Unit (RFU);
- User can press any dye under the **Fluorescence** option in the run monitoring option bar, and the run monitoring diagram will only display the corresponding real-time melting curve;


c. User can press  key below the run monitoring diagram to monitor the fluorescence **Heat Map** of the current experiment, as shown in figure E-4.3.



Figure E-4.3 Run Monitoring Diagram - Heat Map

- The fluorescence heat map displays the 48 sample wells corresponding to the sample block;
 - The color bar on the right side of fluorescence heat map displays the change tendency of fluorescence intensity and its corresponding color;
- If user select the **Temperature** option in the run monitoring option bar, run monitoring diagram will display the temperature program of the current experiment, as shown in figure E-4.4.

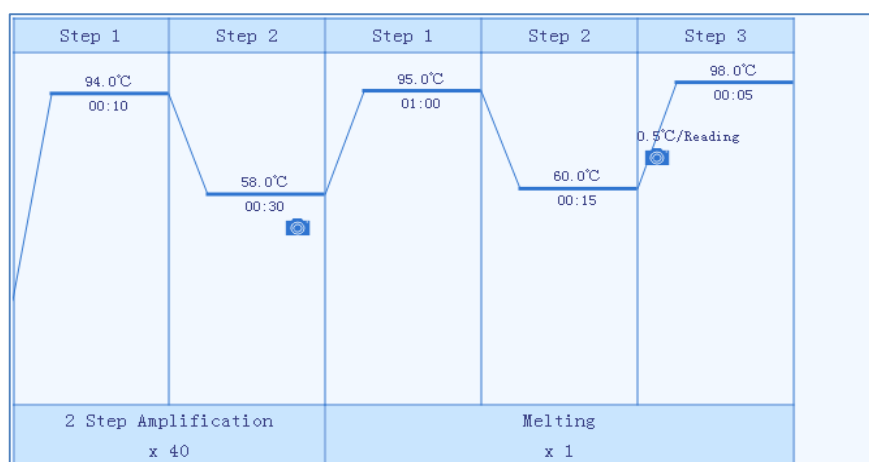


Figure E-4.4 Run Monitoring Diagram - Temperature program

2.4 Result Analysis

After the experiment, user can click the **Result Analysis** tab to analysis the current experiment result.

- The result analysis interface displays the fluorescence heat map of the current experiment by default. User can select a dye in the right **Fluorescence** option column to view the corresponding fluorescence heat map, as shown in figure E-5.1.

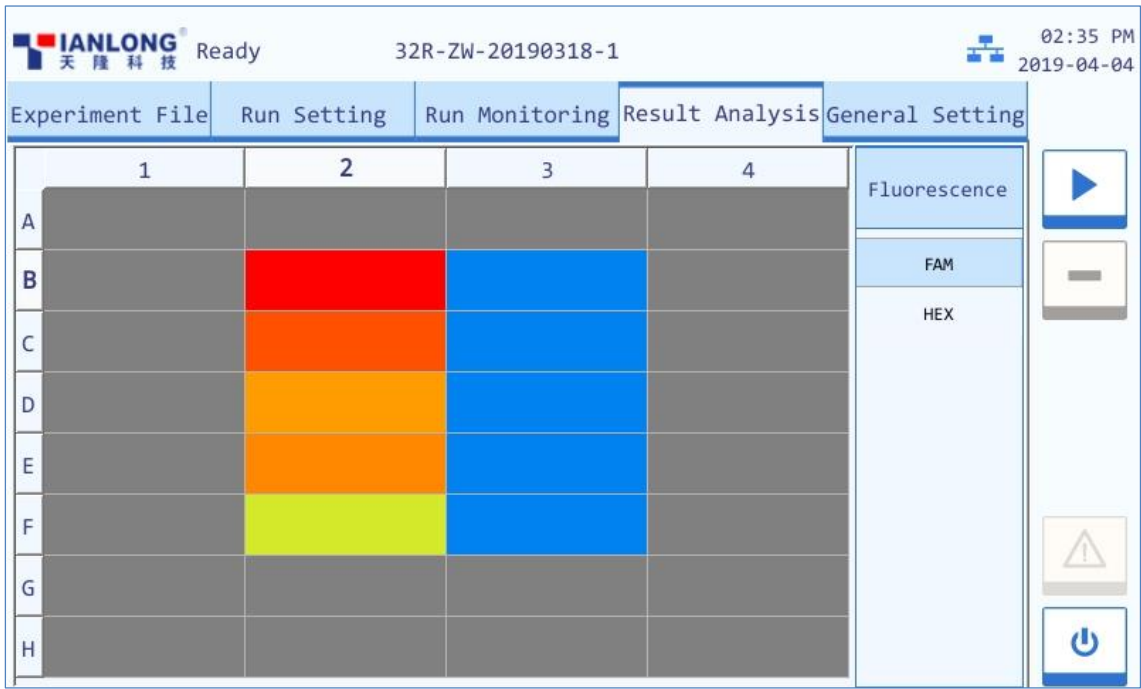


Figure E-5 Run Monitoring Diagram - Fluorescence heat map

- User can press any well on the fluorescence heat map to check the amplification result of this sample, as shown in figure E-5.2.

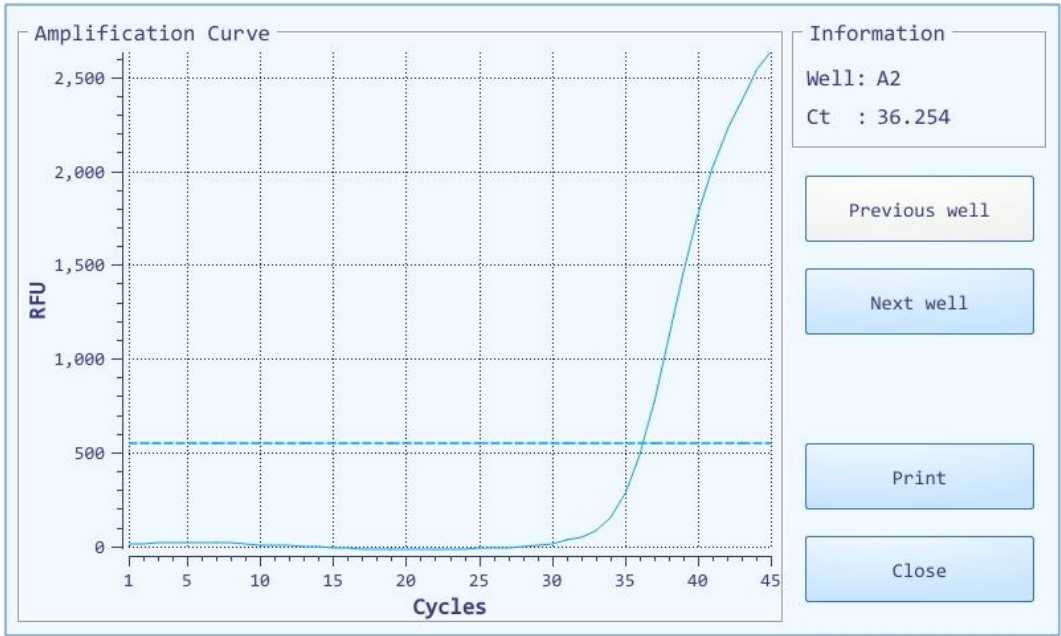


Figure E-5 Run Monitoring Diagram - Amplification result

- The amplification result interface displays the Amplification Curve, Well and Ct value information.

- Use can click **Last Well** to check the amplification result of the last well sample.
- Use can click **Next Well** to check the amplification result of the last well sample.
- Use can click **Print** to print the amplification result of the current sample.
- Use can click **Close** to close the amplification result interface.

2.5 General Setting

The General Setting tab consists of **Instrument**, **Configuration** and **Service** three sub-tabs.

2.5.1 Instrument Sub-tab

The General Setting tab displays the **Instrument** sub-tab by default, as shown in figure E-6.1.

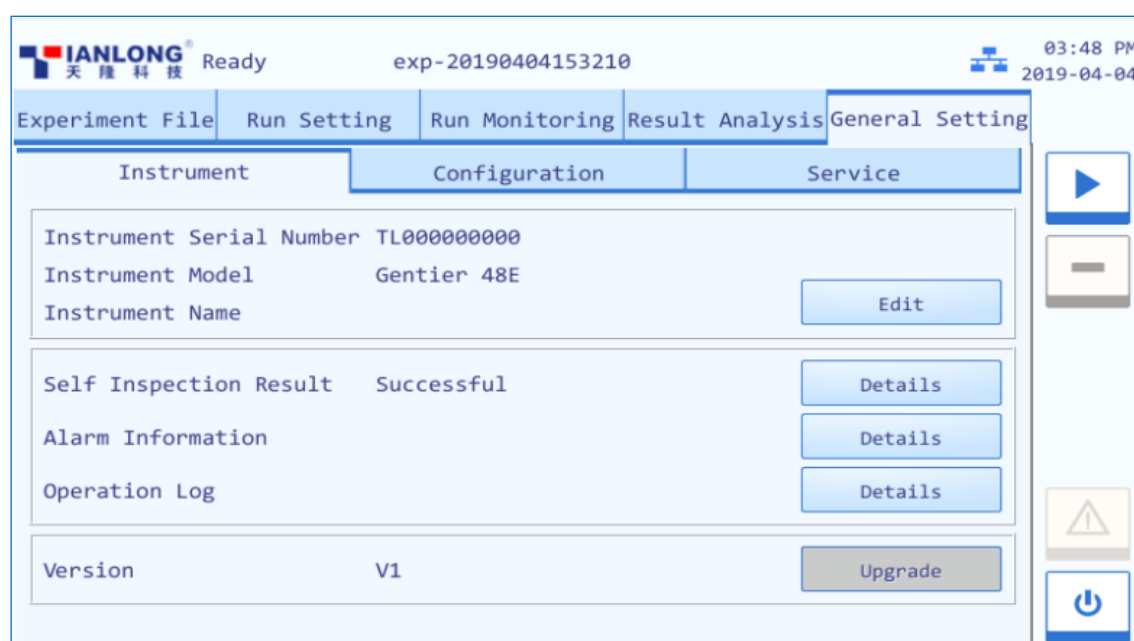
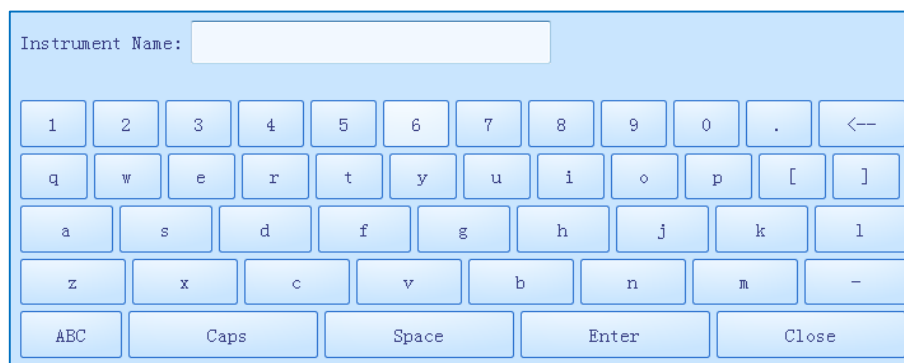


Figure E- 6.1 Instument software – General Setting tab - Instrument sub-tab

Introduction to Instrument sub-tab

- **Instrument Serial Number:** display the factory serial number of the current Gentier instrument.
- **Instrument Model:** display the current Gentier instrument model.
- **Instrument Name:** display the name of the current Gentier instrument, user can press < **Edit** > and the instrument software will pop up the instrument name edit window, as shown in figure E-6.1a.



The image shows a software window titled 'Instrument Name:'. It features a text input field at the top. Below the input field is a virtual keyboard layout with five rows of buttons. The first row contains digits 1-0 and a backspace key. The second row contains letters q-p and brackets. The third row contains letters a-l. The fourth row contains letters z-m and a hyphen/underscore key. The fifth row contains four larger buttons labeled 'ABC', 'Caps', 'Space', 'Enter', and 'Close'.

Figure E-6.1a Instrument sub-tab - Instrument Name edit window

- ▶ **Self Inspection:** display the latest self-inspection result of the current Gentier instrument, user can press < **Details** > to view detailed self-inspection information.
 - ▶ **Alarm Information:** user can press < **Details** > to view the detailed alarm information occurred during the self-inspection and experiment running.
 - ▶ **Operation Log:** user can press < **Details** > to view all the operations have been executed on the current Gentier instrument.
 - ▶ **Version:**
 - User can press < **Details** > to view the system version information of the current Gentier instrument.
 - User can press < **Upgrade** > to upgrade instrument system.
-

2.5.2 Configuration Sub-tab

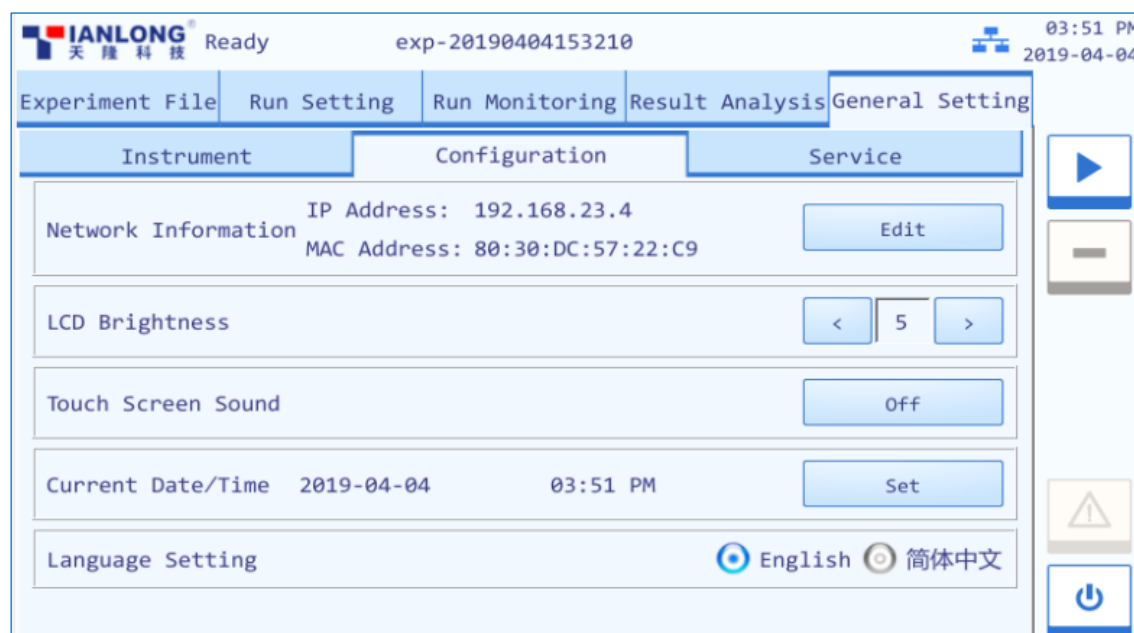


Figure E- 6.2 General Setting tab - Configuration sub-tab



Introduction to Configuration sub-tab

- **Network Information:** display the network information of the current Gentier instrument. User can press < **Edit** > and the instrument software will pop up the network information window, as shown in figure E-6.2a.



Figure E-6.2a Configuration sub-tab - Network Information window

- In the IP Setting box, user can view all the network IP addresses, or press < **Edit** > to modify the corresponding parameter.
- User can press < **Default** > to recover the default network IP settings of the current instrument.

- User can press < **Back** > to turn back to the configuration tab.
- ▶ **LCD Brightness:** user could press  and  icons to adjust the screen brightness (dark: 1 ~ bright: 5).
- ▶ **Touch Screen Sound:** user could press < **Open** > or < **Close** > to decide whether to enable the touch screen sound.
- ▶ **Current Date/Time:** user can press < **Set** > to set the date and time displayed on the current Gentier instrument, the date/time setting window will pop up as shown in E-6.1a.



2017-02-21						
<	<<	2	2017	>>	>	
Sun	Mon	Tue	Wed	Thu	Fri	Sat
29	30	31	1	2	3	4
5	6	7	8	9	10	11
12	13	14	15	16	17	18
19	20	21	22	23	24	25
26	27	28	1	2	3	4
5	6	7	8	9	10	11

15:33:18

Hour : 15  

Minute: 33  

Second: 18  

Date format







Format: yyyy-MM-dd

☒ 24-hour

☐ 12-hour

Confirm Back

Figure E-6.1a Instrument sub-tab - Date/time setting window

- User can press  and  icons to set the displayed year.
- User can press  and  icons to set the displayed month.
- User can set the displayed date on the calendar.
- User can press  and  icons to set the displayed time, including Hour, Minute and Second.
- In the **Date Format** setting box, user can set the displayed date format in the **Format** drop-down list; user can also press **24-hour** or **12-hour** option to set the displayed time format.
- User can press < **Confirm** > to confirm the date and time setting;
- User can press < **Back** > to return to the **Instrument** sub-tab interface.
- ▶ **Language Setting:** user could set the instrument system language, the two options provided are **English** and **Simplified Chinese**.

2.5.3 Service Sub-tab

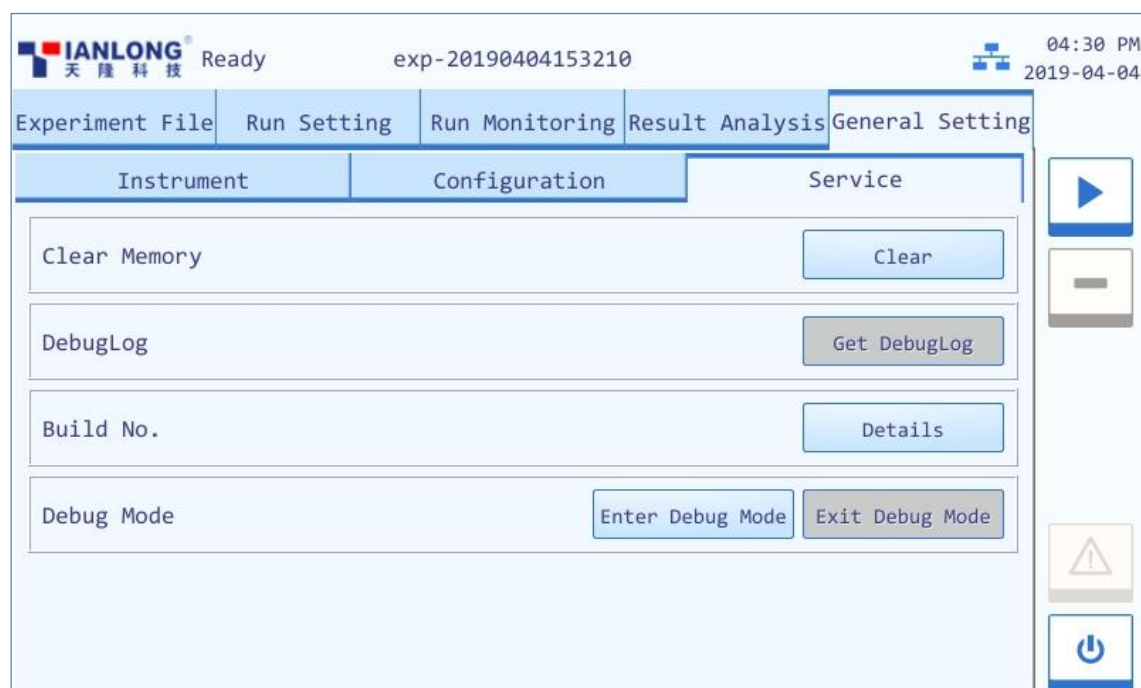


Figure E-6.3 General Setting tab - Service sub-tab

Introduction to Service sub-tab






- ▶ **Lock Instrument for Transportation:** user can press < **Lock** > and the system software will guide the operator to lock the transport lock. For detailed installation steps of transport, please refer to [F-3.2 Install the Transport Lock](#).
- ▶ **Clear Memory:** user can press < **Clear** > to clear all experiment files.
- ▶ **DebugLog:** user could connect an USB drive to the Gentier instrument and press press < **Get DebugLog** > to download all the log file of the current Gentier instrument, this will help our professional engineer to get the relative information in case the instrument require maintenance.
- ▶ **Build No. :** user could press < **Details** > to view the build number of the current instrument.
- ▶ **Debug Mode:** press < **Enter Debug Mode** > and input the password, the instrument software will pop up the debug interface.
- ⚠ **Reminding:** no person except the professional engineers from our company is allowed to enter the debug mode.

F. Instrument Cleaning and Maintenance

Under proper use condition, the Gentier instrument requires little maintenance. However, the Gentier instrument should be cleaned and maintained on a regular schedule for long time and constant use. This section includes the information on cleaning and maintenance of the Gentier instrument.

1. Instrument Cleaning Operation Instructions


The Gentier instrument should be cleaned on a regular schedule (every other month), please carefully read the following instructions before cleaning the instrument.


-  **Prohibit:** never clean the instrument when it is electrified.
-  **Prohibit:** never pour water or other solutions in into the sample block or any interior parts of the instrument. Fluids can cause electrical shock when the instrument is electrified.
-  **Caution:** ethanol is a flammable and volatile liquid, it's exposure may irritate eyes, skin and respiratory tract, and may lead to central nervous system hypofunction and liver damage. Please wear appropriate protective goggles, clothing and gloves, when using ethanol to clean.
-  **High-Temp:** the sample block and hot lid may produce high temperature during running, do not clean the sample block until it reaches the room temperature.
-  **Biohazard:** please regard all samples as potential biohazard materials, universal safety precautions should be taken when handling or processing samples. Samples pills should be immediately disinfected with an appropriate disinfectant solution to avoid equipment contamination or user' personnel injury.

1.1 Clean Instrument Shell

1st step: switch off the instrument and unplug the power cord;

2nd step: clean the instrument shell with a piece of damp, soft cloth, and if needed, please rinse the cloth with mild commercial detergent for cleaning.

-  **Prohibit:** do not spray the detergent directly on the instrument, as malfunctions of the electronics or may occur.


 **Caution:** please do not use organic or strong detergent to clean the instrument shell, which may ruin the surface coating.


1.2 Clean Touch Screen

1st step: switch off the instrument and unplug the power cord;

2nd step: gently wipe the touch screen with a piece of dry, soft cloth to remove dust, oil or fingerprints.

3rd step: if the touch screen is still not clean, use a piece of damp, soft cloth that moistened with low concentration isopropanol or ethanol to clean the touch screen in a gentle motion.

 **Prohibit:** do not spray the detergent directly on the touch screen, as malfunctions of the electronics or may occur.


 **Caution:** do not use abrasive detergent or rough material, as they may scratch the touch screen.


1.3 Clean Sample Block

1st step: switch off the instrument and unplug the power cord;

2nd step: open the top lid and clean the sample block surface with a piece of damp, soft cloth, and if needed, please rinse the cloth with mild commercial detergent.

3rd step: clean the sample block wells with the degreased cotton swab moistened with ethanol to ensure the firm contact and good thermal conductivity between the consumables and the wall of block well.

 **Prohibit:** do not spray the detergent directly on the thermal cycler block, as malfunctions of the electronics or may occur.

 **Caution:** do not close the top lid before the sample block is completely dry.

2. Instrument Maintenance Operation Instructions

2.1 Maintain Sufficient Air Flow

The placing area of Gentier instrument should be checked regularly, as it requires sufficient air flow to precisely reach the correct target temperature. Please ensure that the air flow is unrestricted and other items around the instrument do not interfering with the air flow.

2.2 Maintain Stable Power Supply.


The Gentier instrument requires stable power supply for proper functioning, therefore its power supply should be checked regularly to ensure the consistent of instrument required voltage and the power supply voltage (allowable deviation $\pm 10\%$). And make sure that the rated load of receptacle is no less than the requirement of instrument.


2.3 Maintain Instrument Cleanliness


Contamination of sample block or optical parts can interfere with thermal cycling and data collection.

Avoid contaminating the Gentier instrument:

- Always clean the outside surface of consumables before placing them in the sample block.
- Clean the sample block periodically to prevent the buildup of dirt, biohazardous materials or solutions.
- Always seal the consumables with sealing film before running the experiment.

 **Caution:** Never place a consumable with open or leak cap in the sample block. The reagents may escape during heating and cover the sample block and hot lid.

 **Caution:** Never run a PCR reaction with volatile reagents that could contaminate the sample block and hot lid.

 **Caution:** In case the instrument will not be used for a long time, unplug it and cover the instrument with a soft cloth or plastic bags to prevent dust from entering.

2.4 Replace Fuse Tube

The fuse tube (type 250V~/F10AH) of Gentier instrument is located in the fuse tube box near the power outlet at rear of the instrument. Before replacing the fuse tube, please switch off the instrument and unplug the power plug, use a screwdriver to pry open the fuse tube box. Then substitute the old fuse tube with an equal one, as shown in figure F-1.

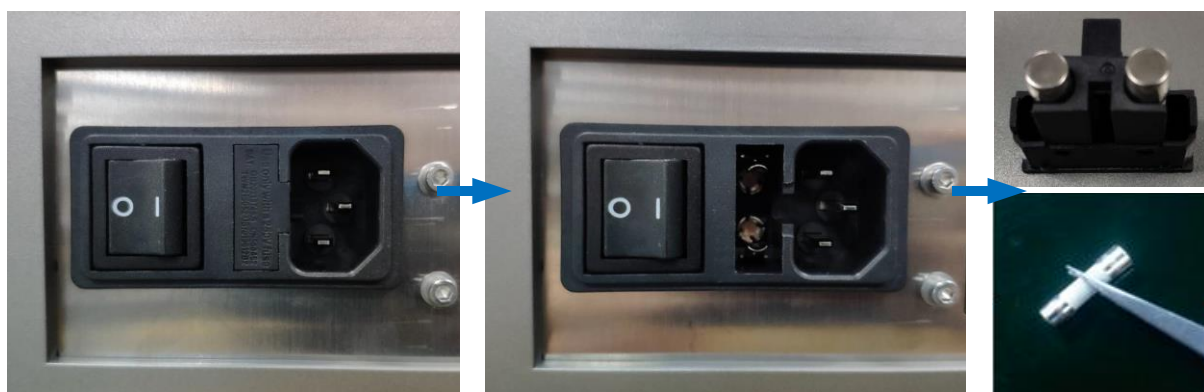


Figure F-1. Replace the fuse tube

Reminding: in case there is no display on the screen after starting up the Genesy thermal cycler, please check the fuse tube.

Caution: improper fuse tube may lead to circuit system damage, even fire.

3. Transportation or Return to Factory

3.1 Instrument Disinfection

In case the Gentier instrument will be moved to another lab or needed to return to the after-sale service department for maintenance, please first disinfection the instrument and fill in the disinfection certificate.


The disinfection process of Gentier instrument is listed as follows:


- ▶ Please wear protective clothing and medical disposable gloves;
- ▶ Open the top lid and get all consumables out of the sample block;
- ▶ Switch off the instrument and unplug the power cord;

- ▶ Prepare the detergent (lab routine disinfectant), wet the cotton cloth and degreased cotton swabs with the prepared detergent;
- ▶ Clean the sample block surface with the cotton cloth, clean the sample block well with the degreased cotton swabs, and leave the top lid open until it is dry;
- ▶ Clean the sample block again with 75% ethanol;
- ▶ Clean the instrument shell with mild detergent;
- ▶ Open the top lid and leave the instrument under UV light for 1 hour disinfection.

3.2 Product Packaging

Please use the original packaging materials to properly pack the Gentier instrument and its accessories in order to prevent the collision and oscillation during transportation.

 **Caution:** The original transport package of Gentier instrument is designed to reduce the instrument damage and ensure its transportation safety. Adopt other packaging materials will broke the warranty, and **Drawell.** will not be responsible for damages as consequences of improper packaging that incurred during the transportation back to maintenance department.

 **Reminding:** the transportation of Gentier instrument can adopt general transport (with awnings).

The specific packing steps of Gentier instrument are as follows:

- 1st Step:** pack the instrument with plastic bag and prepare the instrument accessories, as shown in figure F-4a.
- 2nd Step:** place the bottom protective foam into the carton, as shown in figure F-4b.
- 3rd Step** insert the instrument into the bottom protective foam, as shown in figure F-4c.
- 4th Step:** place the top protective foam on top of the instrument and put the accessories in side the carton and, as shown in figure F-4d F-4e.
- 5th Step:** use wide adhesive tape to seal the carton cover, as shown in figure F-4f.

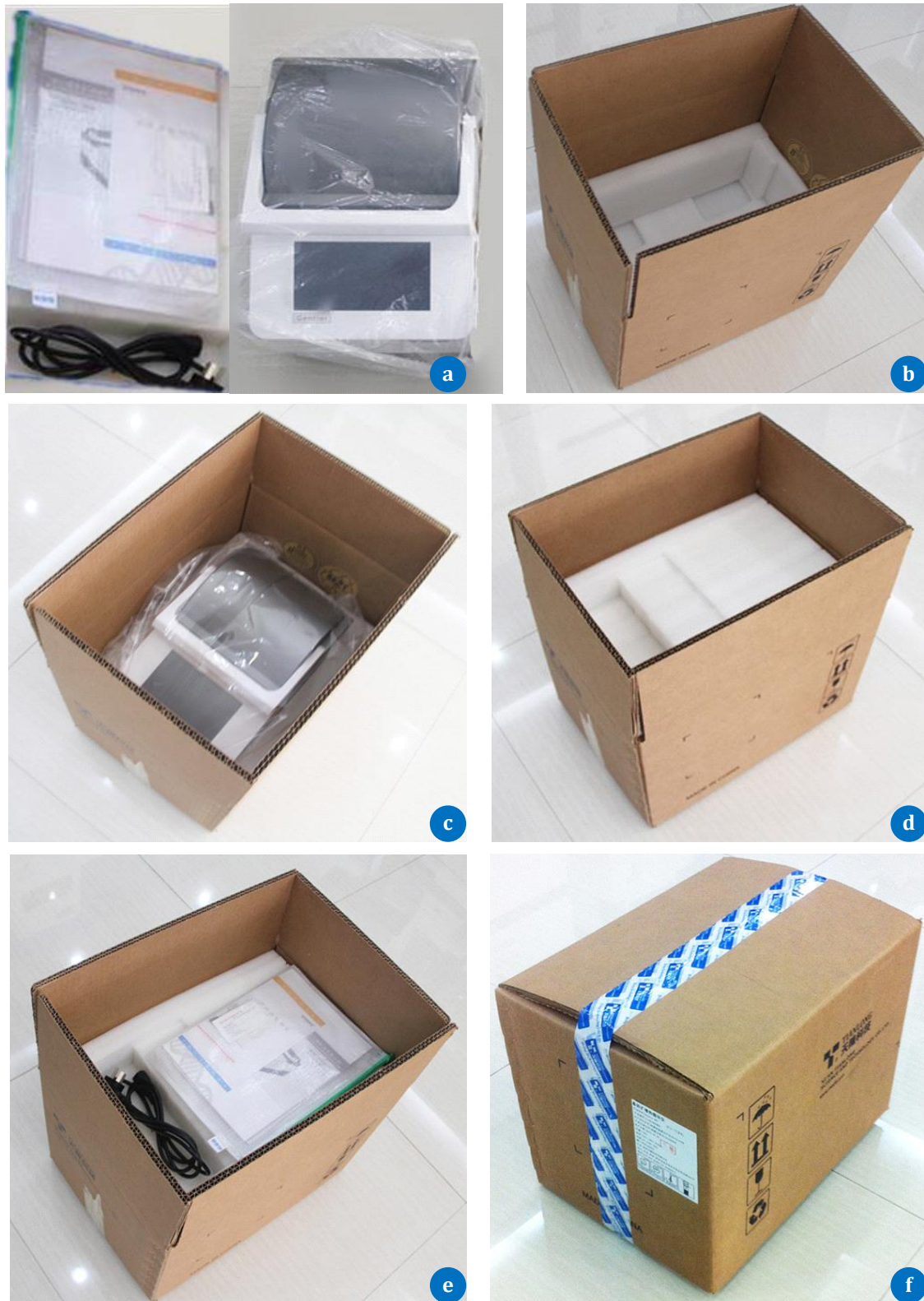


Figure F-4. The packing steps of Gentier instrument


4. Troubleshooting


In general, corrective instructions will be display along with the error messages by the application or instrument software. And under normal circumstances the software running errors can be solved by restarting the computer or the instrument system.

This section describes the main possible errors of the Gentier Instrument together with possible causes and corrective instructions.

No.	Error	Possible Cause	Corrective Instructions
1	No display on the screen	Without power supply	Plug the power supply
		Power switch is on 'off' position	Switch on the power switch
		Unstable power cord connection	Connect the power cord again or renew the power cord
		Inappropriate power voltage	Adjust the scope of power voltage into a normal range
		Fuse tube damaged	Replace the fuse tube
		Others	Contact us
2	Boot screen display error message	The activation of system failed.	Please contact us and consult the maintenance engineer
		Power voltage too low	Ensure there is no other appliance or circuit in the same electric circuit.
3	System crashed or out of control	Improper operation	Restart the instrument system
		Others	Contact us
4	Temperature does not rise while heating	Check temperature control setting	Startup the temperature control
		Others	Contact us
5	No Experiment Results	Wrong operation process	Check the operation process and test again
		Quality problems of reagents	Renew the reagent and run the experiment again.
		Experiment settings do not meet the requirements	Reset the experiment procedure

		(temperature is inappropriate or cycle number is not enough).	
		Others	Contact us
6	Abnormal ramp or incorrect temperature	Air vent is blocked	Clean up the air vent.
		Connecting lines are loose.	Contact us

 **Caution:** in case you cannot judge and eliminate these failures by yourself, please directly contact with our company.

 **Caution:** in case any of following situation occurs, please immediately cut off the power supply and contract us. We will arrange qualified maintenance personnel for processing.

- Any liquid has entered into the instrument;
- Abnormal sound or smell appears inside the instrument;
- Instrument is soaked with water or rain;
- Any housing damage caused by accidently drop of the instrument;
- Obvious functional changes of instrument.

G. After-sale Service Commitments

1. Warranty Service

1. In the condition that user comply with the rules of transportation and operation. Our company guarantees to exchange the Gentier instrument in case any malfunction is caused by the defects of materials and instrument manufacture within 1 month after sale.
2. We will provide 24 months of warranty period beginning from the delivery of Gentier instrument. Within the warranty period our company guarantees the maintenance of the instrument in case any malfunction is caused by the defects of materials and instrument manufacture. User could contact us, and we will arrange maintenance personnel for processing (We promise to provide the list of components and the electric circuit for the maintenance personnel). Or directly send the instrument back to the maintenance department assigned by our company (User shall be responsible for the transportation fees), and we will send back the well maintained instrument to the user for free.
3. For post warranty service, we will charge for the maintenance fee according to specific situation.
4. The life span of Gentier instrument is about 5 years, for any maintenance of the instrument that have been used beyond 5 years, our company will not take any responsibilities.
5. The following circumstances are not within the scope of warranty:
 - The instrument damages caused by improper, negligent operation or force majeure including: war, fire, flood, earthquake, typhoon and any other unforeseen accidents.
 - The parts or components damages caused by abnormal voltage.
 - User did not comply with the rules of transportation and use.
 - User did not comply with the maintenance instructions.
 - The Gentier instrument has been opened or maintained by person, manufacturers or agents that are not authorized by **Drawell**

2. Response Time

We will make response within 24 hours upon receiving the notification, no matter whether the Gentier instrument is within the warranty period or not. For any problems that cannot be solved through the phone, we will provide on-site service within 7 days for the customs located in China.

3. Contact Information

Company Name: Chongqing Drawell Instrument CO.,Ltd

Company Address: Suite 2605,Building No.11,(Evergrande City Square) Shiyou Road No.1,Yuzhong District,Chongqing,China.

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Company Tel: 0086-023-63268643

Email Address: sales05@drawell.com.cn

Website Address: www. drawell.com.cn