resDNASEQ™ Quantitative DNA Kits

USER GUIDE

for use with:
(Genomic DNA quantitation)
resDNASEQ™ Quantitative CHO DNA Kit
resDNASEQ™ Quantitative *E. coli* DNA Kit
resDNASEQ™ Quantitative HEK293 DNA Kit
resDNASEQ™ Quantitative Human DNA Kit
resDNASEQ™ Quantitative MDCK DNA Kit
resDNASEQ™ Quantitative NS0 DNA Kit
resDNASEQ™ Quantitative *Pichia* DNA Kit
resDNASEQ™ Quantitative Sf9 and Baculovirus DNA Kit
resDNASEQ™ Quantitative Vero DNA Kit
(Plasmid DNA quantitation)
resDNASEQ™ Quantitative Plasmid DNA - Kanamycin Resistance Gene Kit

Catalog Numbers 4402085, 4458435, A46014, A26366, 4464335, 4458441, 4464336, A46066, A41797, and A50337

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Revision L

Life Technologies Ltd | 7 Kingsland Grange | Woolston, Warrington WA1 4SR | United Kingdom
For descriptions of symbols on product labels or product documents, go to thermofisher.com/symbols-definition.

Revision history: 4469836 L (English)

Revision	Date	Description
L	28 July 2023	Update for release of AccuSEQ™ Real-Time PCR Detection Software v2.2 and AccuSEQ™ Real-Time PCR Software v3.2.
K	8 November 2021	Update to include the resDNASEQ™ Quantitative Synthetic Vero DNA Kit (Cat. No. A53242).
J	28 April 2021	Update to the control serial dilutions required for the resDNASEQ™ Quantitative Plasmid DNA - Kanamycin Resistance Gene Kit (Cat. No. A50337).

Contents

1 Product information

IMPORTANT! Before using this product, read and understand the information in the “Safety” appendix in this document.

Product description

The resDNASEQ™ Quantitative DNA Kits are used to quantitate residual DNA from CHO, *E. coli*, HEK293, Human, MDCK, NS0, *Pichia*, Sf9 and Baculovirus, and Vero, or plasmid DNA for Kanamycin resistance, in cell lines which are used for production of biopharmaceutical products. Use the kit after you extract host-cell DNA from test samples. For extraction information, see the *PrepSEQ™ Sample Preparation Kits User Guide* (Pub. No. 4469838).

The resDNASEQ™ Quantitative DNA Kits use TaqMan™ quantitative PCR to perform rapid, specific quantitation of sub-picogram levels of residual host-cell or plasmid DNA. The assay is accurate and reliable across a broad range of sample types, from in-process samples to final product.

To generate the standard curve used to quantitate the DNA in test samples, the CHO, MDCK, NS0, Vero, synthetic Vero, and Plasmid DNA - Kanamycin resistance assays require six dilutions and the *E. coli*, HEK293, Human, *Pichia*, and Sf9 and Baculovirus assays require five dilutions. Control DNA for standard curve generation is included in the kits. In addition, the kits use an internal positive control (IPC) to evaluate the performance of each PCR reaction.

IMPORTANT! We are committed to providing an exceptional product experience. Reach out to us for technical support and product training to help you every step of the way, see “Customer and technical support” on page 72.

Contents

Table 1 resDNASEQ™ Quantitative CHO DNA Kit (Cat. No. 4402085)

Contents	Amount	Storage
resDNASEQ™ CHO Real-Time PCR Reagents		
TaqMan™ Environmental Master Mix 2.0	2 × 0.75 mL	–25°C to –15°C before first use, protect from light 2–8°C after first use, protect from light
10X CHO DNA Real-Time PCR Assay Mix	300 µL	–25°C to –15°C, protect from light
Negative Control (Water)	1.0 mL	–25°C to –15°C before first use 2–8°C after first use
resDNASEQ™ CHO DNA Control		
CHO DNA Control, 30 ng/µL	40 µL	–25°C to –15°C
DNA Dilution Buffer	7 mL	2–8°C

Table 2 resDNASEQ™ Quantitative *E. coli* DNA Kit (Cat. No. 4458435)

Contents	Amount	Storage
resDNASEQ™ Real-Time PCR Reagents		
TaqMan™ Environmental Master Mix 2.0	2 × 0.75 mL	–25°C to –15°C before first use, protect from light 2–8°C after first use, protect from light
Negative Control (Water)	1.0 mL	–25°C to –15°C before first use 2–8°C after first use
10X *E. coli* DNA Assay Mix	300 µL	–25°C to –15°C, protect from light
resDNASEQ™ *E. coli* DNA Control		
E. coli DNA Control, 30 ng/µL	40 µL	–25°C to –15°C
DNA Dilution Buffer	7 mL	2–8°C

Table 3 resDNASEQ™ Quantitative HEK293 DNA Kit (Cat. No. A46014)

Contents	Amount	Storage
resDNASEQ™ HEK293 DNA Real-Time PCR Reagents		
TaqMan™ Environmental Master Mix 2.0	2 × 0.75 mL	–25°C to –15°C before first use, protect from light 2–8°C after first use, protect from light
10X HEK293 Assay Mix	300 µL	–25°C to –15°C, protect from light
Negative Control (Water)	1.0 mL	–25°C to –15°C before first use 2–8°C after first use
resDNASEQ™ HEK293 DNA Control		
HEK293 DNA Control, 30 ng/µL	40 µL	–25°C to –15°C
DNA Dilution Buffer	7 mL	2–8°C

Table 4 resDNASEQ™ Quantitative Human DNA Kit (Cat. No. A26366)

Contents	Amount	Storage
resDNASEQ™ Human Real-Time PCR Reagents		
TaqMan™ Environmental Master Mix 2.0	2 × 0.75 mL	–25°C to –15°C before first use, protect from light 2–8°C after first use, protect from light
10X Human DNA Assay Mix	300 µL	–25°C to –15°C, protect from light
Negative Control (Water)	1.0 mL	–25°C to –15°C before first use 2–8°C after first use
resDNASEQ™ Human DNA Control		
Human DNA Control, 30 ng/µL	40 µL	–25°C to –15°C
DNA Dilution Buffer	7 mL	2–8°C

Table 5 resDNASEQ™ Quantitative MDCK DNA Kit (Cat. No. 4464335)

Contents	Amount	Storage
resDNASEQ™ Real-Time PCR Reagents		
TaqMan™ Environmental Master Mix 2.0	2 × 0.75 mL	–25°C to –15°C before first use, protect from light 2–8°C after first use, protect from light
Negative Control (Water)	1.0 mL	–25°C to –15°C before first use 2–8°C after first use
10X MDCK DNA Assay Mix	300 µL	–25°C to –15°C, protect from light

Table 5 resDNASEQ Quantitative MDCK DNA Kit (Cat. No. 4464335) *(continued)*

Contents	Amount	Storage
resDNASEQ™ MDCK DNA Control		
MDCK DNA Control, 30 ng/µL	40 µL	–25°C to –15°C
DNA Dilution Buffer	7 mL	2–8°C

Table 6 resDNASEQ™ Quantitative NS0 DNA Kit (Cat. No. 4458441)

Contents	Amount	Storage
resDNASEQ™ Real-Time PCR Reagents		
TaqMan™ Environmental Master Mix 2.0	2 × 0.75 mL	–25°C to –15°C before first use, protect from light 2–8°C after first use, protect from light
Negative Control (Water)	1.0 mL	–25°C to –15°C before first use 2–8°C after first use
10X NS0 DNA Assay Mix	300 µL	–25°C to –15°C, protect from light
resDNASEQ™ NS0 DNA Control		
NS0 DNA Control, 30 ng/µL	40 µL	–25°C to –15°C
DNA Dilution Buffer	7 mL	2–8°C

Table 7 resDNASEQ™ Quantitative *Pichia* DNA Kit (Cat. No. 4464336)

Contents	Amount	Storage
resDNASEQ™ Real-Time PCR Reagents		
TaqMan™ Environmental Master Mix 2.0	2 × 0.75 mL	–25°C to –15°C before first use, protect from light 2–8°C after first use, protect from light
Negative Control (Water)	1.0 mL	–25°C to –15°C before first use 2–8°C after first use
10X *Pichia* DNA Assay Mix	300 µL	–25°C to –15°C, protect from light
resDNASEQ™ *Pichia* DNA Control		
Pichia DNA Control, 30 ng/µL	40 µL	–25°C to –15°C
DNA Dilution Buffer	7 mL	2–8°C

Table 8 resDNASEQ™ Quantitative Sf9 and Baculovirus DNA Kit (Cat. No. A46066)

Contents	Amount	Storage
resDNASEQ™ Real-Time PCR Reagents		
TaqMan™ Environmental Master Mix 2.0	2 × 0.75 mL	–25°C to –15°C before first use, protect from light 2–8°C after first use, protect from light
Negative Control (Water)	1.0 mL	–25°C to –15°C before first use 2–8°C after first use
10X Sf9 + Baculovirus DNA Assay Mix	300 µL	–25°C to –15°C, protect from light
resDNASEQ™ Sf9 and Baculovirus DNA Control		
Multiplex DNA Control with: • Sf9 DNA Control, 30 ng/µL • Baculovirus DNA Control, 30 ng/µL	40 µL	–25°C to –15°C
DNA Dilution Buffer	7 mL	2–8°C

Table 9 resDNASEQ™ Quantitative Vero DNA Kit (Cat. No. A41797)

Contents	Amount	Storage
resDNASEQ™ Real-Time PCR Reagents		
TaqMan™ Environmental Master Mix 2.0	2 × 0.75 mL	–25°C to –15°C before first use, protect from light 2–8°C after first use, protect from light
Negative Control (Water)	1.0 mL	–25°C to –15°C before first use 2–8°C after first use
10X Vero DNA Assay Mix	300 µL	–25°C to –15°C, protect from light
resDNASEQ™ Vero DNA Control		
Vero DNA Control, 30 ng/µL	40 µL	–25°C to –15°C
DNA Dilution Buffer	7 mL	2–8°C

Table 10 resDNASEQ™ Quantitative Plasmid DNA - Kanamycin Resistance Gene Kit (Cat. No. A50337)

Contents	Amount	Storage
resDNASEQ™ Kanamycin DNA Real-Time PCR Reagents		
TaqMan™ Environmental Master Mix 3.0	2 × 0.75 mL	–25°C to –15°C before first use, protect from light 2–8°C after first use, protect from light
10X KanR Assay Mix	300 µL	–25°C to –15°C, protect from light
Yeast tRNA (10 mg/mL)	500 µL	–25°C to –15°C

Table 10 resDNASEQ Quantitative Plasmid DNA - Kanamycin Resistance Gene Kit (Cat. No. A50337) *(continued)*

Contents	Amount	Storage
Negative Control (Water)	1.0 mL	–25°C to –15°C before first use 2–8°C after first use
resDNASEQ™ Kanamycin DNA Control		
KanR DNA Control, 3 x 10^7 copies/µL	44 µL	–25°C to –15°C
DNA Dilution Buffer	7 mL	2–8°C

Required materials not supplied

Unless otherwise indicated, all materials are available through **thermofisher.com**. "MLS" indicates that the material is available from **fisherscientific.com** or another major laboratory supplier.

Item	Source
Instrument	
QuantStudio™ 5 Real-Time PCR System with AccuSEQ™ Real-Time PCR Software v3.1 or later	Contact your local sales representative.
Consumables	
MicroAmp™ Optical 96-Well Reaction Plate with Barcode, 20 plates, 0.1-mL wells; for use with Applied Biosystems™ 7500 Fast Real-Time PCR System	4346906
MicroAmp™ 96-Well Base	N8010531, 10 bases
MicroAmp™ Optical Adhesive Film	4311971, 100 covers 4360954, 25 covers
MicroAmp™ Adhesive Film Applicator	4333183, 5 applicators
Miscellaneous items	
Disposable gloves	Major lab supplier (MLS)
Pipettes	MLS
Aerosol-resistant micropipette tips	MLS
For the PCR plate: Fisher Scientific™ Mini Plate Spinner Centrifuge, 120- or 230-volt	14-100-143 (120-volt), 14-100-141 (230-volt)
Nonstick, RNase-free Microfuge Tubes, 1.5 mL (1 box; 250 tubes/box)	AM12450

Workflows

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2 Genomic DNA quantitation

Prepare the control DNA serial dilutions for the standard curve

Guidelines for standard dilutions

- Prepare the standard curve and the test samples in different areas of the lab.
- Use different sets of pipettors for test sample preparation and for standard curve preparation and aliquoting to avoid cross-contamination of test samples.
- Vortex each tube to mix the contents thoroughly before each dilution step.
- Briefly centrifuge to collect all the liquid at the bottom before making the next dilution.

Prepare the control serial dilutions (Genomic DNA)

1. Label nonstick 1.5-mL microfuge tubes: **NTC, SD1**, **SD2**, **SD3**, **SD4**, **SD5**.
   For CHO, Vero, MDCK, and NS0 kits, label an additional tube with **SD6**.
2. Add 50 µL of DNA Dilution Buffer to tube NTC. Put aside.
3. Add 990 µL of DNA Dilution Buffer to tube SD1.
4. Add 450 µL of DNA Dilution Buffer to tubes SD2, SD3, SD4, SD5, and (for CHO, Vero, MDCK, and NS0 only) SD6.
5. Remove the tube of DNA control (30 ng/µL) from the freezer.
6. After the DNA thaws, vortex it gently for 2 seconds, then briefly centrifuge.
7. Perform the serial dilutions:
   a. Add 10 µL of the DNA control to the tube that is labeled SD1, then vortex thoroughly and briefly centrifuge.
   b. Transfer 50 µL of the DNA from tube SD1 to tube SD2, then vortex thoroughly and briefly centrifuge.

c. Continue to transfer 50 µL of DNA from the previous dilution tube to the next dilution tube until you add DNA to tube SD5 (*E. coli*, HEK293, Human, *Pichia*, and Sf9 and Baculovirus) or SD6 (CHO, Vero, MDCK, and NS0). Final dilutions are shown in the table. After each transfer, vortex thoroughly, then centrifuge briefly.

Serial dilution (SD) tube	Dilution	pg DNA/reaction (10 µL of the diluted DNA used in final 30 µL of PCR reaction)
Control	DNA control tube	300,000 pg
SD 1	10 µL DNA control + 990 µL DDB	3,000 pg
SD 2	50 µL SD1 + 450 µL DDB	300 pg
SD 3	50 µL SD2 + 450 µL DDB	30 pg
SD 4	50 µL SD3 + 450 µL DDB	3 pg
SD 5	50 µL SD4 + 450 µL DDB	0.3 pg
SD 6 (for CHO, Vero, MDCK, and NS0 only)	50 µL SD5 + 450 µL DDB	0.03 pg

8. Store the DNA dilution tubes:

Temperature	For use
4°C	Same day
–20°C	≤1 week
–20°C	SD1 in single-use aliquots ≤6 months

Prepare the PCR reaction mix

1. Determine the number of reactions needed for the controls and test samples that you will quantify.
2. Thaw all kit reagents completely at room temperature, thoroughly mix reagents, and briefly centrifuge.
3. Prepare a PCR reaction mix using the reagents and volumes shown in the table below.
   - Multiply the PCR reaction volume for one reaction (30 µL) by the number of reactions that you need to run.
   - Use 10% excess volume to compensate for pipetting losses.

Note: Use reagents from the same lot for all reactions.

Kit reagents	Volume for 1 30-µL reaction	Volume for 36 30-µL reactions (includes 10% overage)
Negative Control (Water)	2 µL	79.2 µL
10X DNA assay mix appropriate for the cell line being tested	3 µL	118.8 µL
TaqMan™ Environmental Master Mix 2.0	15 µL	594 µL
DNA template	10 µL	Add DNA template to each well separately, not as part of Master Mix
Total	**30 µL**	**792 µL** of Master Mix

Prepare the PCR plate

Plate setup differs slightly for each AccuSEQ™ System. See your software user guide for specific instructions. Place samples, NTCs, and standards in different quadrants of the plate.

A factory default template is available for the Quantitative Sf9 and Baculovirus DNA in AccuSEQ™ Real-Time PCR Software v3.2 or later.

Figure 1 Quantitative Sf9 and Baculovirus DNA template default sample plate layout in AccuSEQ™ Real-Time PCR Software v3.2 or later

① Toolbar (in order: Undo, Redo, Copy, Paste, Delete, View)

② Select Item to highlight (Sample, Target, or Task).

Note: If more samples are added, the Target displays as an S in the wells (experiment type–Quantitation).

③ Select Item. For example, Sample 1. Sample 1 replicates are highlighted.

Table 11 Quantitative Sf9 and Baculovirus DNA template plate well descriptions in AccuSEQ™ Real-Time PCR Software v3.2 or later

Name	Description
SC1 to SC6	Standard curve dilutions in triplicate
Matrix	24 samples in triplicate
NTC	No template control in triplicate

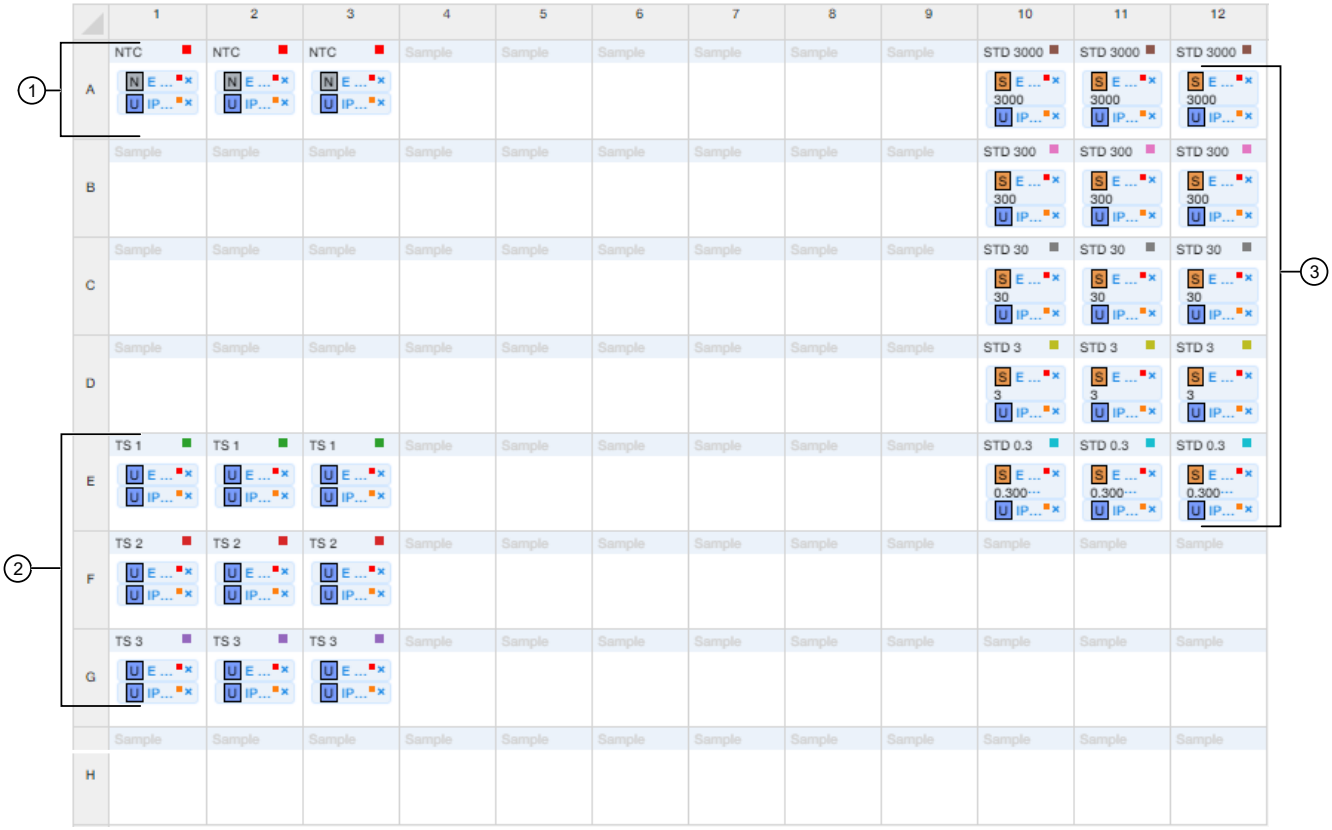


Figure 2 Default plate setup in the resDNA_5Std template in AccuSEQ™ Real-Time PCR Software v3.1 or later

① No template controls
② Samples
③ Standard curve

1. Add 20 µL PCR reaction mix to each well.
2. Add 10 µL of PCR NTC to the appropriate wells.
3. Add 10 µL each of extracted sample DNA to the appropriate wells.

   Note: If you prepared samples with the automated protocol, use a multichannel pipette to transfer the extracted sample.

4. Add 10 µL of standard dilutions to the appropriate wells.

   Note: Use different sets of pipettors to dispense test sample and standard curve dilutions to avoid cross-contamination of test samples.

5. Use a film applicator to seal the plate with optical film, then briefly centrifuge with a miniplate centrifuge that is compatible with 96-well plates.

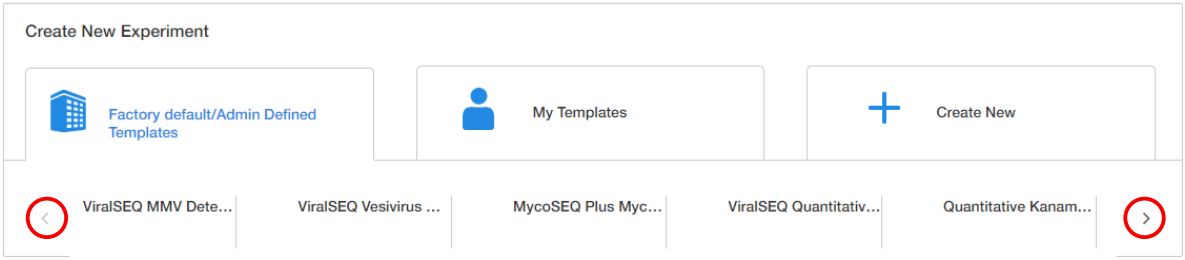
Setup, run, and analyze samples with AccuSEQ™ Software on the QuantStudio™ 5 Real-Time PCR Instrument

Create a resDNASEQ™ experiment

1. In the **Home** screen, click the **Factory default/Admin Defined Template** tab, then select a **resDNASEQ_5std** or **resDNASEQ_6std** template.

   Note: AccuSEQ™ Software v3.2 or later has a **Factory default/Admin Defined Template** for the Quantitative Sf9 and Baculovirus DNA Assay, see "Create a Quantitative Sf9 and Baculovirus DNA experiment" on page 21.

   To navigate to through the available templates click the arrows to either side of the available templates.



Serial Dilutions (Standards)	Template	Assays
5	_5Std	*E. coli*, HEK293, Human, *Pichia*
6	_6Std	CHO, Vero, MDCK, NS0

2. In the **Experiment Properties** pane of the **Setup** tab:
   a. (*Optional*) Change the system-generated name of the experiment.
   b. (*Optional*) Enter the plate **Barcode**, then add **Comments**.

      Default resDNASEQ™ settings (cannot be changed)
      - **Experiment Type** is **Quantitation-Standard Curve**
      - **Chemistry** is **TaqMan™ Reagents**
      - **Ramp Speed** is **Standard - 2hrs**

   c. Click **Next**.

      Note: Experiment names cannot be changed after this step.

3. In the **qPCR Method** pane of the **Setup** tab, view the default volume and cycling conditions (cannot be changed).

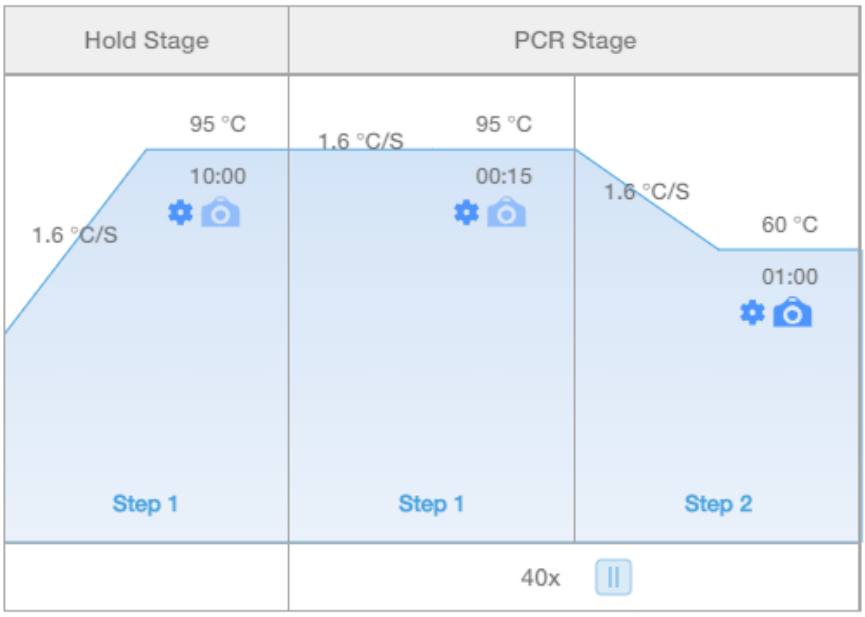
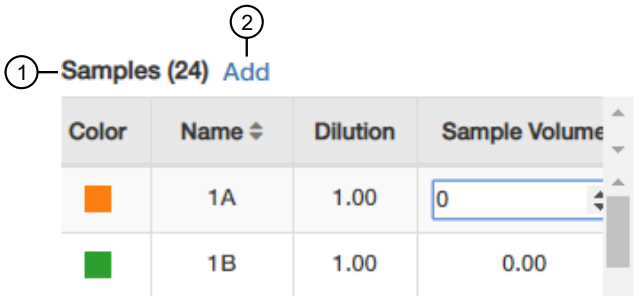


Figure 3 resDNASEQ™ template default cycling conditions

4. Click **Next**.

5. In the **Samples** pane of the **Setup** tab, enter the sample **Name** and **Dilution**. **Sample Volume** is not needed. Add more **Samples** if needed.

IMPORTANT! Do not change the **Targets**.



① **Samples** pane
② **Add**—adds more samples

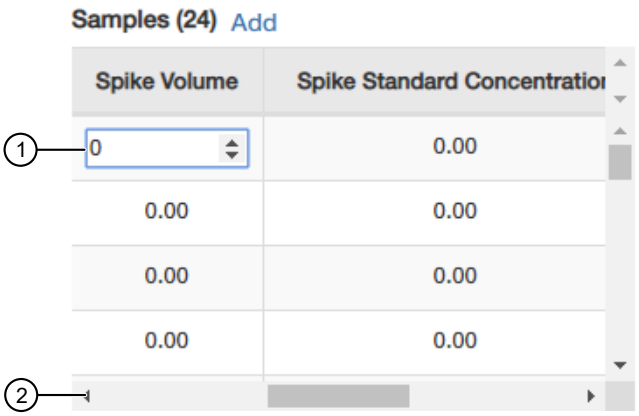
6. In the **Samples** pane of the **Setup** tab, scroll to the right, then enter the spike information.

   For more information about plate setup, see the *AccuSEQ™ Real-Time PCR Software v3.2 User Guide* (Pub. No. MAN0029199).

   - **Sample Volume**—not needed; leave as default (0).
   - **Spike Volume**—volume of DNA added to the PCR (set to 10).
   - **Spike Standard Concentration**—expected spike amount per reaction (for example, 10 pg/reaction or 30 copies/reaction).
   - **Reference**—the non-spiked sample; the mean quantity of reference is subtracted during % recovery calculation.
   - **Spike Input**—automatically calculated (double check if correct).

   Note: If not correct, ensure **Spike Volume** is set to 10 and **Spike Standard Concentration** is the expected pg spike per PCR reaction.

   - (Optional) **Comments**
   - **Protein Concentration**—Sample protein concentration (if Total DNA in pg DNA/mg Protein is needed).



① Textbox—type in the value, or use the up and down arrows
② Scroll bar—scroll to find the spike parameter

7. Click **Next**.
   The **Run** tab is displayed.

8. Experiments are auto-saved in the software. To save, exit the experiment. The software prompts you to save changes. Click **Yes**.

   Note: Clicking **Save As** creates a copy of the experiment.

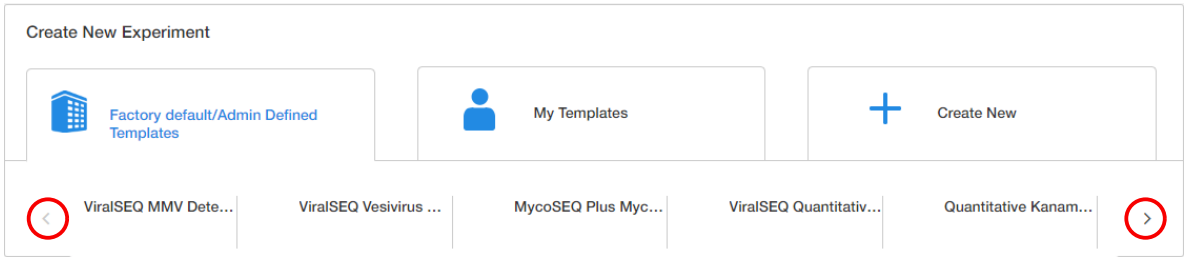
9. (*Optional*) Click **Print** to print the plate layout for use in preparing the reactions.

Create a Quantitative Sf9 and Baculovirus DNA experiment

Note: The **Quantitative Sf9 and Baculovirus DNA** template is available with AccuSEQ™ Software v3.2 or later.

1. In the **Home** screen, click the **Factory default/Admin Defined Template** tab, then select a **Quantitative Sf9 and Baculovirus DNA** template.

   To navigate to through the available templates click the arrows to either side of the available templates.



2. In the **Experiment Properties** pane of the **Setup** screen:

   a. (*Optional*) Change the system-generated name of the experiment.

   Note: Names must be unique. Deleted experiment names can not be reused.

   b. (*Optional*) Enter the plate barcode in the **Barcode** field, then add comments in the **Comments** field.

   Note: Comments are not editable post analysis.

   Default Quantitative Sf9 and Baculovirus DNA settings (cannot be changed)

   - **Experiment Type—Quantitation-Standard Curve**
   - **Chemistry—TaqMan™ Reagents**
   - **Ramp Speed—Standard - 2hrs**

   c. Click **Next**.

   Note: Experiment names cannot be changed after this step.

3. In the **qPCR Method** pane of the **Setup** screen, view the default volume and cycling conditions (cannot be changed).

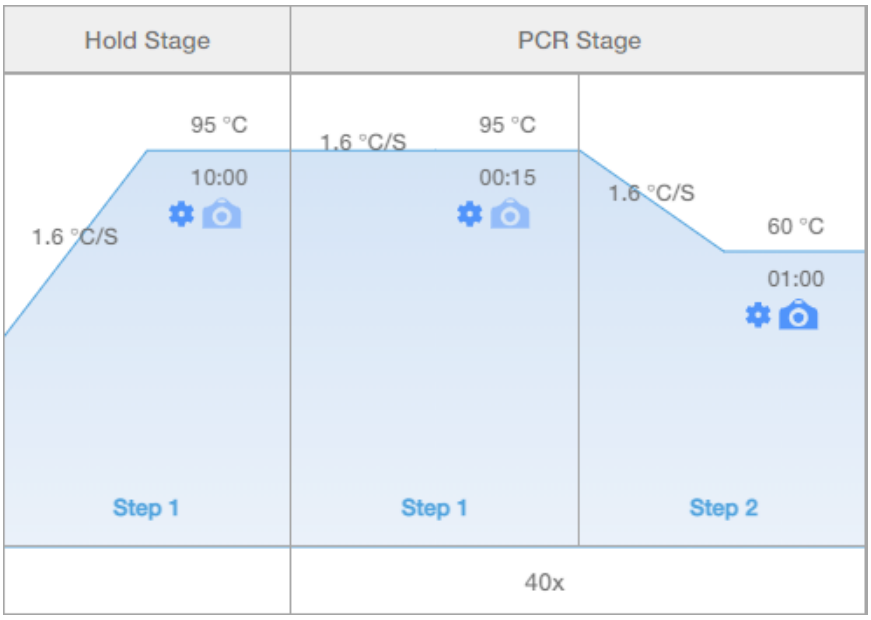
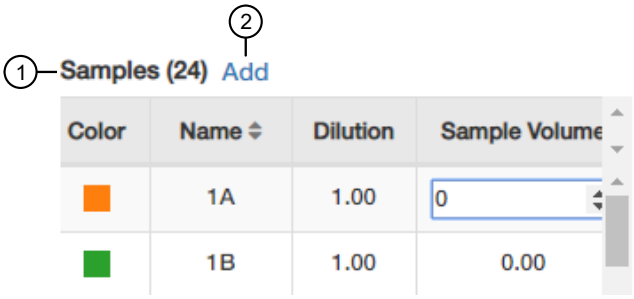


Figure 4 Quantitative Sf9 and Baculovirus DNA template default cycling conditions

4. Click **Next**.

5. In the **Samples** table in the **Plate Setup** pane, enter the sample name and the sample dilution. Sample volume is not needed. Click **Add** to add more samples if needed.

IMPORTANT! Do not change the **Targets**.



① **Samples** table
② **Add**—adds more samples

6. In the **Samples** table of the **Plate Setup** pane, scroll to the right, then enter the spike information.

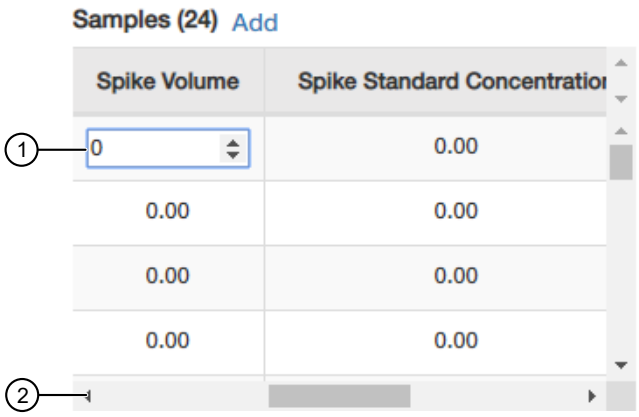
   For more information about plate setup, see *AccuSEQ™ Real-Time PCR Software v3.2 User Guide* (Pub. No. MAN0029199).

   - **Sample Volume**—not needed; leave as default (0).
   - **Spike Volume**—volume of DNA added to the PCR (set to 10).
   - **Spike Standard Concentration**—expected spike amount per reaction (for example, 10 pg).
   - **Reference**—the non-spiked sample; the mean quantity of reference is subtracted during % recovery calculation.
   - **Spike Input**—calculated (double check if correct).

   Note: If not correct, ensure **Spike Volume** value is set to 10 and **Spike Standard Concentration** value is the expected pg spike per PCR reaction.

   - (Optional) **Comments**
   - **Protein Concentration**—Drug substance protein concentration (if Total DNA in pg DNA/mg Protein is needed).

Note: The software calculates the pg of DNA in the initial sample and the percentage of recovered DNA



① Textbox—type in the value, or use the up and down arrows
② Scroll bar—scroll to find the spike parameter

Figure 5 Quantitative Sf9 and Baculovirus DNA template default sample plate layout

① Toolbar (in order: Undo, Redo, Copy, Paste, Delete, View)

② Select Item to highlight (Sample, Target, or Task).

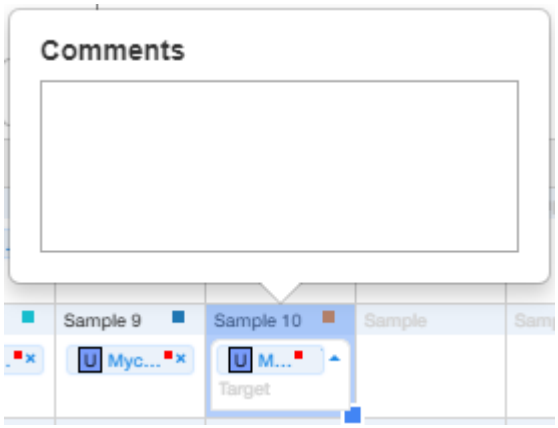
Note: If more samples are added, the Target displays as an S in the wells (experiment type–Quantitation).

③ Select Item. For example, Sample 1. Sample 1 replicates are highlighted.

Table 12 Plate well descriptions

Name	Description
SC1 to SC6	Standard curve dilutions in triplicate
Matrix	24 samples in triplicate
NTC	No template control in triplicate

7. (*Optional*) Double-click wells to add comments. Comments can also be added post-analysis.



8. Click **Next**.
   The **Run** screen is displayed.

9. Experiments are auto-saved in the software. To save, exit the experiment. The software prompts you to save changes. Click **Yes**.

   Note: Clicking **Save As** creates a copy of the experiment.

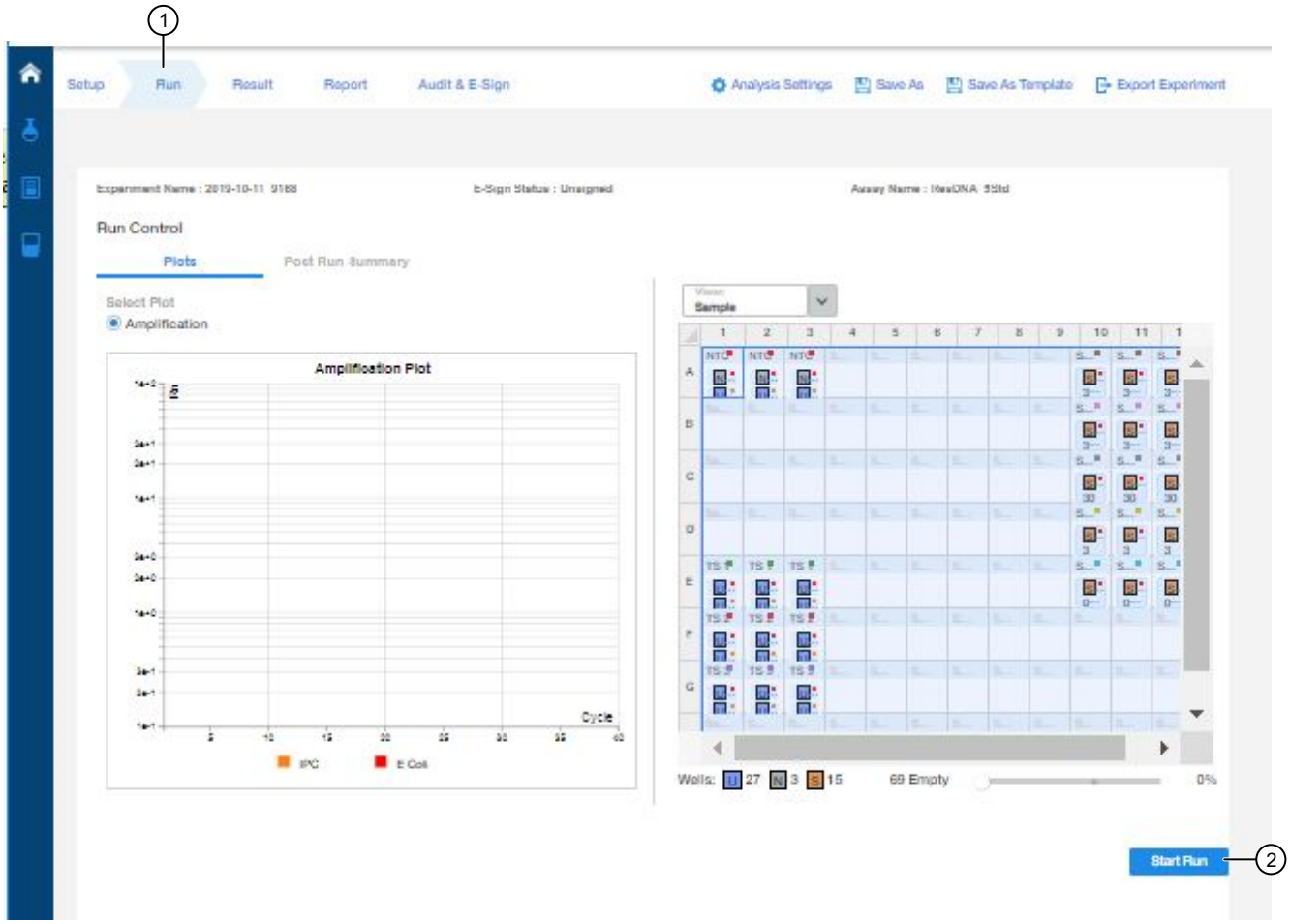
10. (*Optional*) Click **Print** to print the plate layout for use in preparing the reactions.

11. Assemble the PCR reactions using the instructions in this guide.

Start the run

Start the run in the AccuSEQ™ Software.

Option	Description
If the experiment is open	Click **Start Run**.
If the experiment is closed	a. Open the experiment. b. Click the **Run** tab. c. Click **Start Run**.



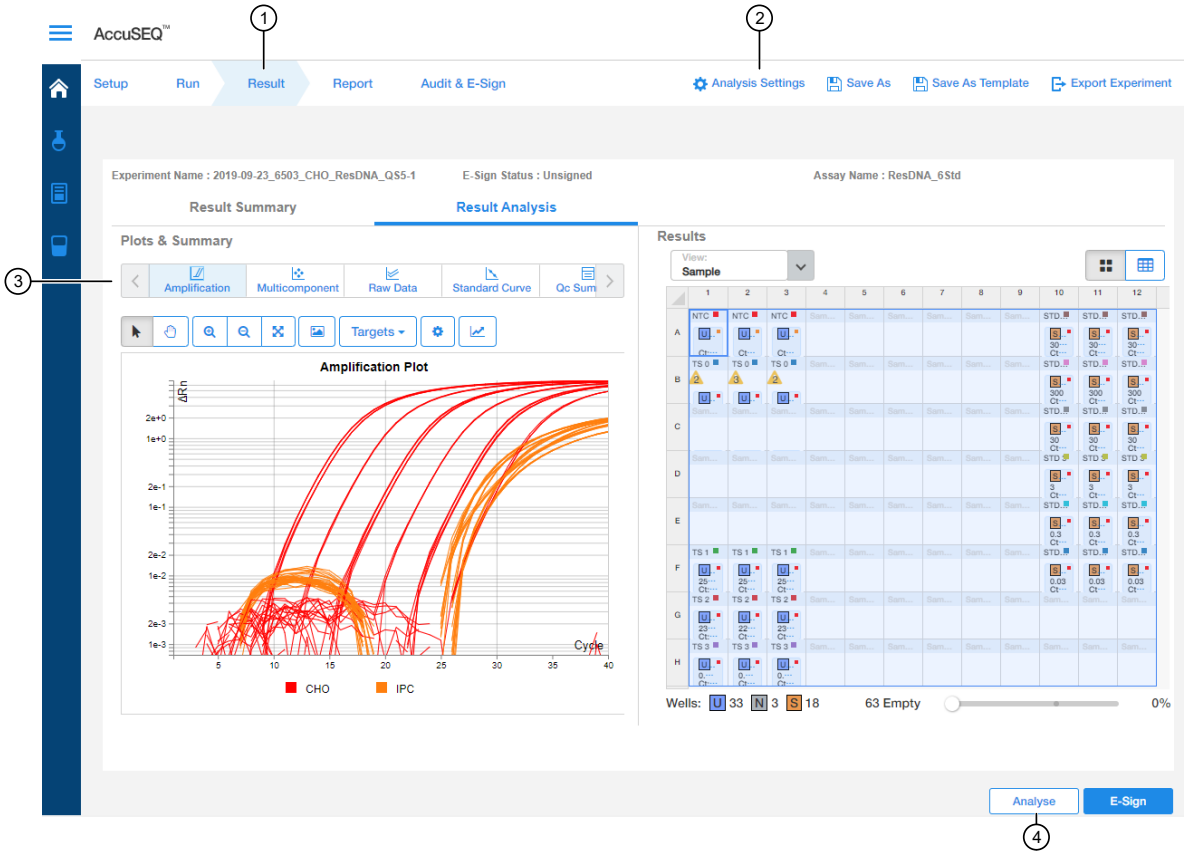
1 **Run** tab
2 **Start Run** button

A message stating **Run has been started successfully** is displayed when the run has started.

Analyze the results

After the qPCR run is finished, use the following general procedure to analyze the results. For more detailed instructions see the *AccuSEQ™ Real-Time PCR Software v3.2 User Guide* (Pub. No. MAN0029199).

1. In the AccuSEQ™ Real-Time PCR Software, open the experiment, then navigate to the **Result** tab.



① **Result** tab
② **Analysis Settings** button
③ Plot horizontal scrollbar
④ **Analyze** button

2. In the **Result Analysis** tab, review the Amplification Curve plots for amplification profiles in the controls, samples, and the standard curve.

3. In the **Result Analysis** tab, review the **QC Summary** for any flags in wells.

4. In the **Result Analysis** tab, review the **Standard Curve** plot. Verify the values for the Slope, Y-intercept, R^2, and Efficiency.

   Note: The **Standard Curve** efficiency should be between 90-110% and the R^2>0.99. If these criteria are not met, up to two points, not in the same triplicate, can be removed from the standard curve data, and the analysis repeated.

5. (*Optional*) Navigate to the **Report** tab to generate a report of the experiment, or to export results.

3 Plasmid DNA quantitation

Sample residual plasmid DNA is measured in copy number, not concentration. To manually convert the copy number output to a mass measurement, multiply the copy number given by the AccuSEQ™ Real-Time PCR Software to the average molecular weight of your plasmid.

Prepare the control DNA serial dilutions for the standard curve

Guidelines for standard dilutions

- Prepare the standard curve and the test samples in different areas of the lab.
- Use different sets of pipettors for test sample preparation and for standard curve preparation and aliquoting to avoid cross-contamination of test samples.
- Vortex each tube for 15-30 seconds to mix the contents thoroughly before each dilution step.
- Vortex for 15-30 seconds, then tap down standards, before adding the standards to the PCR plate (plasmid standards are more fragile than genomic DNA standards).

Prepare the control serial dilutions (Plasmid DNA - Kanamycin Resistance)

1. Label nonstick 1.5-mL microfuge tubes: **NTC, SD1**, **SD2**, **SD3**, **SD4**, **SD5**, and **SD6**.
   The dilution **SD6** will not be used for the standard curve. It will be used to confirm the limit of detection (LOD).
2. Add 70 µL of DNA Dilution Buffer to tube NTC. Put aside.
3. Add 990 µL of DNA Dilution Buffer to tube Dilution 1.
4. Add 180 µL of DNA Dilution Buffer to tubes SD1, SD2, SD3, SD4, and SD5.
5. Add 100 µL of DNA Dilution Buffer to tube SD6.
6. Remove the tube of KanR DNA control (3.0×10^7 copies/µL) from the freezer.
7. After the DNA thaws, vortex it thoroughly for 15-30 seconds, then tap gently to bring contents to the bottom of the tube. Do not centrifuge.
8. Perform the serial dilutions:
   a. Add 10 µL of the KanR DNA control (3.0×10^7 copies/µL) to the tube that is labeled Dilution 1, vortex thoroughly for 15-30 seconds, then tap gently to bring contents to the bottom of the tube.

b. Transfer 20 µL of the DNA from tube Dilution 1 to tube SD1, vortex thoroughly for 15-30 seconds, then tap gently to bring contents to the bottom of the tube..

c. Continue to transfer 20 µL of DNA from the previous dilution tube to the next dilution tube until you add DNA to tube SD5.

d. Transfer 100 µL of DNA from SD5 to SD6, then vortex thoroughly. Final dilutions are shown in the table. After each transfer, vortex thoroughly for 15-30 seconds, then tap gently to bring contents to the bottom of the tube. Do not centrifuge.

Serial dilution (SD) tube	Dilution	Concentration (copy number/ µL)	Copy number/ PCR reaction
Control	DNA control tube	3.0×10^7	N/A
Dilution 1	10 µL DNA control + 990 µL DDB	300,000	N/A
SD 1	20 µL Dilution 1 + 180 µL DDB	30,000	300,000
SD 2	20 µL SD1 + 180 µL DDB	3,000	30,000
SD 3	20 µL SD2 + 180 µL DDB	300	3,000
SD 4	20 µL SD3 + 180 µL DDB	30	300
SD 5	20 µL SD4 + 180 µL DDB	3	30
SD 6 (LOD)	100 µL SD5 + 100 µL DDB	1.5	15

9. Store the DNA dilution tubes:

Temperature	For use
4°C	≤2 days
–20°C	Dilution 1 ≤1 week
–20°C	Dilution 1 in single-use aliquots ≤6 months

Prepare the samples (Kanamycin resistance assay only)

Extract samples with the PrepSEQ™ Residual DNA Sample Preparation Kit (Cat. No. 4413686).

(Recommended) Use **Yeast tRNA**, supplied in the PrepSEQ™ Residual DNA Sample Preparation Kit or the resDNASEQ™ Quantitative Plasmid DNA - Kanamycin Resistance Gene Kit for the kanamycin resistance and assay.

1. Dilute the **Yeast tRNA**.

Table 13 Diluted Yeast tRNA

Component	Volume
Yeast tRNA (10mg/mL)	5 µL
PBS (1X), pH 7.2	245 µL
Total	**250 µL**

2. Add 5 µL **Diluted Yeast tRNA** to 370 µL of each test sample. This is sufficient for triplicate 100 µL extractions.

Prepare the PCR reaction mix

1. Determine the number of reactions needed for the controls and test samples that you will quantify.
2. Thaw all kit reagents completely at room temperature, thoroughly mix reagents, and briefly centrifuge.
3. Prepare a PCR reaction mix using the reagents and volumes shown in the table below.
   - Multiply the PCR reaction volume for one reaction (30 µL) by the number of reactions that you need to run.
   - Use 10% excess volume to compensate for pipetting losses.

Note: Use reagents from the same lot for all reactions.

Kit reagents	Volume for 1 30-µL reaction	Volume for 48 30-µL reactions (includes 10% overage)
Negative Control (Water)	2 µL	106 µL
10X DNA assay mix appropriate for the cell line being tested	3 µL	159 µL
TaqMan™ Environmental Master Mix 2.0	15 µL	795 µL
DNA template	10 µL	Add DNA template to each well separately, not as part of Master Mix
Total	**30 µL**	**1060 µL** of Master Mix

Prepare the PCR plate

Plate setup differs slightly for each AccuSEQ™ System. See your software user guide for specific instructions. Place samples, NTCs, and standards in different quadrants of the plate.

A factory default template is available for the Quantitative Kanamycin Assay in AccuSEQ™ Real-Time PCR Software v3.2 or later.

Figure 6 Quantitative Kanamycin template default sample plate layout in AccuSEQ™ Real-Time PCR Software v3.2 or later

① Toolbar (in order: Undo, Redo, Copy, Paste, Delete, View)

② Select Item to highlight (Sample, Target, or Task).

Note: If more samples are added, the Target displays as an S in the wells (experiment type–Quantitation).

③ Select Item. For example, Sample 1. Sample 1 replicates are highlighted.

Table 14 Plate well descriptions for Quantitative Kanamycin template in AccuSEQ™ Real-Time PCR Software v3.2 or later

Name	Description
SC1 to SC6	Standard curve dilutions in triplicate
X1 to X3, D1 to D4	24 samples in triplicate
NTC	No template control in triplicate

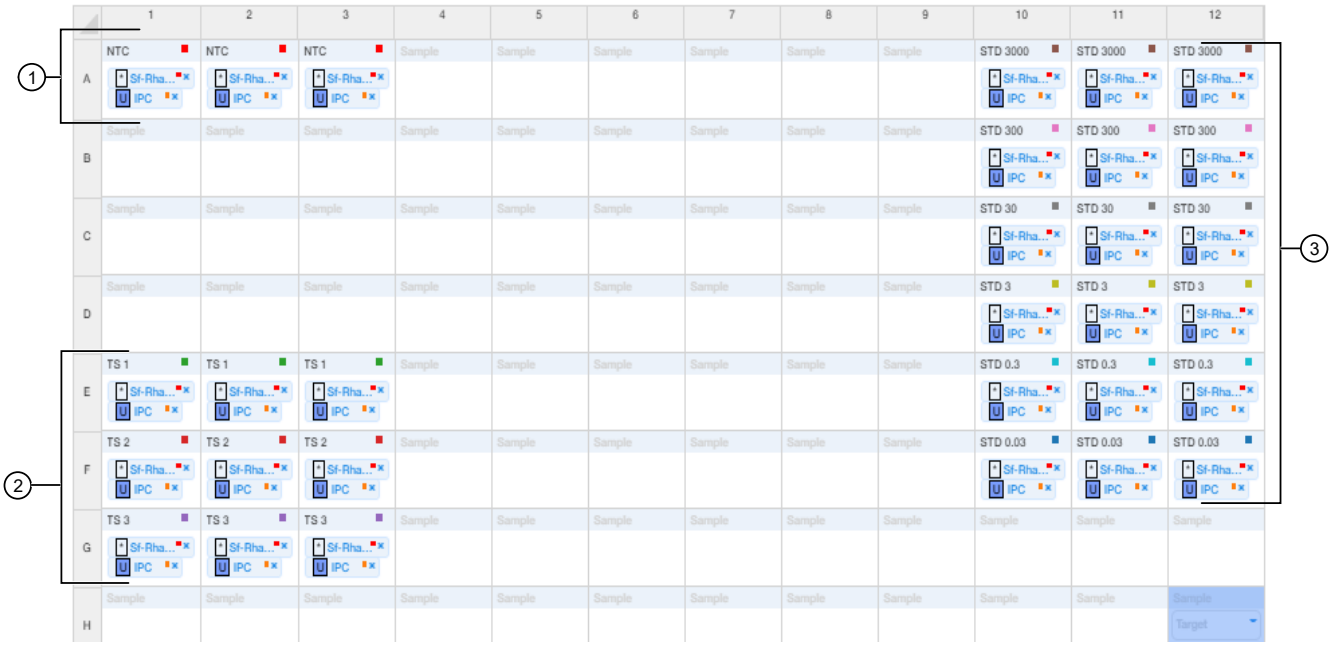


Figure 7 Default plate setup in the resDNA_5Std template in AccuSEQ™ Real-Time PCR Software v3.1

(1) No template controls
(2) Samples
(3) Standard curve

1. Add 20 µL PCR reaction mix to each well.
2. Add 10 µL of PCR NTC to the appropriate wells.
3. Add 10 µL each of sample DNA (with tRNA added) to the appropriate wells.

   IMPORTANT! Vortex samples for 15-30 seconds before adding the samples to the **PCR Plate**, then tap to bring contents to the bottom of the wells.

   If you prepared samples with the automated protocol, use a multichannel pipette to transfer the extracted sample.
4. Add 10 µL of standard dilutions to the appropriate wells.

   IMPORTANT! Vortex samples for 15-30 seconds before adding the samples to the **PCR Plate**, then tap to bring contents to the bottom of the wells.

   If you prepared samples with the automated protocol, use a multichannel pipette to transfer the extracted sample.
5. Use a film applicator to seal the plate with optical film, then briefly centrifuge with a miniplate centrifuge that is compatible with 96-well plates.

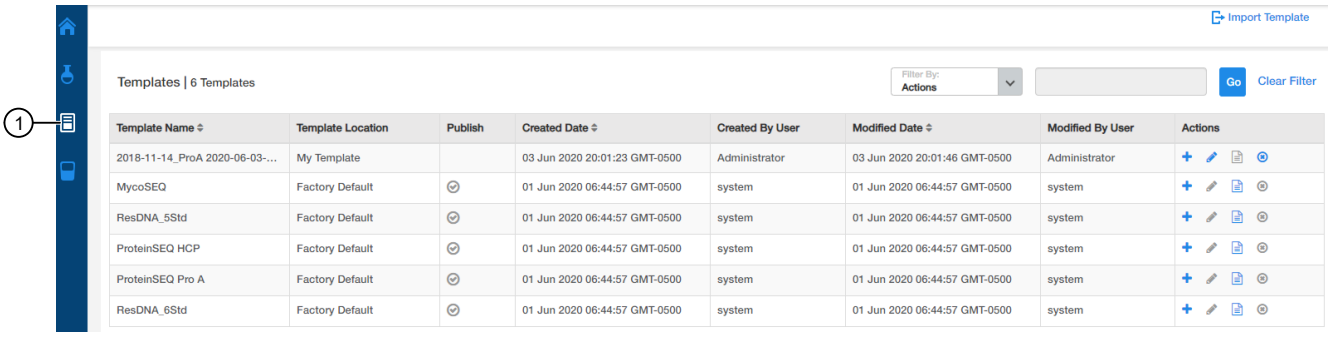
Setup, run, and analyze samples with AccuSEQ™ Software on the QuantStudio™ 5 Real-Time PCR Instrument

Create a resDNASEQ™ template (Plasmid DNA)

Plasmid DNA resDNASEQ™ assays in AccuSEQ™ Software v3.1 do not use a **Factory default/Admin Defined Template**. Create a new template before the first use of these assays.

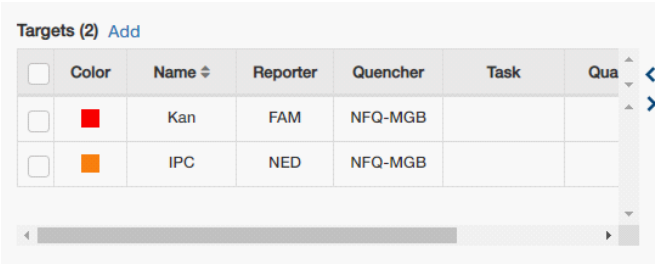
Note: AccuSEQ™ Software v3.2 or later has a **Factory default/Admin Defined Template** for the Quantitative Kanamycin Assay, see "Create a Quantitative Kanamycin experiment" on page 38.

1. In the **Home** screen, click **Templates** in the left navigation pane.



1 **Templates** icon

2. Click **Create New** next to the **ResDNA_5Std** factory default template.
3. Click **Next** to move to the **qPCR Method** screen.
4. Click **Next** to move to the **Plate Setup** screen.
5. In the **Plate Setup** screen, add the **Targets** and **Reporters**.



For the resDNASEQ™ Quantitative Plasmid DNA - Kanamycin Resistance Gene Kit, this is the FAM™ dye for the kanamycin (Kan) target and the NED™ dye for the IPC.

6. Click **Save as Template**.
7. Enter a **Template Name** and description, then select **Admin Defined** and **Locked**. Click **Save**.

   The template is saved, and can be accessed from **Templates** in the **Home** screen.

8. Click **Templates** in the **Home** screen, then open the new template.

   Note: The template must be saved prior to editing the **Analysis Settings**

9. Click **Analysis Settings**, then deselect **Default Settings**.

10. Enter new thresholds for the targets.

    Table 15 resDNASEQ™ Quantitative Plasmid DNA - Kanamycin Resistance Gene Kit

Target	Threshold
Kanamycin	0.04
IPC	0.02

11. Click Apply, then close the template.

12. In the **Home** screen, click **Templates** in the left navigation pane, then **Publish** the template. See *AccuSEQ™ Real-Time PCR Software v3.1 User Guide* (Pub. No. 100094287).
    The template is listed in the **Factory default/Admin Defined Templates**.

Create a resDNASEQ™ experiment (Plasmid DNA)

Plasmid DNA resDNASEQ™ assays in AccuSEQ™ Software v3.1 do not use a **Factory default/Admin Defined Template**. Create a new template before the first use of these assays.

Note: AccuSEQ™ Software v3.2 or later has a **Factory default/Admin Defined Template** for the Quantitative Kanamycin Assay, see “Create a Quantitative Kanamycin experiment” on page 38.

1. In the **Home** screen, click **Factory default/Admin Defined Templates**, then select the custom plasmid DNA **resDNASEQ** template created in “Create a resDNASEQ™ template (Plasmid DNA)” on page 34.

   Note: To create an experiment from an existing resDNASEQ™ experiment, see *AccuSEQ™ Real-Time PCR Software v3.1 User Guide* (Pub. No. 100094287).

2. In the **Experiment Properties** pane of the **Setup** tab:
   a. (*Optional*) Change the system-generated name of the experiment.

      Note: Names must be unique. Deleted experiment names can not be reused.

   b. (*Optional*) Enter the plate **Barcode**, then add **Comments**.

      Note: Comments are not editable post analysis.

      Default resDNASEQ™ settings (cannot be changed)

      - **Experiment Type** is **Quantitation-Standard Curve**
      - **Chemistry** is **TaqMan™ Reagents**
      - **Ramp Speed** is **Standard - 2hrs**

c. Click **Next**.

Note: Experiment names cannot be changed after this step.

3. In the **qPCR Method** pane of the **Setup** tab, view the default volume and cycling conditions (cannot be changed).

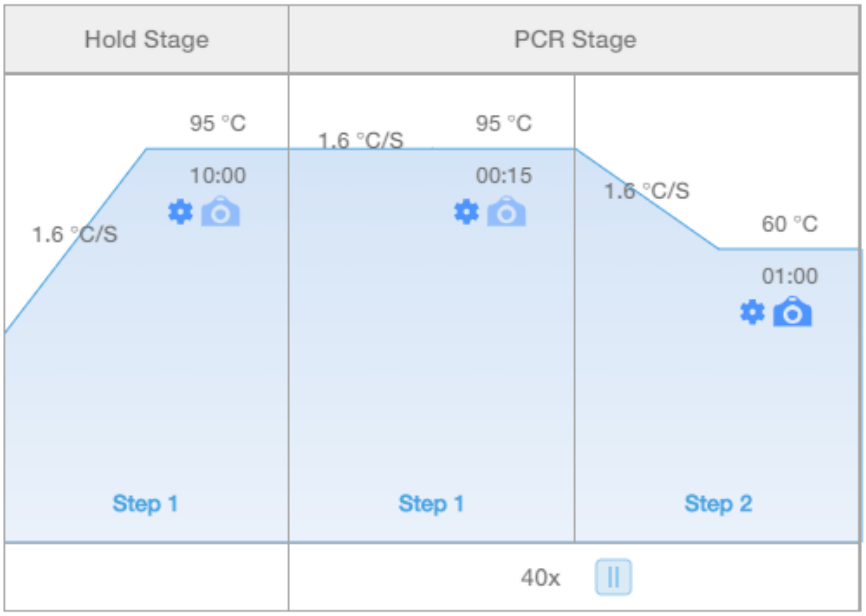
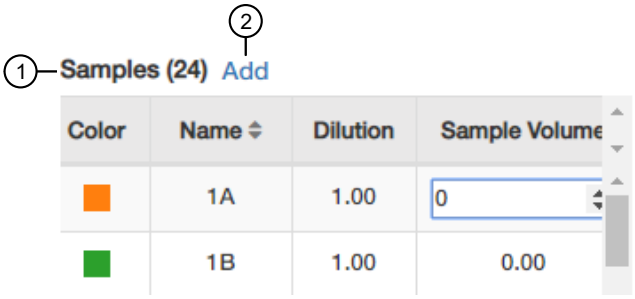


Figure 8 resDNASEQ™ template default cycling conditions

4. Click **Next**.

5. In the **Samples** pane of the **Setup** tab, enter the sample **Name** and **Dilution**. **Sample Volume** is not needed. Add more **Samples** if needed.

IMPORTANT! Do not change the **Targets**.



① **Samples** pane
② **Add**—adds more samples

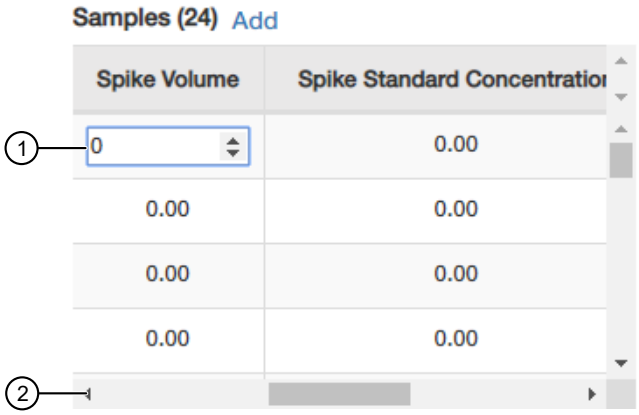
6. In the **Samples** pane of the **Setup** tab, scroll to the right, then enter the spike information.

   For more information about plate setup, see *AccuSEQ™ Real-Time PCR Software v3.1 User Guide* (Pub. No. 100094287).

   - **Sample Volume**—not needed; leave as default (0).
   - **Spike Volume**—volume of DNA added to the PCR (set to 10).
   - **Spike Standard Concentration**—expected spike amount per reaction.
   - **Reference**—the non-spiked sample; the mean quantity of reference is subtracted during % recovery calculation.
   - **Spike Input**—automatically calculated (double check if correct).

   Note: If not correct, ensure **Spike Volume** is set to 10 and **Spike Standard Concentration** is the expected pg spike per PCR reaction.

   - (Optional) **Comments**
   - **Protein Concentration**—Drug substance protein concentration (if Total DNA in pg DNA/mg Protein is needed).



① Textbox—type in the value, or use the up and down arrows

② Scroll bar—scroll to find the spike parameter

For more information about plate setup, see *AccuSEQ™ Real-Time PCR Software v3.1 User Guide* (Pub. No. 100094287).

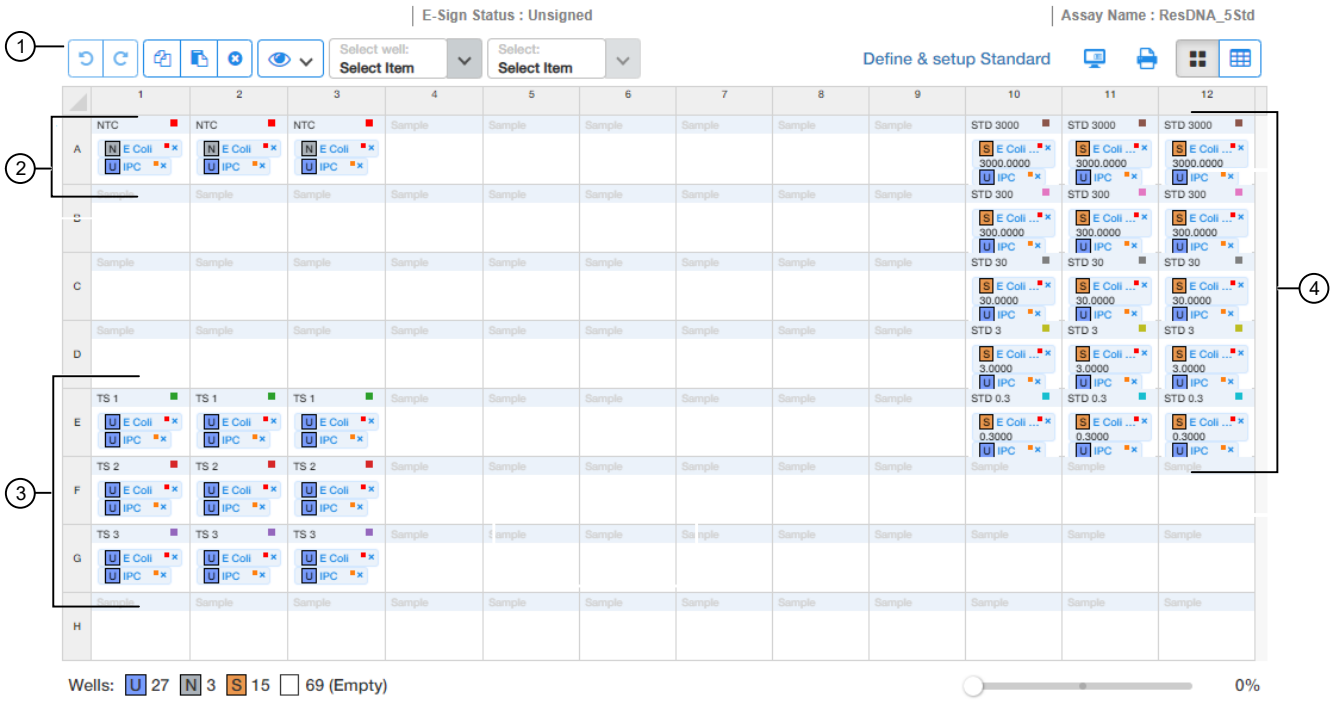


Figure 9 resDNASEQ™_5Std template default sample plate layout

① Toolbar (in order: Undo, Redo, Copy, Paste, Delete, View)
② 3 No Template Control (NTC) samples
③ 3 default **Samples**
④ Standard curve dilutions (S) in triplicate

7. Click **Next**.
   The **Run** tab is displayed.

8. Experiments are auto-saved in the software. To save, exit the experiment. The software prompts you to save changes. Click **Yes**.

   Note: Clicking **Save As** will create a copy of the experiment.

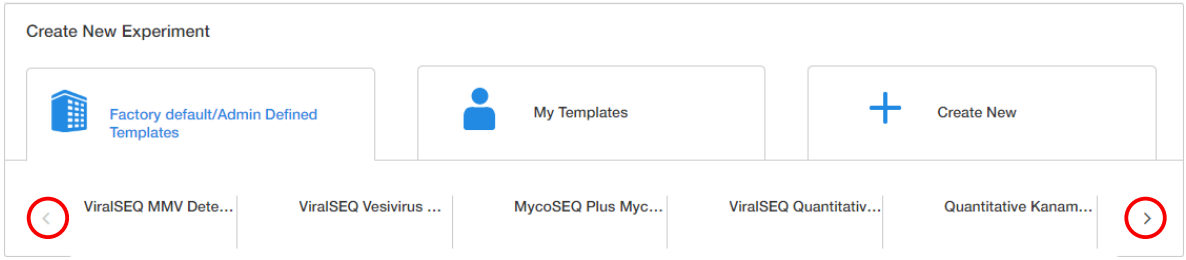
9. (*Optional*) Click **Print** to print the plate layout for use in preparing the reactions.

10. Assemble the PCR reactions using the manufacturer's instructions for the reagents and following the plate layout set up in the template.

Create a Quantitative Kanamycin experiment

Note: The **Quantitative Kanamycin** template is available with AccuSEQ™ Software v3.2 or later.

1. In the **Home** screen, click the **Factory default/Admin Defined Template** tab, then select a **Quantitative Kanamycin** template.
   To navigate to through the available templates click the arrows to either side of the available templates.



2. In the **Experiment Properties** pane of the **Setup** screen:
   a. (*Optional*) Change the system-generated name of the experiment.

   Note: Names must be unique. Deleted experiment names can not be reused.

   b. (*Optional*) Enter the plate barcode in the **Barcode** field, then add comments in the **Comments** field.

   Note: Comments are not editable post analysis.

   Default Quantitative Kanamycin settings (cannot be changed)

   - **Experiment Type—Quantitation-Standard Curve**
   - **Chemistry—TaqMan™ Reagents**
   - **Ramp Speed—Standard - 2hrs**

   c. Click **Next**.

   Note: Experiment names cannot be changed after this step.

3. In the **qPCR Method** pane of the **Setup** screen, view the default volume and cycling conditions (cannot be changed).

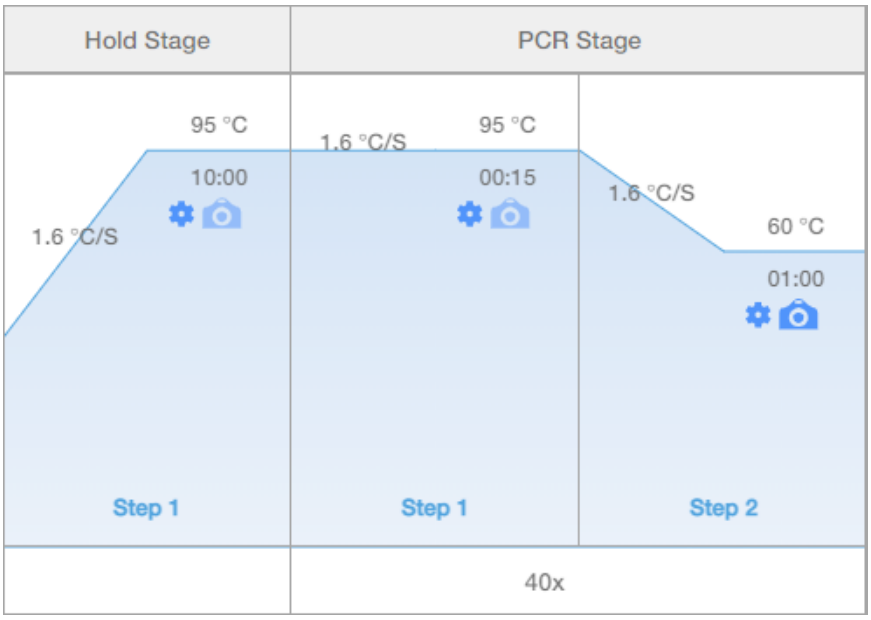
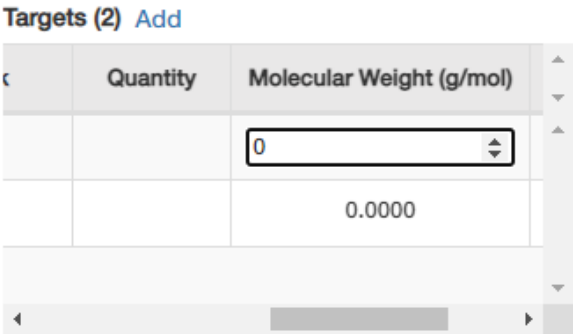


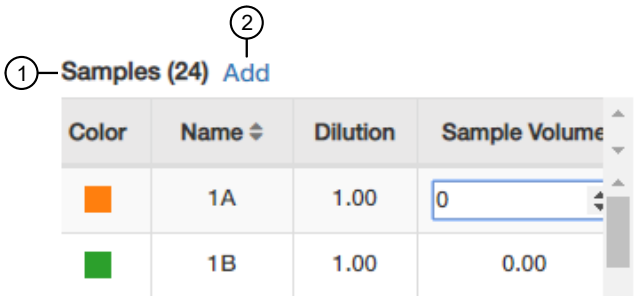
Figure 10 Quantitative Kanamycin template default cycling conditions

4. Click **Next**.

5. In the **Targets** table of the **Plate Setup** pane, enter the molecular weight (g/mol) of the targets.



6. In the **Samples** table in the **Plate Setup** pane, enter the sample name and the sample dilution. Sample volume is not needed. Click **Add** to add more samples if needed.



① **Samples** pane
② **Add**—adds more samples

7. In the **Samples** table in the **Plate Setup** pane, scroll to the right, then enter the spike information.

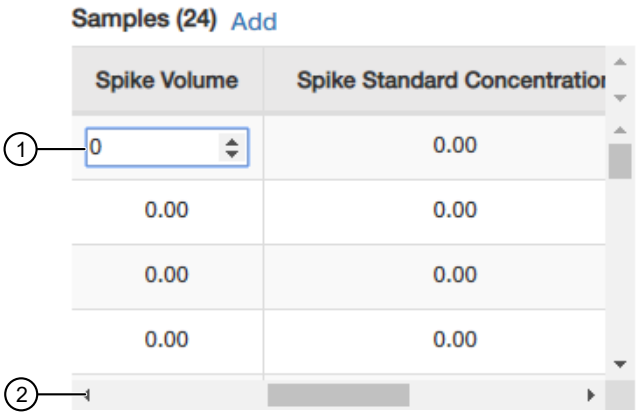
   For more information about plate setup, see *AccuSEQ™ Real-Time PCR Software v3.2 User Guide* (Pub. No. MAN0029199).

   - **Sample Volume**—not needed; leave as default (0).
   - **Spike Volume**—volume of DNA added to the PCR (set to 10).
   - **Spike Standard Concentration**—expected spike amount per reaction (for example, 10 copies).
   - **Reference**—the non-spiked sample; the mean quantity of reference is subtracted during % recovery calculation.
   - **Spike Input**—calculated (double check if correct).

     Note: If not correct, ensure **Spike Volume** is set to 10 and **Spike Standard Concentration** is the expected copies spike per PCR reaction.

   - **Protein Concentration**—Drug substance protein concentration (if total DNA in copies DNA/mg protein is needed).
   - (Optional) **Comments**

Note: The software calculates the mass (g) and % recovery.



① Textbox—type in the value, or use the up and down arrows
② Scroll bar—scroll to find the spike parameter

Figure 11 Quantitative Kanamycin template default sample plate layout

① Toolbar (in order: Undo, Redo, Copy, Paste, Delete, View)
② Select Item to highlight (Sample, Target, or Task).

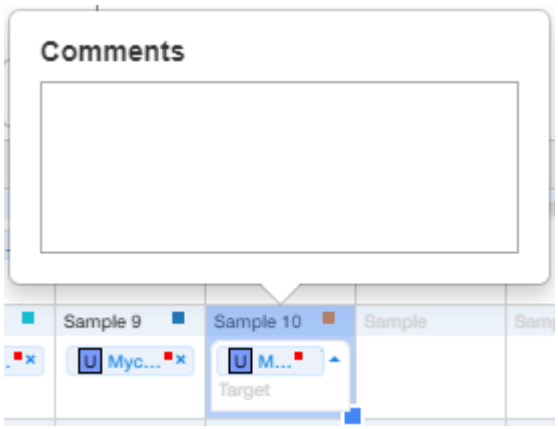
Note: If more samples are added, the Target displays as an S in the wells (experiment type–Quantitation).

③ Select Item. For example, Sample 1. Sample 1 replicates are highlighted.

Table 16 Plate well descriptions

Name	Description
SC1 to SC6	Standard curve dilutions in triplicate
X1 to X3, D1 to D4	24 samples in triplicate
NTC	No template control in triplicate

8. (*Optional*) Double-click wells to add comments. Comments can also be added post-analysis.



9. Click **Next**.
   The **Run** screen is displayed.

10. Experiments are auto-saved in the software. To save, exit the experiment. The software prompts you to save changes. Click **Yes**.

Note: Clicking **Save As** creates a copy of the experiment.

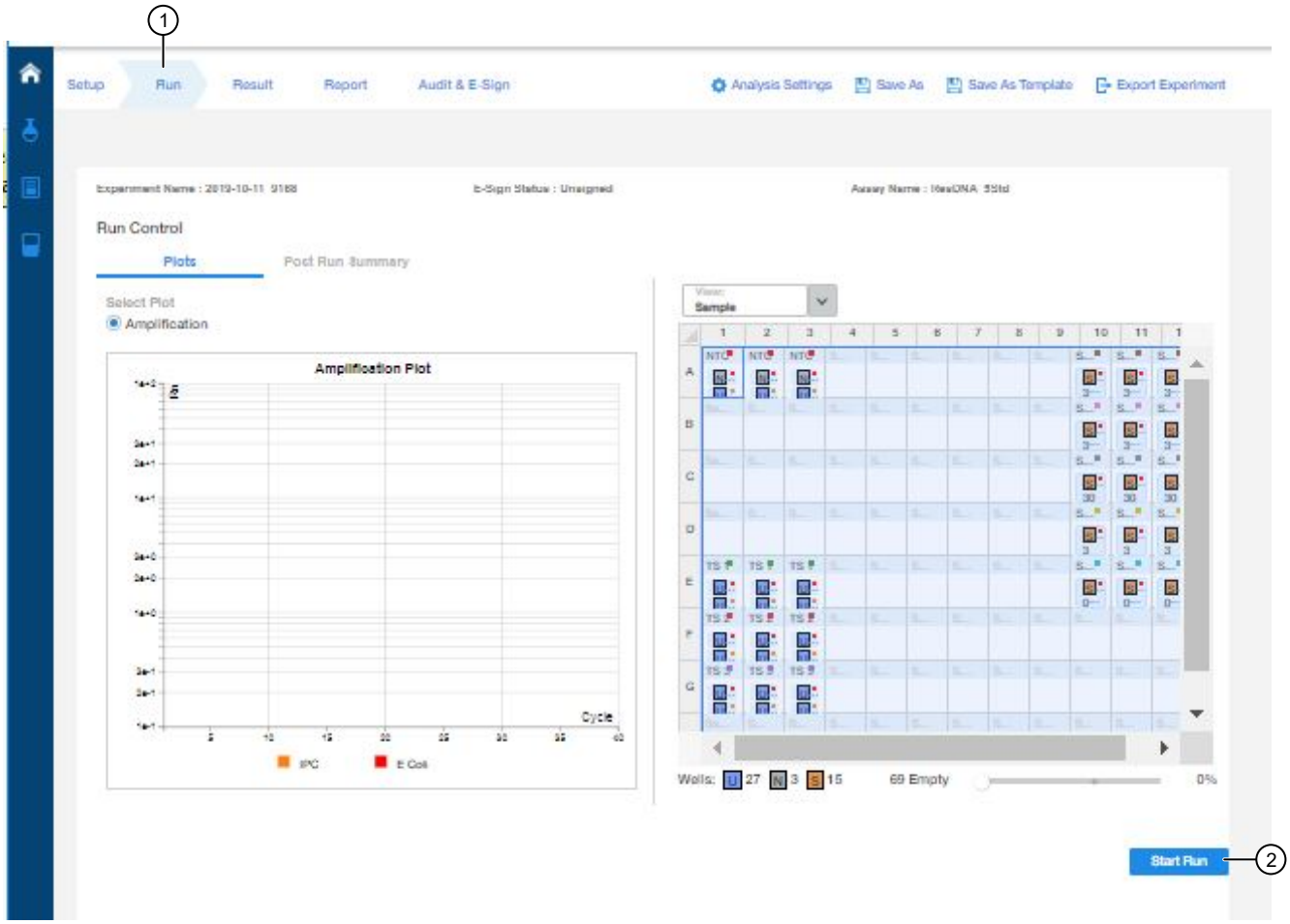
11. (*Optional*) Click **Print** to print the plate layout for use in preparing the reactions.

12. Assemble the PCR reactions using the instructions of the manufacturer for the reagents and following the plate layout set up in the template.

Start the run

Start the run in the AccuSEQ™ Software.

Option	Description
If the experiment is open	Click **Start Run**.
If the experiment is closed	a. Open the experiment. b. Click the **Run** tab. c. Click **Start Run**.



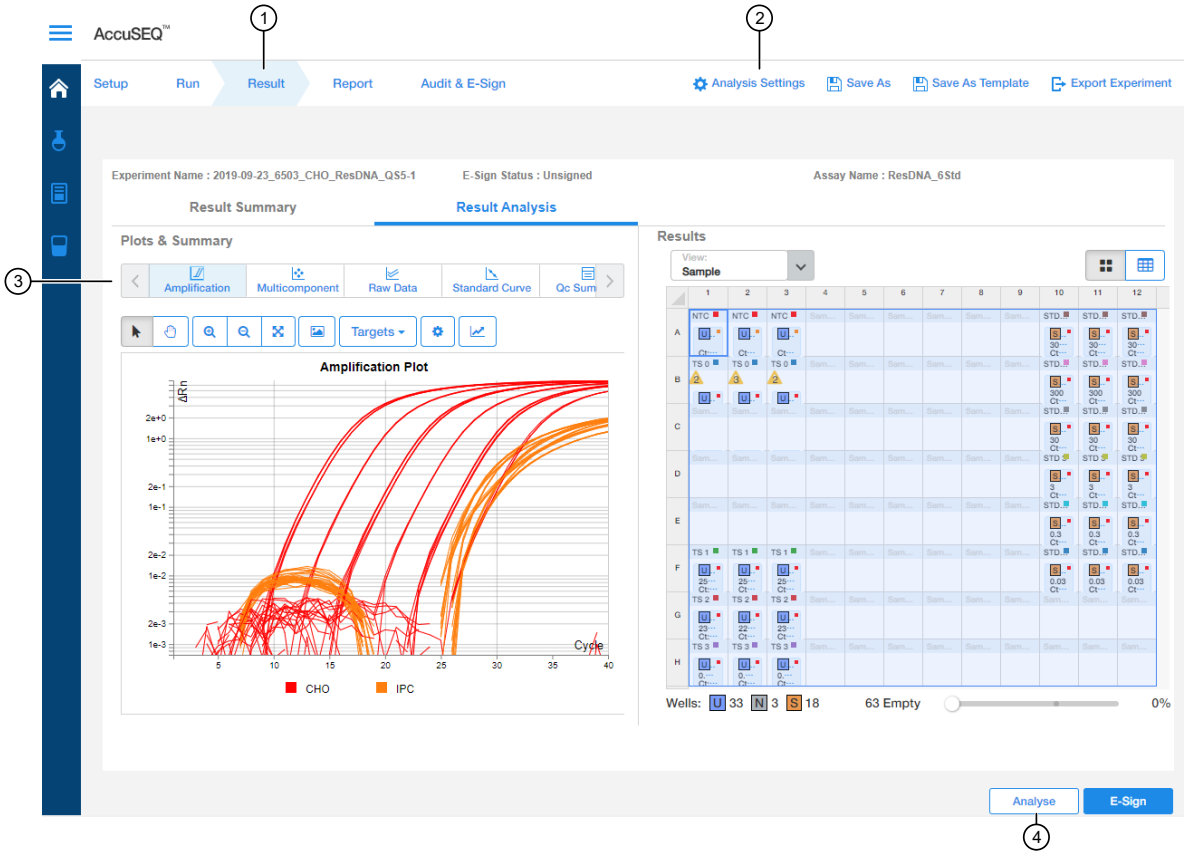
1 **Run** tab
2 **Start Run** button

A message stating **Run has been started successfully** is displayed when the run has started.

Analyze the results

After the qPCR run is finished, use the following general procedure to analyze the results. For more detailed instructions see the *AccuSEQ™ Real-Time PCR Software v3.2 User Guide* (Pub. No. MAN0029199).

1. In the AccuSEQ™ Real-Time PCR Software, open the experiment, then navigate to the **Result** tab.



① **Result** tab
② **Analysis Settings**
③ Plot horizontal scrollbar
④ **Analyze** button

2. In the **Result Analysis** tab, review the Amplification Curve plots for amplification profiles in the controls, samples, and the standard curve. Ensure that auto threshold is selected.

Note: For AccuSEQ™ Real-Time PCR Software v3.1 Plasmid- KanR assay only, ensure that the threshold was set to 0.04 for Kanamycin target and 0.02 for IPC.

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3. In the **Result Analysis** tab, review the **QC Summary** for any flags in wells.

4. In the **Result Analysis** tab, review the **Standard Curve** plot. Verify the values for the Slope, Y-intercept, R^2, and Efficiency.

   Note: The **Standard Curve** efficiency should be between 90-110% and the R^2>0.99. If these criteria are not met, up to two points, not in the same triplicate, can be removed from the standard curve data, and the analysis repeated.

5. (*Optional*) Navigate to the **Report** tab to generate a report of the experiment, or to export results.

6. (*Optional* for AccuSEQ™ Real-Time PCR Software v3.1) To manually convert the copy number output to a mass measurement, multiply the copy number given by the AccuSEQ™ Real-Time PCR Software to the average molecular weight of the plasmid, then divide by the Avogadro constant.

$$Mass\ (g) = \frac{Copy\ number\ x\ Molecular\ weight\ (gmol^{-1})}{6.0221\ X\ 10^{23}\ mol^{-1}}$$

   Note: AccuSEQ™ Real-Time PCR Software v3.2 and later calculates mass (g) and % Recovery for the **Quantitative Kanamycin** template in the **Results** table.

A

Troubleshooting

Observation	Possible cause	Action
Slope for the standard curve is outside the typical range, or R_2 value is significantly less than 0.99	When applying detectors for standards, the Task and Quantity were applied to the wrong detector. *or* The incorrect Quantity was entered. *or* Adjust baseline settings. *or* Poor standard curve preparation technique (forgot to mix, inaccurate pipetting).	**1.** In the SDS software, from the plate document, double-click a well containing a DNA standard to view the Well Inspector. **2.** Ensure that the correct Task and Quantity are applied to the correct detector, then reanalyze. **3.** Compare std curve statistics using autobaseline or manual baseline. The upper limit of the manual baseline setting must be 2 cycles before uptick in amplification. Verify in Rn vs C_t linear view.
Δ Rn and C_t values are inconsistent with replicates	Evaporation of reaction mixture from some wells occurred because the optical adhesive cover was not correctly sealed to the reaction plate or due to over-drying the eluates in PrepSEQ™.	**1.** Select the **Component** tab. Confirm that affected wells generated significantly less fluorescence than unaffected replicates. **2.** Check the amount of solution in each well of the reaction plate. Confirm that the wells affected by evaporation contained less solution than unaffected wells, and corresponded with the inconsistent results. **3.** For subsequent runs, ensure that the optical adhesive cover is correctly sealed to the reaction plate.
	Incorrect volume of PCR reaction mix was added to some reactions.	**1.** Select the **Component** tab. Confirm that affected wells generated significantly less fluorescence than unaffected replicates. **2.** Select the **Spectra** tab. Confirm that the wells with the incorrect volume of PCR reaction mix generated significantly different amounts of fluorescence than the unaffected wells. **3.** For subsequent runs, ensure the correct volume of PCR reaction mix.
Jagged amplification plots	Weak lamp or incorrect replacement.	Replace the lamp or ensure that the existing replacement is correct.

(continued)

Observation	Possible cause	Action
No defined amplification plots	An incorrect detector was selected on the amplification plot. *or* An incorrect detector was applied to the reactions when setting up the plate document.	**1.** Ensure that the correct detector was selected on the amplification plot. **2.** If the correct detector was not selected, then in the plate document, double-click a well to view the Well Inspector, verify that the detector settings are correct, and reanalyze.
Abnormal ΔRn values or negative ΔR_n values	Incorrect passive reference was selected when setting up the plate document.	**1.** From the plate document, double-click a well to view the Well Inspector. **2.** Ensure that ROX™ dye was selected as the Passive Reference.
Standard curve for plasmid DNA assays is outside of the 90–110% efficiency range	Incomplete vortexing of low level standards.	Repeat reactions, ensuring that samples and standards are vortexed for 15-30 seconds.
Wide variance of C_t values of plasmid DNA samples	Incomplete vortexing of samples.	

B

Use the kit with the 7500 Fast Real-Time PCR Instrument and AccuSEQ™ software v2.x

Required materials not supplied

Unless otherwise indicated, all materials are available through **thermofisher.com**. "MLS" indicates that the material is available from **fisherscientific.com** or another major laboratory supplier.

Item	Source
Instrument	
7500 Fast Real-Time PCR System with AccuSEQ™ software v2.x	Contact your local sales representative
Generic consumables	
Disposable gloves	MLS
Aerosol-resistant pipette tips	MLS
Pipettors: • Positive-displacement • Air-displacement • Multichannel	MLS
Consumables for the 7500 Fast Real-Time PCR System	
MicroAmp™ Fast Optical 96-Well Reaction Plate with Barcode, 0.1 mL, 20 plates; for use with 7500 Fast Real-Time PCR System	4346906
MicroAmp™ Optical 96-Well Reaction Plate with Barcode & Optical Adhesive Films, 100 plates with covers	4314320
MicroAmp™ Optical 8-Cap Strips, 300 strips	4323032
MicroAmp™ Optical Adhesive Film Kit, 20 covers, 1 applicator, 1 optical cover compression pad	4313663
MicroAmp™ Optical Adhesive Film	4360954

Create a plate layout in the AccuSEQ™ v2.0 or v2.1 software

Note: AccuSEQ™ Software v2.2 or later has factory default **SEQ Experiments** for the Quantitative Sf9 and Baculovirus DNA and Quantitative Kanamycin Assays, see "Create a Quantitative Sf9 and Baculovirus DNA Assay experiment" on page 53 and "Create a Quantitative Kanamycin Assay experiment" on page 56.

Use this workflow for Quantitative Sf9 and Baculovirus DNA and Quantitative Kanamycin Assays in AccuSEQ™ Software v2.0 or v2.1.

1. In the home screen, select **Create Custom Experiment**.
2. In the **Experiment name** field, enter a unique name for the experiment.
3. Specify experiment information.
   a. Select experiment type **Quantitation – Standard Curve**.
   b. Select reagents **TaqMan™ Reagents**.
   c. Select ramp speed **Standard**.
4. In the **Plate Setup** screen, select the **Define Targets and Samples** tab.
5. Specify target information.
   a. Click **Add New Target**.
   b. Enter a host cell name in the target name field.
   c. Select reporter **FAM™** dye and quencher **NFQ_MGB** dye.

      Note: For the Sf9 and Baculovirus assay, select reporter VIC™ dye for the Sf9 target and quencher **NFQ_MGB** dye. Add an additional target, and select reporter FAM™ dye for the baculovirus target and quencher **NFQ_MGB** dye.

   d. Select a color for this target.
6. Specify IPC target information.
   a. Click **Add New Target**.
   b. Enter **IPC** in the target name field.
   c. Select reporter **VIC™** dye and quencher **NFQ_MGB** dye.

      Note: For the Sf9 and Baculovirus assay and the Plasmid DNA - Kanamycin Resistance assay, select the reporter NED™ dye and quencher NFQ_MGB dye.

   d. Select a color for this target.
7. Define new samples.
   a. Click **Add New Sample**.

b. In **Sample Name**, add the names of the samples you want to define.

c. Click **Next**, or select the **Assign Targets and Samples** tab.

8. In the **Assign Targets and Samples** tab, define new targets.
   a. Follow the instructions in the top of the tab to set up the standards, unknowns, and negative controls.

   b. Click **Define and Set Up Standards** to open the **Define and Set Up Standards** dialog box to enter the appropriate settings and define the standard curve. When defined, click **Apply** and **Close**. The new standard curve is applied to the plate layout screen.

9. Assign the IPC to the standard curve wells.

10. In the **Run Method** screen, in the **Graphical View** tab.
    a. In **Reaction Volume Per Well**, enter **30 µL**

    b. Right-click the left column named **Holding Stage 1** and select **Delete Selected**. This 50°C hold stage is not needed.

11. Click the **Analysis** button in the left panel. In the **Analysis Settings** window on the right, set the default settings.
    a. On the **Ct Settings** tab, click **Edit Default Settings**. Then set **Threshold** to 0.2, set to **Automatic Baseline**, and then click **Save Changes**.

    Note: For CHO samples, a manual baseline of 3–12 is more appropriate.
    For the Plasmid DNA - Kanamycin Resistance assay, do not use the default settings. Set the threshold to 0.04 for KanR and 0.02 for IPC.

    b. Select (highlight) both targets.

    c. In the right-hand window, select **Use Default Settings**.

    d. Click **Apply Analysis Settings**.

Note: You can analyze the assays using Automatic or Manual Baseline, use the setting that yields the best standard curve. For CHO, the upper limit threshold for manual baseline analysis is 12.

12. Select **File ▸ Save as**, ensure that the file is named “resDNA_Template”, then select **Save as a template file** in the dropdown list and close the template plate document.

Note: You can reuse the plate template document to run an assay by opening the template file and choosing **Save As** to save the file with the experiment name.

Run the plate

Note: AccuSEQ™ Software v2.2 or later has factory default **SEQ Experiments** for the Quantitative Sf9 and Baculovirus DNA and Quantitative Kanamycin Assays, see "Create a Quantitative Sf9 and Baculovirus DNA Assay experiment" on page 53 and "Create a Quantitative Kanamycin Assay experiment" on page 56.

Use this workflow for Quantitative Sf9 and Baculovirus DNA and Quantitative Kanamycin Assays in AccuSEQ™ Software v2.0 or v2.1.

1. In the toolbar, select **File ▸ Open**, navigate to the **resDNA_Template** file, then click **Open**.
2. In the **Experiment Name** field, enter the appropriate experiment name, then click **Finish**.
3. Make any necessary changes to the test sample labels.
   - **Sample Volume**—not needed; leave as default (0).
   - **Spike Volume**—volume of DNA added to the PCR (set to 10).
   - **Spike Standard Concentration**—expected spike amount per reaction (for example, 10 pg).
   - **Reference**—the non-spiked sample; the mean quantity of reference is subtracted during % recovery calculation.
   - **Spike Input**—automatically calculated (double check if correct).

   Note: If not correct, ensure **Spike Volume** is set to 10 and **Spike Standard Concentration** is the expected pg spike per PCR reaction.
4. Select **Save As** to save the new experiment as an EDS experiment file with the same name as entered in the **Experiment Name** field.
5. Load the plate into the instrument.
6. Click **Start Run**.
7. Select a run screen (**Amplification** plot, **Temperature** plot, or **Run method**) to monitor the progress of the run.

Create a Quantitative Sf9 and Baculovirus DNA Assay experiment

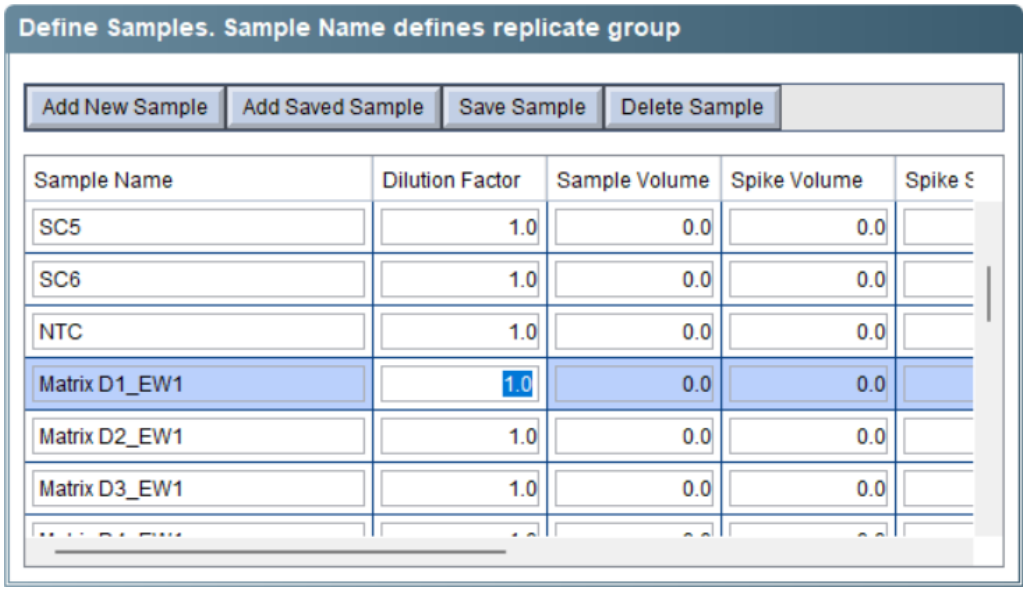
Note: The **Quantitative Sf9 and Baculovirus DNA Assay** SEQ experiment is available with AccuSEQ™ Software v2.2 or later.

1. Click **Create SEQ Experiment** in the home screen.
2. Select **Quantitative Sf9 and Baculovirus DNA Assay** from the assay list, then click **Next** . The **Plate Setup** screen opens.
3. In the **Experiment Menu** navigation bar, click **Experiment Properties**.
4. Enter the experiment name.

   Note: The experiment name can be up to 100 letters and numbers. Spaces are not allowed.

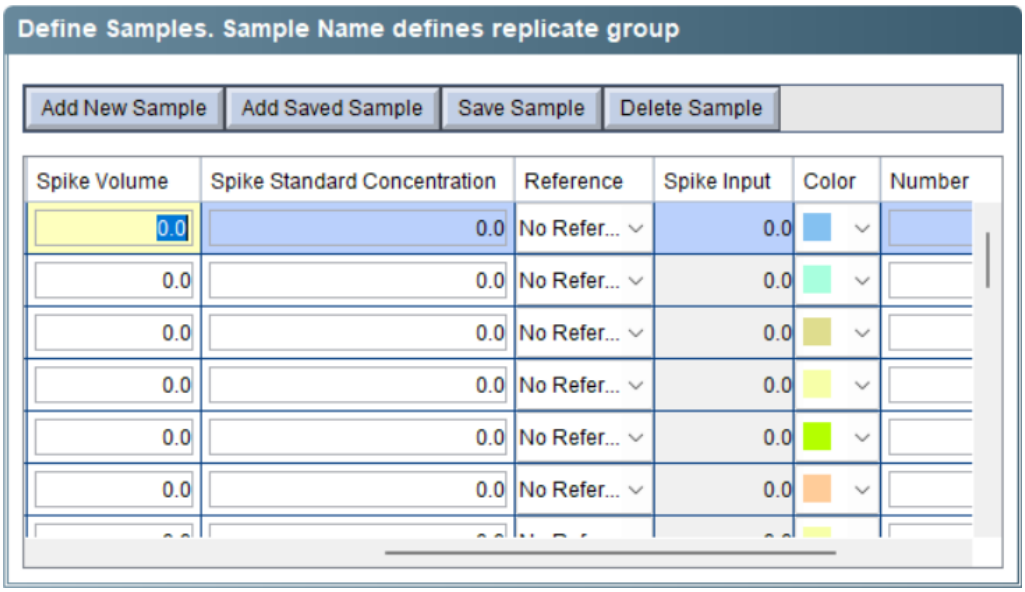
5. *(Optional)* Enter a plate barcode and comments.
6. In the **Experiment Menu** navigation bar, click **Plate Setup**.
7. In the **Define Samples** pane of the **Define Targets and Samples** tab of the **Plate Setup** pane, enter the sample name and the sample dilution. Sample volume is not needed. Click **Add New Sample** to add more samples if needed.

   IMPORTANT! Do not change the **Targets**.



For information about adding and defining samples, see *AccuSEQ™ Real-Time PCR Software v2.2 User Guide* (Pub. No. MAN0029201)

8. In the **Define Samples** pane in the **Define Targets and Samples** tab, scroll to the right, then enter the spike information.



- **Sample Volume**—not needed; leave as default (0).
- **Spike Volume**—volume of DNA added to the PCR (set to 10).
- **Spike Standard Concentration**—expected spike amount per reaction (for example, 10 pg).
- **Reference**—the non-spiked sample; the mean quantity of reference is subtracted during % recovery calculation.
- **Spike Input**—calculated (double check if correct).

  Note: If not correct, ensure **Spike Volume** value is set to 10 and **Spike Standard Concentration** value is the expected pg spike per PCR reaction.

- (Optional) **Comments**
- **Protein Concentration**—Drug substance protein concentration (if Total DNA in pg DNA/mg Protein is needed).

Note: The software calculates the pg of DNA in the initial sample and the percentage of recovered DNA

9. Click the **Assign Targets and Samples** tab.
The default plate layout is shown.

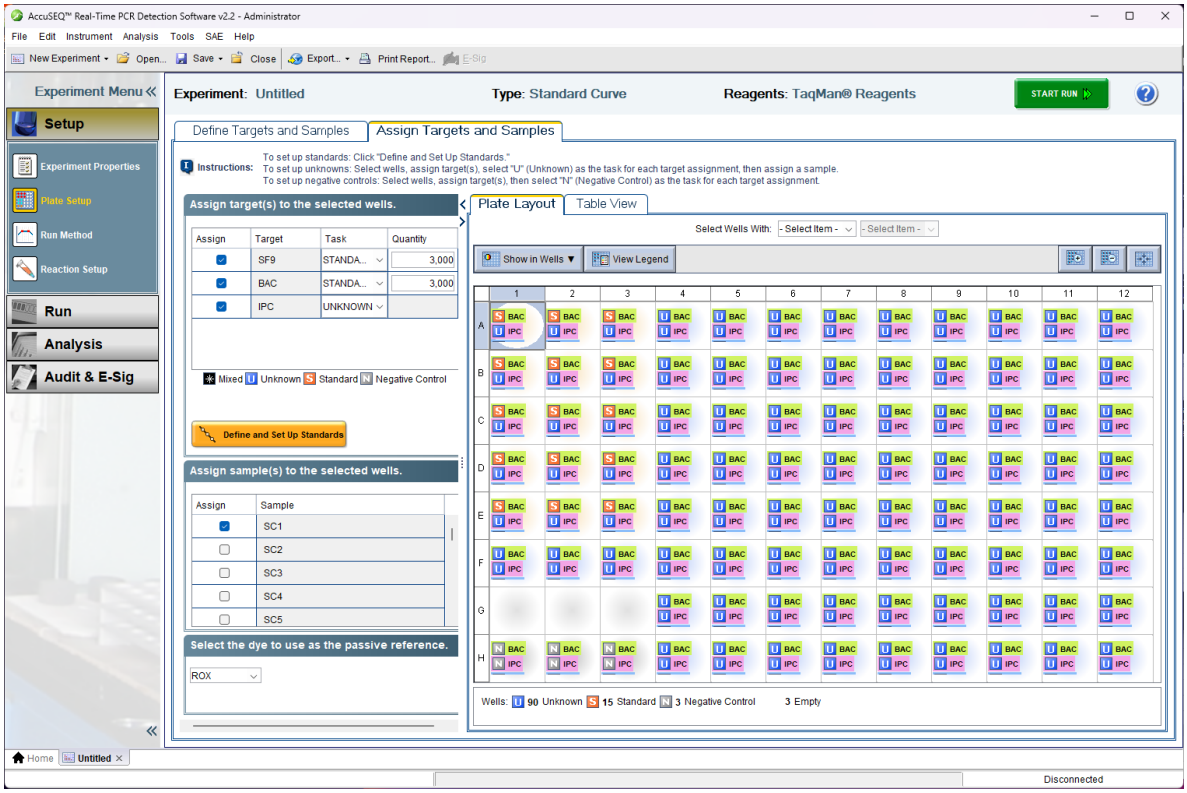


Table 17 Plate well descriptions

Name	Description
SC1 to SC6	Standard curve dilutions in triplicate
Matrix	3 samples in 6 matrices
NTC	No template control in triplicate

10. *(Optional)* Edit the plate.
To change the plate layout display, see *AccuSEQ™ Real-Time PCR Software v2.2 User Guide* (Pub. No. MAN0029201).
   a. Assign the targets to wells in the plate layout.
   b. Assign samples to wells in the plate layout.

11. Confirm that ROX™ dye is selected as the passive reference.

12. *(Optional)* Save the plate layout as an image.
   a. Right-click the plate layout.
   b. Select **Save As**.

c. Specify a file name and location for the JPEG file, then click **Save**.

Note: Electronic signature information is not included in an image file. Use Print Report to include Electronic signature information with the plate layout.

13. *(Optional)* Print the plate layout
    a. Right-click the plate layout.
    b. Click **Print Preview** or **Print**.
14. In the **Experiment Menu** navigation bar, click **Run Method**.
15. In the **Run Method** screen, view the default volume and cycling conditions.

IMPORTANT! Do not modify the run method for factory default SEQ experiments.

16. In the **Experiment Menu** navigation bar, click **Reaction Setup**.
17. In the **Reaction Setup** screen, complete the reaction setup, see *AccuSEQ™ Real-Time PCR Software v2.2 User Guide* (Pub. No. MAN0029201).

    Assemble the PCR reactions using the instructions of the manufacturer for the reagents and following the plate layout.

Create a Quantitative Kanamycin Assay experiment

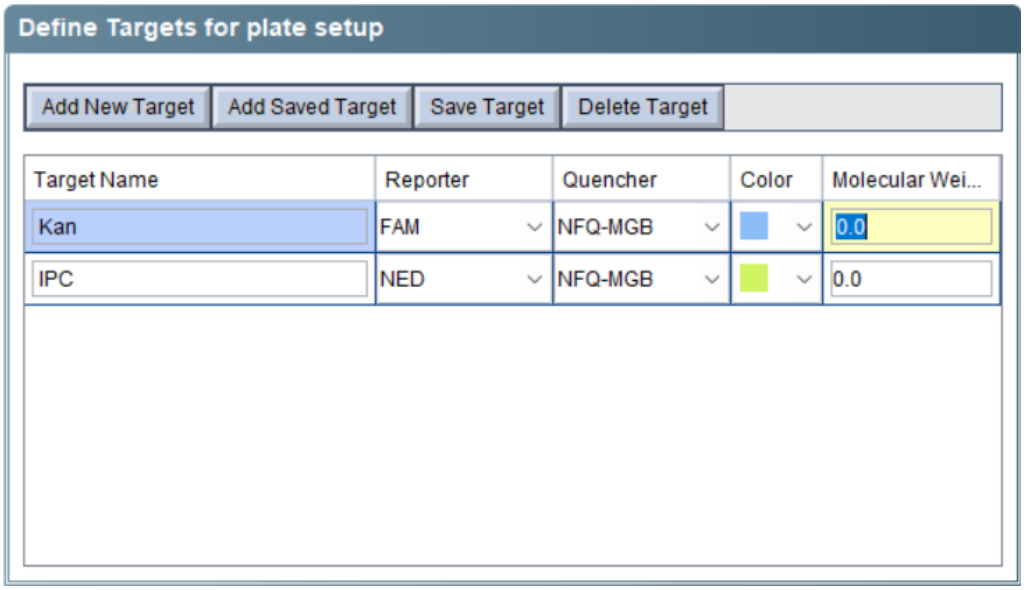
Note: The **Quantitative Kanamycin Assay** SEQ experiment is available with AccuSEQ™ Software v2.2 or later.

1. Click **Create SEQ Experiment** in the home screen.
2. Select **Quantitative Kanamycin Assay** from the assay list, then click **Next** .
   The **Plate Setup** screen opens.
3. In the **Experiment Menu** navigation bar, click **Experiment Properties**.
4. Enter the experiment name.

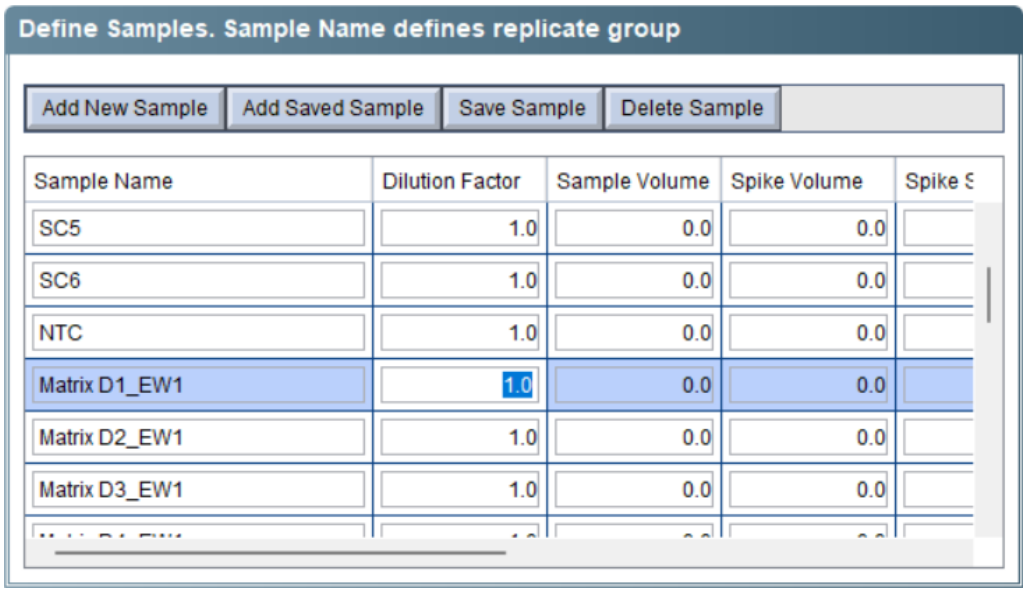
Note: The experiment name can be up to 100 letters and numbers. Spaces are not allowed.

5. *(Optional)* Enter a plate barcode and comments.
6. In the **Experiment Menu** navigation bar, click **Plate Setup**.
7. In the **Define Targets** pane of the **Define Targets and Samples** pane, enter the molecular weight (g/mol) of the targets to calculate mass (g) in the results.

IMPORTANT! Do not change the **Targets**.

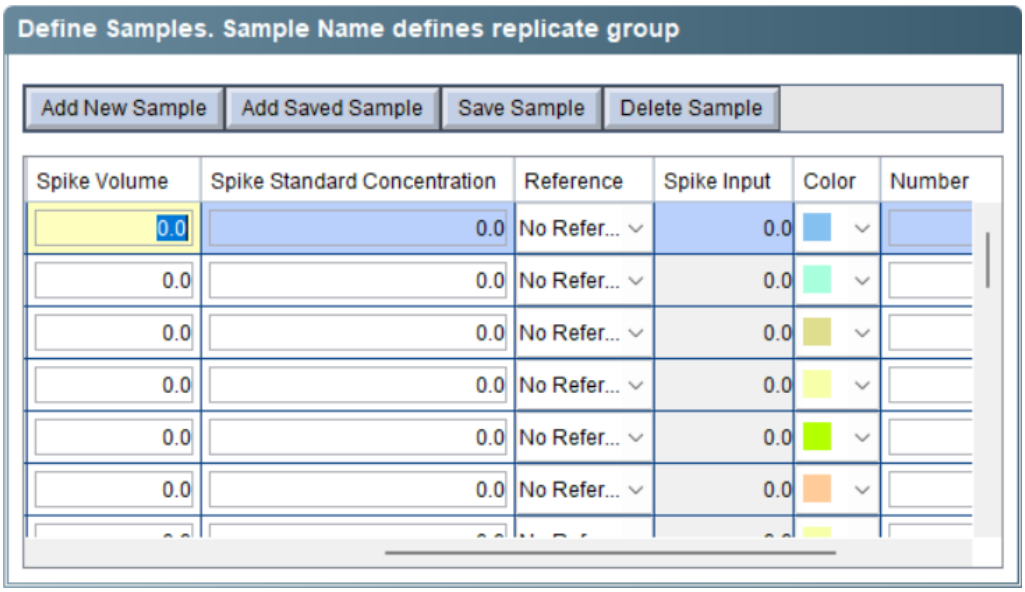


8. In the **Define Samples** pane of the **Define Targets and Samples** tab of the **Plate Setup** pane, enter the sample name and the sample dilution. Sample volume is not needed. Click **Add New Sample** to add more samples if needed.



For information about adding and defining samples, see *AccuSEQ™ Real-Time PCR Software v2.2 User Guide* (Pub. No. MAN0029201).

9. In the **Define Samples** pane in the **Define Targets and Samples** tab, scroll to the right, then enter the spike information.



- **Sample Volume**—not needed; leave as default (0).
- **Spike Volume**—volume of DNA added to the PCR (set to 10).
- **Spike Standard Concentration**—expected spike amount per reaction (for example, 10 copies).
- **Reference**—the non-spiked sample; the mean quantity of reference is subtracted during % recovery calculation.
- **Spike Input**—calculated (double check if correct).

  Note: If not correct, ensure **Spike Volume** is set to 10 and **Spike Standard Concentration** is the expected copies spike per PCR reaction.

- **Protein Concentration**—Drug substance protein concentration (if total DNA in copies DNA/mg protein is needed).
- (Optional) **Comments**

Note: The software calculates the mass (g) and % recovery in the **Table View** tab of the **Analysis** screen.

10. Click the **Assign Targets and Samples** tab.
The default plate layout is shown.

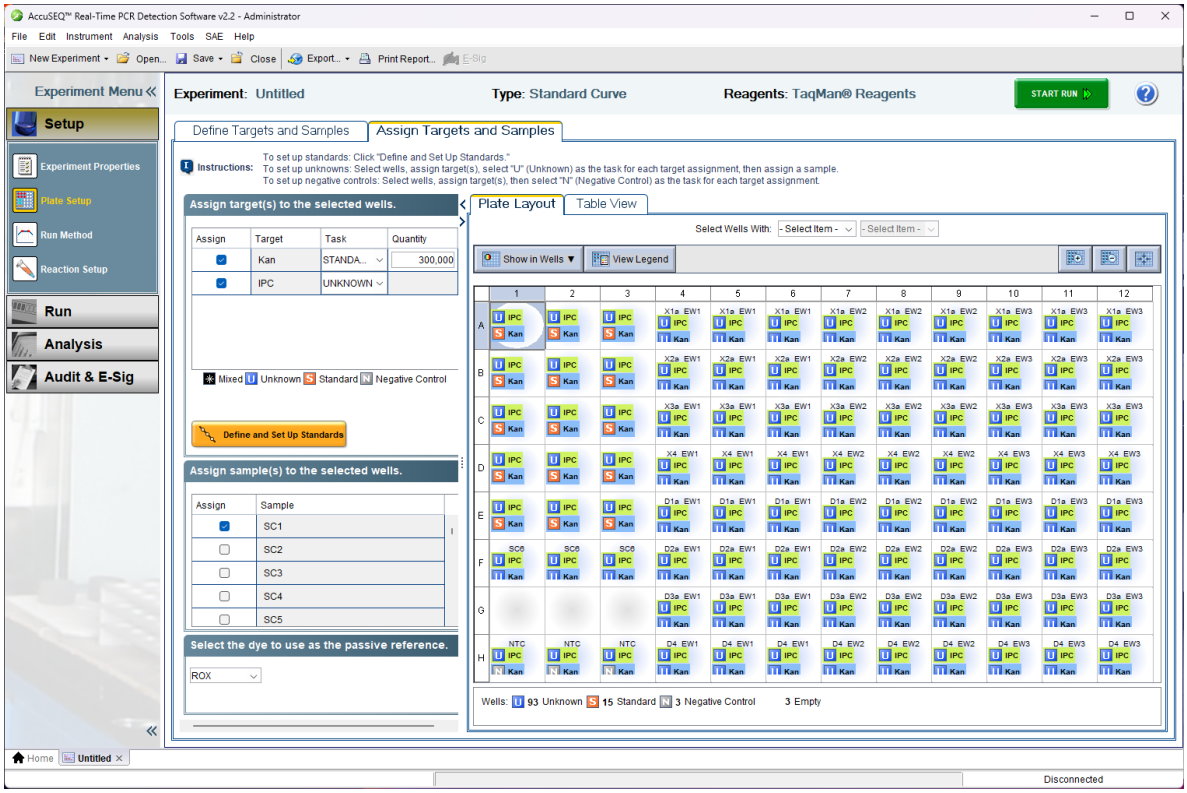


Table 18 Plate well descriptions

Name	Description
SC1 to SC6	Standard curve dilutions in triplicate
X1 to X3, D1 to D4	24 samples in triplicate
NTC	No template control in triplicate

11. *(Optional)* Edit the plate.
To change the plate layout display, see *AccuSEQ™ Real-Time PCR Software v2.2 User Guide* (Pub. No. MAN0029201).
    a. Assign the targets to wells in the plate layout.
    b. Assign samples to wells in the plate layout.

12. Confirm that ROX™ dye is selected as the passive reference.

13. *(Optional)* Save the plate layout as an image.
    a. Right-click the plate layout.
    b. Select **Save As**.

c. Specify a file name and location for the JPEG file, then click **Save**.

Note: Electronic signature information is not included in an image file. Use Print Report to include Electronic signature information with the plate layout.

14. *(Optional)* Print the plate layout
    a. Right-click the plate layout.
    b. Click **Print Preview** or **Print**.
15. In the **Experiment Menu** navigation bar, click **Run Method**.
16. In the **Run Method** screen, view the default volume and cycling conditions.

IMPORTANT! Do not modify the run method for factory default SEQ experiments.

17. In the **Experiment Menu** navigation bar, click **Reaction Setup**.
18. In the **Reaction Setup** screen, complete the reaction setup, see *AccuSEQ™ Real-Time PCR Software v2.2 User Guide* (Pub. No. MAN0029201).

    Assemble the PCR reactions using the instructions of the manufacturer for the reagents and following the plate layout set up in the template.

Run and monitor an experiment

To load and start the instrument, see *Applied Biosystems™ 7300/7500/7500 Fast Real-Time PCR System Getting Started Guide: Absolute Quantitation using Standard Curve* (Pub. No. 4347825).

At the top of every run screen, the run status is shown.

During an experiment run, progress bars reflect run status. During the third stage of a the experiment, there is a delay between melting and completion of the third stage during which the system analyzes the data.

1. In the experiment menu, select **Run**.
2. If the run is not already in progress, click **Start Run**.
3. If the run is in progress, select a run screen from the navigation pane to monitor the progress.
   - **Amplification Plot**–Select to view the change in normalized reporter signal (ΔRn) for each cycle.

     Note: Shown only for experiments with amplification.
   - **Temperature Plot**–Select to view the temperature (°C) plotted against time.
   - **Run Method**–Select to view the run method status during the instrument run.

   To modify the plots in the **Run** screen, see *AccuSEQ™ Real-Time PCR Software v2.2 User Guide* (Pub. No. MAN0029201).

4. *(Optional)* Click **Stop Run** at the top-left of the run screen to stop the run.
   If the computer is connected to the instrument, the software can stop the instrument run.

   IMPORTANT! You cannot resume a run after you stop it.

Analyze the results

After the run is finished, use the following general procedure to analyze the results:

1. In the toolbar, select **Analysis** ▸ **Analysis Settings**.
2. Click 

(**Analyze**).
3. Select **Analysis** ▸ **QC Summary** in the left panel of the screen. Review the flag summary.
4. In the left panel, select **Analysis** ▸ **Standard Curve**. Verify the values for the Slope, Y-Intercept, R2, and Efficiency.
5. Select **File** ▸ **Print Report** to generate a hardcopy of the experiment, or click **Print Preview** to view and save the report as a PDF or HTML file.
6. Optional: Select **File** ▸ **Export**. In the **Export Data** menu, select file type ***.xls**. Click **Start Export**.

Use the kit with 7500 System SDS Software v1.5.1

Create the plate document, run the plate, and analyze the results with 7500 Fast SDS software

The following instructions apply only to the Applied Biosystems™ 7500 Fast instrument with SDS v1.x software. If you use a different instrument or software, refer to the applicable instrument or software documentation. Genomic residual DNA is measured in concentration, while plasmid residual DNA is measured in copy number.

Create a plate document

Residual DNA assays are duplex assays, containing sample DNA and Internal Positive Control (IPC).

Plate document: settings

If you have run the assay frequently, you can use the table below to enter settings in Plate Document fields. If you are a new user, follow the detailed instructions in the following sections.

Summary of settings for the Plate Document		
In this field…		**Use these settings**
Detector	resDNASEQ™ kit target cell lines	• Single target assays: FAM™ dye • Sf9 and Baculovirus assay: VIC™ and FAM™ dyes (Select **NFQ_MGB** dye for quencher)
	IPC	• Single target assays: VIC™ dye • Sf9 and Baculovirus assay and Plasmid DNA - Kanamycin Resistance assay: NED™ dye (Select **NFQ_MGB** for quencher)
PCR	Hold	Temp: **95**°C Time: **10:00**

(continued)

Summary of settings for the Plate Document		
In this field…		**Use these settings**
PCR	Cycling (Standard Mode)	Cycles: **40** Temp: **95**°C Time: **0:15** Temp: **60**°C Time: **1:00**
Analysis	CHO, *E. coli*, HEK293, Human, MDCK, NS0, *Pichia*, Plasmid DNA - Kanamycin resistance, Sf9 and Baculovirus, and Vero	**Automatic Baseline** *or* **Manual Baseline**[1] Threshold: **0.2** **Note:** For CHO, the upper limit for manual baseline analysis is 12. For the Plasmid DNA - Kanamycin Resistance assay, change the threshold to 0.04 for KanR and 0.02 for IPC.

[1] You can analyze the assay using Automatic or Manual Baseline, use the setting that yields the best standard curve.

Plate document: procedure

Use the SDS v1.x software on the Applied Biosystems™ 7500 Fast instrument to perform this procedure.

1. In the template **Assay** dropdown list, select **Absolute Quantification**.
2. In the **Run Mode** dropdown list, select **Standard 7500**.
3. Enter **resDNA_*Template*** in the **Plate** name field, then click **Next**.
4. Click **New Detector**:
   a. Enter the name of the target cell line in the **Name** field.
   b. Select reporter **FAM™** dye and quencher **NFQ_MGB** dye.

      Note: For the Sf9 and Baculovirus assay, select reporter VIC™ dye for the Sf9 target and quencher **NFQ_MGB** dye. Add an additional target, and select reporter FAM™ dye for the baculovirus target and quencher **NFQ_MGB** dye.

   c. Select a color for the detector, then click **Create Another**.
5. Click **New Detector**:
   a. Enter **IPC** in the **Name** field.
   b. Select reporter **VIC™** dye and quencher **NFQ_MGB** dye.

      Note: For the Sf9 and Baculovirus assay and the Plasmid DNA - Kanamycin Resistance assay, select the reporter NED™ dye and the quencher **NFQ_MGB** dye.

   c. Select a color for the detector, then click **OK**.

d. Select the detectors, then click **Add>>** to add the detectors to the document (plate).

6. Select **ROX™** dye as the passive reference dye, then click **Next**.

7. Select the applicable set of wells for the samples, then select the target cell line and **IPC** detectors for each well. The following figure shows an example plate layout:

										Standard Curve (pg)		
	1	2	3	4	5	6	7	8	9	10	11	12
A	TS1	TS1	TS1		TS1 ERC	TS1 ERC	TS1 ERC			NTC	NTC	NTC
B	TS2	TS2	TS2		TS2 ERC	TS2 ERC	TS2 ERC					
C	TS3	TS3	TS3		TS3 ERC	TS3 ERC	TS3 ERC			0.03 pg	0.03 pg	0.03 pg
D										0.3 pg	0.3 pg	0.3 pg
E										3 pg	3 pg	3 pg
F	NEG	NEG	NEG							30 pg	30 pg	30 pg
G										300 pg	300 pg	300 pg
H										3,000 pg	3,000 pg	3,000 pg

8. Set tasks for each sample type by clicking on the **Task Column** dropdown list:
   a. NTC: target cell line detector task = **NTC**
   b. NEG, test samples, and ERC wells: target DNA detector task = **Unknown**
   c. IPC = **Unknown** for all wells

9. Set up the standard curve:
   a. Select the wells.
   b. Assign the tasks (target DNA = **Standard**) and enter the appropriate Quantity for each set of triplicates.

Tube label	Row-wells	Task	Quantity	Label (pg)
SD 1	H-10, 11, 12	Standard	3,000	3,000 pg
SD 2	G-10, 11, 12	Standard	300	300 pg
SD 3	F-10, 11, 12	Standard	30	30 pg
SD 4	E-10, 11, 12	Standard	3	3 pg
SD 5	D-10, 11, 12	Standard	0.3	0.3 pg
SD 6 (for CHO, Vero, MDCK, and NS0 only)	C-10, 11, 12	Standard	0.03	0.03 pg

Note: The Plasmid DNA - Kanamycin Resistance assay creates a standard curve based on copy number (300,000 to 30 copies per reaction).

10. Select the **Instrument** tab, then set thermal-cycling conditions:
    - Set the **thermal cycling reaction volume** to **30 µL**.
    - Set the reaction to **Standard Mode**.
    - Set the temperature and the time as shown in the following table:

Step	AmpliTaq Gold™ enzyme activation	PCR	
	Hold	Cycle (40 Cycles)	
		Denature	Anneal/extend
Temp (°C)	95	95	60
Time (mm:sec)	10:00	0:15	1:00

    Refer to the applicable 7500 Fast Real-Time PCR Systems instrument manual for additional information.

11. In the **Analysis Settings** window, enter the following settings, then click **OK**:
    a. Select **Manual Ct**.
    b. In **Threshold**, enter **0.2**.

    Note: For the Plasmid DNA - Kanamycin Resistance assay, change the threshold to 0.04 for KanR and 0.02 for IPC.

    c. Select **Automatic Baseline** or **Manual Baseline**.

Note: You can analyze the assays using Automatic or Manual Baseline, use the setting that yields the best standard curve. For CHO, the upper limit threshold for manual baseline analysis is 12.

12. Select **File ▸ Save as**, confirm that the file is named "resDNA_Template", then select **SDS Templates (*.sdt)** in the **Save as type** dropdown list and close the template plate document.

Note: You can reuse the plate template document whenever you run the assay.

13. Close the saved template file.

Run the plate

1. In the SDS software, select **File ▸ New**, navigate to the **resDNA_Template** file (created in "Plate document: procedure" on page 63), then click **Open**.
2. In **Plate Name**, enter an appropriate experiment name, then click **Finish**.
3. Make any necessary changes to the test sample labels.
4. Select **Save As** to save the new experiment as an SDS experiment file.

5. Load the plate on the instrument.

6. Select the **Instrument** tab, save the document, then click **Start** to start the real-time qPCR run.

Analyze the results

After the qPCR run is finished, use the following general procedure to analyze the results.

1. Select the **Results** tab.

2. Select the **Amplification Plot** tab.

3. Verify the analysis settings, change as appropriate, then click **Analyze**.

4. Select the **Results tab ▸ Standard Curve** tab, then verify the Slope, Intercept, and R2 values.

5. Right-click the **Standard Curve**, select **Export as JPEG**, then click **OK**. Alternatively, press **PrintScreen**, then paste the image in a WordPad file.

6. Select the **Report tab ▸ Report**, then review the mean quantity and standard deviation for each sample.

7. Optional: Select **File ▸ Export ▸ Results**. In the **Save as type** drop-down list, select **Results Export Files (*.csv)**, then click **Save**.

D

Good laboratory practices

Work area setup and lab design

The sensitivity of this kit (and other PCR-based tests) enables amplification of minute quantities of DNA, necessitating precautions to avoid contamination of samples yet to be amplified (Kwok and Higuchi, 1989).

Process samples carefully to prevent contamination by human DNA. Wear gloves at all times and change them frequently. Close sample tubes when not in use. Limit aerosol dispersal by handling sample tubes and reagents carefully.

Good laboratory practices for PCR and RT-PCR

- Wear clean gloves and a clean lab coat.
  - Do not wear the same gloves and lab coat that you have previously used when handling amplified products or preparing samples.
- Change gloves if you suspect that they are contaminated.
- Maintain separate areas and dedicated equipment and supplies for:
  - Sample preparation and reaction setup.
  - Amplification and analysis of products.
- Do not bring amplified products into the reaction setup area.
- Open and close all sample tubes carefully. Avoid splashing or spraying samples.
- Keep reactions and components capped as much as possible.
- Use a positive-displacement pipettor or aerosol-resistant barrier pipette tips.
- Clean lab benches and equipment periodically with 10% bleach solution or DNA decontamination solution.

Avoiding false positives due to cross-contamination

To avoid false positives due to cross-contamination:

- Do not open tubes after amplification.
- Use different sets of pipettors when pipetting negative control, unknown, and positive control samples.

Note: Refer to “Prepare the PCR plate” on page 16 for best practice.

Safety

WARNING! GENERAL SAFETY. Using this product in a manner not specified in the user documentation may result in personal injury or damage to the instrument or device. Ensure that anyone using this product has received instructions in general safety practices for laboratories and the safety information provided in this document.

- Before using an instrument or device, read and understand the safety information provided in the user documentation provided by the manufacturer of the instrument or device.
- Before handling chemicals, read and understand all applicable Safety Data Sheets (SDSs) and use appropriate personal protective equipment (gloves, gowns, eye protection, and so on). To obtain SDSs, see the “Documentation and Support” section in this document.

Chemical safety

WARNING! GENERAL CHEMICAL HANDLING. To minimize hazards, ensure laboratory personnel read and practice the general safety guidelines for chemical usage, storage, and waste provided below. Consult the relevant SDS for specific precautions and instructions:

- Read and understand the Safety Data Sheets (SDSs) provided by the chemical manufacturer before you store, handle, or work with any chemicals or hazardous materials. To obtain SDSs, see the "Documentation and Support" section in this document.
- Minimize contact with chemicals. Wear appropriate personal protective equipment when handling chemicals (for example, safety glasses, gloves, or protective clothing).
- Minimize the inhalation of chemicals. Do not leave chemical containers open. Use only with sufficient ventilation (for example, fume hood).
- Check regularly for chemical leaks or spills. If a leak or spill occurs, follow the manufacturer cleanup procedures as recommended in the SDS.
- Handle chemical wastes in a fume hood.
- Ensure use of primary and secondary waste containers. (A primary waste container holds the immediate waste. A secondary container contains spills or leaks from the primary container. Both containers must be compatible with the waste material and meet federal, state, and local requirements for container storage.)
- After emptying a waste container, seal it with the cap provided.
- Characterize (by analysis if needed) the waste generated by the particular applications, reagents, and substrates used in your laboratory.
- Ensure that the waste is stored, transferred, transported, and disposed of according to all local, state/provincial, and/or national regulations.
- **IMPORTANT!** Radioactive or biohazardous materials may require special handling, and disposal limitations may apply.

WARNING! HAZARDOUS WASTE (from instruments). Waste produced by the instrument is potentially hazardous. Follow the guidelines noted in the preceding General Chemical Handling warning.

WARNING! 4L Reagent and Waste Bottle Safety. Four-liter reagent and waste bottles can crack and leak. Each 4-liter bottle should be secured in a low-density polyethylene safety container with the cover fastened and the handles locked in the upright position.

Biological hazard safety

WARNING! Potential Biohazard. Depending on the samples used on this instrument, the surface may be considered a biohazard. Use appropriate decontamination methods when working with biohazards.

WARNING! BIOHAZARD. Biological samples such as tissues, body fluids, infectious agents, and blood of humans and other animals have the potential to transmit infectious diseases. Conduct all work in properly equipped facilities with the appropriate safety equipment (for example, physical containment devices). Safety equipment can also include items for personal protection, such as gloves, coats, gowns, shoe covers, boots, respirators, face shields, safety glasses, or goggles. Individuals should be trained according to applicable regulatory and company/ institution requirements before working with potentially biohazardous materials. Follow all applicable local, state/provincial, and/or national regulations. The following references provide general guidelines when handling biological samples in laboratory environment.

- U.S. Department of Health and Human Services, *Biosafety in Microbiological and Biomedical Laboratories (BMBL)*, 6th Edition, HHS Publication No. (CDC) 300859, Revised June 2020
  https://www.cdc.gov/labs/pdf/CDC-BiosafetymicrobiologicalBiomedicalLaboratories-2020-P.pdf
- Laboratory biosafety manual, fourth edition. Geneva: World Health Organization; 2020 (Laboratory biosafety manual, fourth edition and associated monographs)
  www.who.int/publications/i/item/9789240011311

F

Documentation and support

Related documentation

Document	Publication number	Description
resDNASEQ™ Quantitative DNA Kits Quick Reference	4469837	For brief instructions about using the resDNASEQ™ Quantitative DNA Kits.
resDNASEQ™ Quantitative E1A DNA Fragment Length Kit User Guide	MAN0025643	For instructions about running the resDNASEQ™ Quantitative E1A DNA Fragment Length Kit (Cat. No. A51969).
PrepSEQ™ Sample Preparation Kits User Guide	4469838	For information about preparing samples for extraction.
PrepSEQ™ Residual DNA Sample Preparation Kit Quick Reference	4469839	For brief instructions about preparing samples for extraction.
AccuSEQ™ Real-Time PCR Software v3.1 User Guide	100094287	For information about AccuSEQ™ Real-Time PCR Software v3.1 with the QuantStudio™ 5 Real-Time PCR System
AccuSEQ™ Real-Time PCR Software v3.1 Quick Reference	100094288	For basic information about AccuSEQ™ Real-Time PCR Software v3.1 with the QuantStudio™ 5 Real-Time PCR System
AccuSEQ™ Real-Time PCR Software v3.2 User Guide	MAN0029199	For information about AccuSEQ™ Real-Time PCR Software v3.2 with the QuantStudio™ 5 Real-Time PCR System
AccuSEQ™ Real-Time PCR Software v3.2 Quick Reference	MAN0029200	For basic information about AccuSEQ™ Real-Time PCR Software v3.2 with the QuantStudio™ 5 Real-Time PCR System
AccuSEQ™ Real-Time PCR Software v2.2 User Guide	MAN0029201	For information about AccuSEQ™ Real-Time PCR Software v2.2 with the 7500 Fast instrument.
Applied Biosystems™ 7300/7500/7500 Fast Real-Time PCR System Getting Started Guide: Absolute Quantitation using Standard Curve	4347825	For information about the 7500 Fast instrument.
AccuSEQ™ software: Custom Quick Reference Card	4425585	For information about AccuSEQ™ software with the 7500 Fast instrument.

Customer and technical support

Single Point of Contact	Contact Information
Technical Support North America	+1 800 955 6288, select option 2, then option 1, then option 2. techsupport@thermofisher.com
Technical Support Europe, the Middle East, and Africa (EMEA)	+00 800 5345 5345 eurotech@thermofisher.com
Technical Support Asia-Pacific (APAC)	Contact your local Thermo Fisher Scientific representative

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- Product documentation
  - User guides, manuals, and protocols
  - Certificates of Analysis
  - Safety Data Sheets (SDSs; also known as MSDSs)

  Note: For SDSs for reagents and chemicals from other manufacturers, contact the manufacturer.

Limited product warranty

Life Technologies Corporation and/or its affiliate(s) warrant their products as set forth in the Life Technologies' General Terms and Conditions of Sale at **www.thermofisher.com/us/en/home/global/terms-and-conditions.html**. If you have any questions, please contact Life Technologies at **www.thermofisher.com/support**.