

Dispensing Software User Guide

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Introduction to the dispenser

How the dispenser works

The dispenser provides fast and reliable dispensing performance across a large dynamic range, eliminating the need for slow, error-prone and wasteful serial dilutions. The dispenser also enables assay miniaturization through low-volume assay component addition.

The dispenser provides dispensing across a wide range of life science applications:

- Dose-response curves for small molecule and biological-based drug discovery research
- Drug-drug synergy experiments for drug discovery
- Low-volume assay component dispensing of a wide range of fluids into microplates
- Complex enzyme kinetic studies can be made simple using the Enzyme Profile function
- Microplate PCR setups are easily designed and dispensed using the PCR function
- DMPK standards
- Single cell isolation
- Single cell proteomics, genomics, and transcriptomics
- Cell line development

Using single-use dispense heads to help avoid the possibility of cross-contamination, the dispenser offers non-contact digital dispensing in the picoliter to microliter range. It combines walk up convenience, ready-to-run performance and flexible, intuitive experiment design software to increase research productivity, reliability of results and the speed of life science research including single cell isolation.

Even the most complex experimental plate layouts, such as drug-drug interaction (synergy) studies, can be made simple. Titrating one drug down the plate while another is titrated orthogonally across the plate enables two or more drug doses to be independently generated in every well. In addition, plate layouts can be randomized to reduce the impact of edge effects, increasing the integrity of the experimental data. The dispenser offers a convenient solution to address the most challenging experiments.

Advantages of the dispenser

The dispenser adds breakthrough capability to life science research workflows by dispensing picoliter droplets directly into assay-ready plates.

Dispenser advantages

- **Elimination of Serial Dilution:** Directly and independently dispense doses in each well using the proper number of picoliter droplets. Use finely spaced sets of doses to improve dose response curve fits. Digitally randomize plate layouts to reduce edge effects and increase data integrity.
- **Resource Savings:** Save labor by eliminating serial dilutions. Save 90% of your precious compounds. Save numerous pipette tips and intermediate plates to reduce your waste stream. Avoid ergonomic injuries and rework associated with repetitive hand pipetting processes.
- **Ease of Use:** Enjoy a simple, intuitive user experience with quick do-it-yourself experiment design software. Boost productivity with walk-up, just-in-time bench top convenience. Avoid carry-over and washing with single use, non-contact dispensing supplies. Add all low-volume assay components with one system.
- **Single cell research workflow solution:** Simple isolation of single cells ranging from 10-25 μm coupled with reagent dispensing in up to 384-well plates enables comprehensive workflow for single cell research.

Safety

The dispenser should be used in accordance with all applicable safety rules that are in effect in the lab in which it is installed. Usage of the dispenser should also follow common-sense safety guidelines for chemical, mechanical, and electrical safety.



NOTE: If this unit requires servicing, or you are unsure about any aspect of operation or safety, contact the Service Help Desk. The contact information can be found in the dispenser **Service and Support** document by navigating to the **File → Documentation** menu of the software.



CAUTION: Do not remove any of the dispenser covers. Contact your Service Help Desk if the dispenser is not working properly. Do not attempt to repair the dispenser yourself.

General safety rules

- Place the dispenser on a work surface that supports all four feet securely.
- Plug the dispenser into a standard, grounded, electrical outlet. Note the power requirement listed on the back label of the dispenser.
- Avoid the moving parts of the dispenser during operation.

Personal protective equipment

The dispenser dispenses chemicals normally used by your company. The same safety precautions and personal protective equipment (PPE) specified by your employer for fluid transfer operations, such as pipetting into bio-assay well plates, should be used when operating the dispenser and handling these materials.

Possible inhalation hazard

The dispenser was designed with chemical safety in mind; however, it does dispense picoliter droplets of fluid and generates a small amount, less than 1% by volume, of very small aerosol droplets that might not be trapped within the well plate that you are dispensing into. If the fluids are known or suspected to be an inhalation or contact hazard, consider using the dispenser in an exhausted enclosure such as a lab fume hood, or a Class 1 or 2 biosafety cabinet.

Ergonomic safety

The dispenser typically requires a single manual pipetting operation for each dispensed compound. In most situations, using the dispenser in the intended manner decreases the frequency of shoulder, arm, and wrist motions and so decreases the stress on these joints compared to traditional methods.

As with any work task, workstation setup is important. To minimize ergonomic risks associated with the use of the dispenser, consider the following work area design ideas:

- Place the dispenser on a work bench at a height that allows for proper visual and physical access for operators. Having the dispenser too high can lead to greater shoulder elevation when pipetting into the dispense head and having the dispenser too low can lead to increased neck and back flexion.
- If an operator is too short to comfortably access the dispenser, consider moving the dispenser to a lower bench height. Alternatively, use a footstool or high lab stool that adjusts to the proper work height.
- Place the dispenser close to the front of the work surface to minimize the distance the operator must reach to pipette into the dispense head.
- Place the pipette tip disposal receptacle close to the dispenser.

- Place the computer keyboard close to the dispenser to minimize reach distance.
- Adjust the computer monitor height and screen angle for good physical and visual access during dispenser operation.
- Perform data analysis at a separate ergonomically designed workstation whenever possible rather than at the dispenser computer.

Additional assistance

Contact the Service Help Desk for additional assistance. The contact information can be found in the dispenser **Service and Support** document by navigating to the **File → Documentation** menu of the software.

System specifications

Intended use

The dispenser and dispense heads are intended only for the following, wherein all conditions must be met:

- Dispensing of DMSO solvent or water containing surfactant with or without small molecule compounds or biomolecule(s).
- Dispensing of aqueous fluids without the need for surfactant with or without small molecule compounds or biomolecule(s).
- Dispensing of single cells ranging in size from 10-25 μm
- For research use only.
- Single-usage (i.e., non-reusable) operation of a dispense head.

In addition to the requirement of Intended Use, the products should not be used for any of the following: (a) applications in Bio Safety Level 3 or 4 (BSL-3 or-4), food, animal or human testing or therapeutic delivery (except solely for therapeutic delivery for in vitro research uses) (b) to dispense any of: bacteria, viruses, pathogens, contagions, infectious agents, potent toxins or oncogenes, radioactive substances; and (c) in uses that are unsafe, bypassing any Product safety features, and/or disobeying any safety instruction.

System requirements

- Microsoft Windows 8 or greater required; Windows 10 recommended.
- Microsoft .Net Framework 4.8 or greater required.
- 8 GB system RAM and 256 GB drive (minimum) recommended.
- Minimum screen resolution of 1280x720 pixels.
- Adobe Acrobat Reader – www.adobe.com/downloads.

Compatible fluids

See [Appendix E–Dispense specifications](#) for details of compatible fluids.

Additional specifications

See the dispenser **Operating Manual** for more system specifications. The **Operating Manual** can be found by navigating to the **File** → **Documentation** menu of the software.

Software overview

The dispensing software lets you:

- Design and build protocols for dispensing fluids into well plates.
- Run the protocols on the dispenser to dispense fluids into well plates.

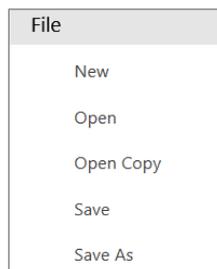
A **protocol** is a file which describes the concentrations or volumes to be dispensed by the dispenser. You create protocols with the tools and functions described in this manual.

The **Main Operating Window** is the first window displayed when you start the software. It gives you control over protocol design functionality. From the Main Operating Window, you create, set up, edit, and run your dispensing protocols.

Loading an example protocol

A good way to become familiar with the dispensing software is to load a sample protocol. Experiment with the features and functionality discussed in this manual at your own pace. Try some or all of these with a focus on the functionality that will be most applicable to your work with the dispenser.

- Click the **File** drop-down in the upper left corner. Click **Open** to browse the default protocols folder and load the **Example Protocol** from your default protocols folder.



Select the **Example Protocol** and then click **Open**.

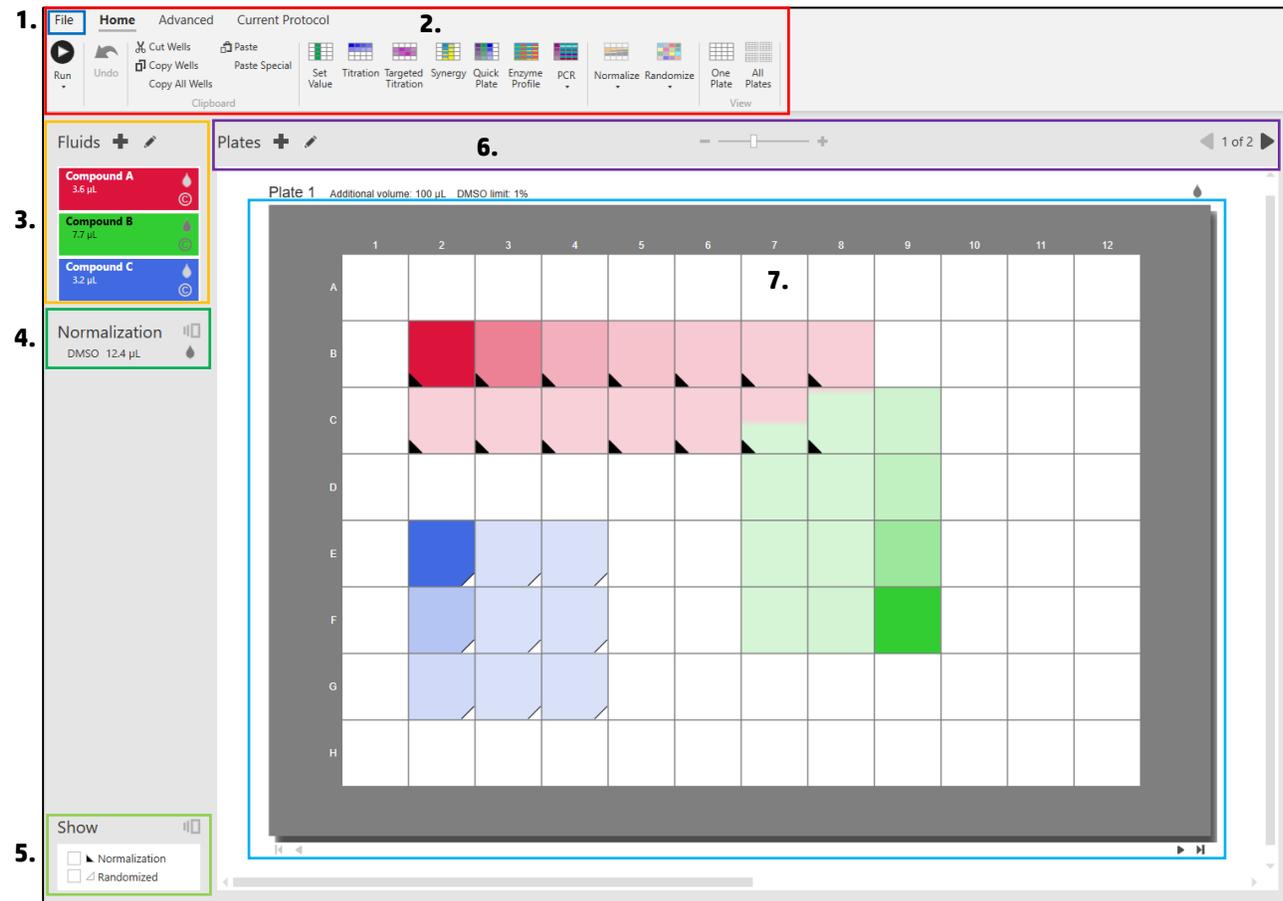
- The **Example Protocol** is then loaded into the Main Operating Window.



TIP: To open a copy of a protocol, use **Open Copy** from the File menu. A copy of the selected protocol will be opened, with a unique file name.

Main operating window - layout

Below is an illustration of the main operating window with a loaded example protocol with the functional areas highlighted. Descriptions of the functional areas follow below.



1. **File Menu:** Allows you to perform file operations common to Windows-based software (New, Save, Open, Exit, etc.) and to adjust system-wide settings.
2. **Ribbon and Tabs:**

The Main Operating Window **ribbon**, located across the top of the window, organizes commonly used tools and functions under three tabs. Access the **functions** associated with a tab by clicking one of the **tabs**:

Home: This tab has most of the basic tools and functions necessary for building a protocol, including undo, copy and paste, and plate viewing, and provides easy access to all the different ways to program the desired titrations.

Advanced: The functions on this tab are related to manipulating the fluids table with copy and paste, well lock, manipulating well plates, and device utilities.

Current Protocol: The functions on this tab are specific to the current protocol being worked on. This tab holds functions for setting the desired concentration or volume units and specifying plate shaking parameters for mixing.

In all tabs, some functions may be “grayed-out” (unavailable) depending on the current items (wells, well plates, fluids, protocol) you are working on. This is the case for functions which require selection prior to

performing the action. The tools and functions available on the tabs are detailed in subsequent chapters of this manual. Most ribbon functions can also be accessed from “right-click” contextual menus.

3. **Fluid List:** The Fluid List contains the fluids you have added to the protocol. The Fluid List indicates the fluid name and total volume of the fluid used in the entire protocol. Icons indicate whether the fluid is dispensed by concentration or volume and whether the fluid will be dispensed when the protocol is run. If the mouse is moved over a fluid, more information is displayed.

You can add to, change, or delete from this list (see [Fluid list](#) section for details).

4. **Normalization:** When a plate has been normalized (see [Normalize](#) section), this display is an indicator that a normalization scheme is part of this protocol.

When the mouse is moved over the  icon, normalization information is displayed, including the volume of all normalization fluids used in the protocol and the desired dispense order for normalization fluids.

Total volume	Dispense order
DMSO 2.0 µL	No priority <input type="button" value="v"/>
a+B 2.0 µL	

5. **Show:** The Plate View area of the Main Operating Window permits turning on or off viewing of normalization and randomization, if those have been applied to the current protocol. Additionally, moving the mouse over the  icon allows you to filter the set of fluids displayed on the plate. See [Show](#) section for details.
6. **Plate Navigation:** The Plate Navigation area allows you to add well plates to the current protocol and to navigate through each of the well plates in a multi-plate protocol. This area also allows you to zoom in or out of the well plate grid. See [Multi-plate protocols and plate navigation](#) section for details.
7. **Well plate grid:** The Main Operating Window contains graphical depictions of the well plates in the current protocol. Each gridded area represents a single well plate, and each square represents a well. The grid has well rows alphabetized on the Y-axis and well columns numbered along the X-axis. The software identifies wells in each plate by a row-column designation – A21, B14, C16, etc.

Information about each plate is displayed just above each plate graphic. You can change the plate settings by double-clicking the information area or selecting Edit Plate from the right-click context menu. A “Don’t dispense” droplet icon is used to skip a plate during a protocol run. See [Plate-associated information](#) section for details.

The well plate receives the fluids that are dispensed by the dispenser. Based on the currently loaded protocol, each well is color-coded to indicate the fluid(s), as well as their respective concentration or volume – as indicated by the color saturation – in that well.

When you move your pointing device over a well, information about the content of that well is displayed, including the concentration and volume of each fluid to be added to the well, and including normalization fluids if applicable.

Since fluid is dispensed in increments of dispense head drops, for low volumes the well content information will display both requested and dispensed concentrations and volumes. The first concentration/volume is the requested concentration or volume. The actual concentration and volume is shown in brackets.

Plate 1 Well C7	
DMSO Volume: 1,010 nL (1,000 %)	
Aqueous Volume: 0.0328 nL	
Additional Volume: 100.0 µL	
Total Volume: 101.0 µL	
<hr/>	
 Compound A  DMSO	
Concentration: 0.00309 µM [0.00396 µM]	
Volume: 0.0312 nL [0.0400 nL]	
 Compound B  a+Tw	
Concentration: 0.00163 µM [0.00163 µM]	
Volume: 0.0328 nL [0.0330 nL]	
 DMSO Normalization	
Volume: 1,010 nL	
<hr/>	
Normalize DMSO to highest volume: 1,010 nL	

See [Appendix D—Concentration calculations](#) for more detail about concentration calculations.



NOTE: In some instances, the concentration/volume you program into a well will not be in the range of the dispenser’s capability (see [Appendix D—Concentration calculations](#)). In such cases, the well contents will display a warning message such as “Below minimum dispense volume.” Additionally, the report file will show only the concentration/volume actually dispensed.

Building a simple protocol

Creating a new protocol file

You can create a new protocol in one of two ways:

Via the initial main operating window

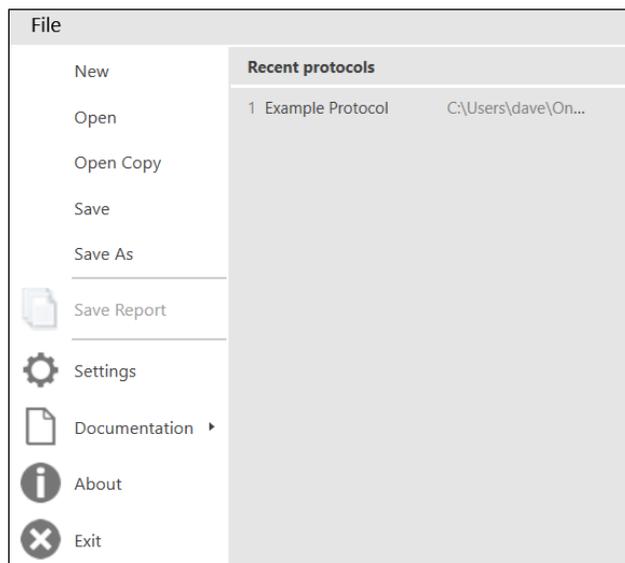
The following procedure will create a new protocol:

- Start the dispensing software.
- The initial “empty” Main Operating Window is displayed.
- Work on that empty protocol, then save your work and name the protocol.

Via the File menu

When the software is running, create a new protocol as follows:

- In the File Menu, click **New**.



A new protocol is created using the default protocol settings.

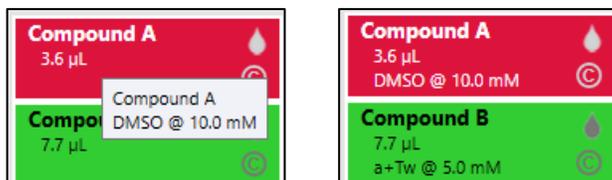
Fluid list

The **Fluid List**, located on the left side of the Main Operating Window, is the place to define all the fluids that can be used in a protocol. Each fluid must be given a unique name, and the fluid concentration (stock concentration) must be identified. Setting up the Fluid List is one of the first steps in creating a protocol.

In a new protocol, the Fluid List is empty. Note that you can copy a list of fluids from another protocol using the [Copy/Paste Fluids from the fluid list](#) function discussed later in this manual. You can also build the Fluid List from scratch.

Show more information

For each fluid, information is displayed in the list, with more detail shown when the mouse is moved over a fluid. To show more information for each fluid, right-click on the list and select **Show More** from the menu. You can also change the default fluid view from the Display section of Settings.



Fluid dispense mode

Each fluid is specified to be dispensed by concentration or by volume.

When dispensed by **Concentration**, the software uses the fluid stock concentration and total well volume to calculate what volume of the fluid is needed to achieve the desired concentration. The software maintains the programmed concentration in the well if other parameters, such as assay volume or stock concentration, are changed.

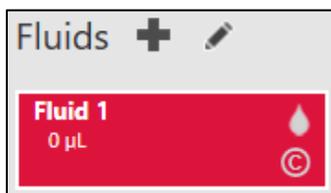
When dispensed by **Volume**, the software maintains the programmed volume in the well if other parameters are changed. If the stock concentration of a fluid is unknown, it must be dispensed by volume; the stock concentration may be left unspecified. If you adjust the stock concentration of a fluid dispensed by volume, the resulting concentration in the programmed wells will be adjusted because the dispensed volume will be maintained.

Adding a fluid

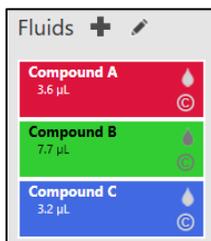
The figure below shows an empty Fluid List. To add a fluid to the list, click the plus sign (+) to the right of the **Fluids** heading. Note that the fluid group, fluid class, concentration, and dispense by concentration/volume default to the values specified in [Settings](#). You can change the default settings.



A preliminary entry is created that can be used as is or modified if desired. To modify any of the parameters, click the  button; you can now **Edit** the entry of any fluid. You can also **Edit** the entry by double clicking.



An example Fluid List is shown below.



For each fluid, the name and total volume of the fluid used in the entire protocol is shown.



NOTE: If the total dispense volume of a fluid exceeds the maximum dispense volume (see [Appendix E–Dispense specifications](#)), the volume will be split between two or more dispense heads.

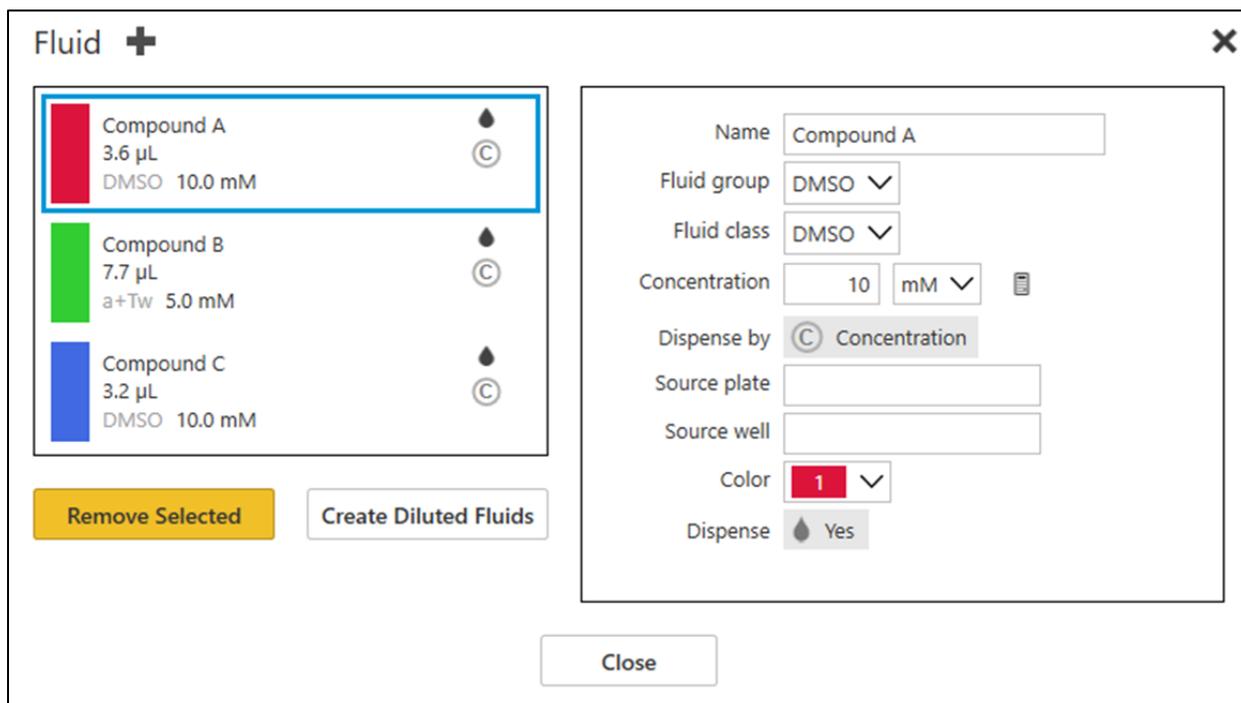
Icons indicate whether the fluid is dispensed by concentration or volume and whether the fluid will be dispensed when the protocol is run. When the mouse is moved over a fluid the fluid class, stock concentration and other information are shown.

Compound B
DMSO 7.5 mM
2007-C / A3

Editing fluids

Click to bring up the **Fluid Edit** dialog box. To edit a fluid, click on the fluid in the list.

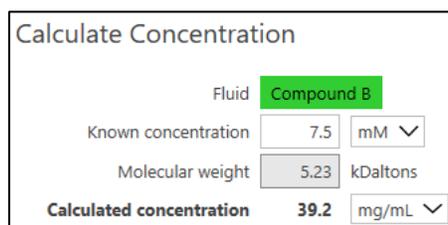
You can also add a new fluid by clicking **+** just as you can do from the main fluid list. Note that this method also copies the concentration, fluid class, dispense mode and source plate from the current fluid to the new fluid.



If desired, provide the fluid a different **name** and edit the stock **concentration**, **dispense** mode, or specify a different **fluid group** and **fluid class**.

The fluid group is used to adjust dispense characteristics for different types of fluids. In addition to DMSO fluids, there are additional classes based on aqueous fluids, acetonitrile, master mixes, and cells. Within each fluid group additional specifications can be defined with the fluid class selection drop down menu. Refer to [Appendix E—Dispense specifications](#) for fluid class details.

The  button can be used to calculate a concentration, converting between units using molecular weight.



Calculate Concentration		
Fluid	Compound B	
Known concentration	7.5	mM
Molecular weight	5.23	kDaltons
Calculated concentration	39.2	mg/mL



NOTE: Concentration can be specified in molarity or mass/volume units, but well concentrations cannot be converted between unit types. If unit type is changed and the fluid is used in the protocol, well fluid dispense values are maintained (concentration or volume depending on the [Fluid dispense mode](#)).

You can also fill in any or all of the following optional information as desired. This information is used in the fluids list and will be included in the report file:

- Source plate
- Source well
- The color used to identify the fluid in the protocol can be changed

If **Dispense** is set to **Don't Dispense** by clicking the droplet icon next to a fluid, you can associate the fluid with wells like the other fluids, but it will not be dispensed. This may be useful when you want a fluid to be used in Normalization calculations or to track controls that will be dispensed by a method other than with the dispenser.

If a fluid is changed, an **Apply** button will appear. You will also be prompted to save when closing the dialog box or selecting a different fluid to edit.

Small molecules

For aqueous fluids with stock concentration specified in mass units, the **small molecule** option is displayed. For this purpose, small molecules are defined as molecules under 1000 Da.

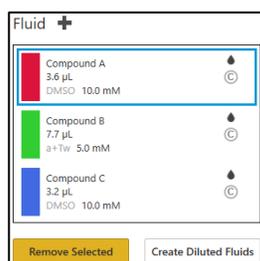


Class	Aqueous+Tween 20	
Concentration	2	mg/mL
Small molecule	<input type="checkbox"/>	

If the source fluid is a small molecule, this option can be checked to prevent the system from applying biomolecule dispense limits based on stock concentration. See [Dispensing considerations](#) for more information.

Removing fluids

Fluids can be removed from the protocol by selecting a fluid and pressing the **Remove Selected** button below the list.



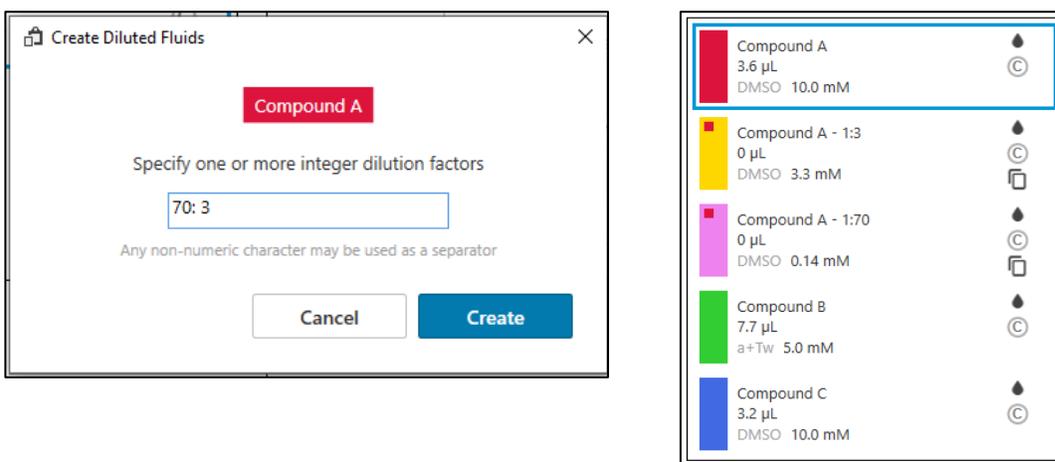
Rearranging fluids

Fluids can be moved up and down the list in the fluid editor by selecting a fluid and dragging it to the desired location with the cursor.

Creating diluted fluids

Diluted fluids can be used to extend the dynamic range of the dispenser. Diluted versions of a fluid can be created using the **Create Diluted Fluids** button. Multiple dilutions can be created at the same time.

The new fluids will be inserted into the fluid list after the original fluid.



NOTE: If a fluid is derived from another fluid using [Creating diluted fluids](#) or as the result of a [Fluid shift or extend](#), some of the information cannot be edited and may change automatically when the source fluid is changed.

Plate parameters

A new protocol is created with a single plate using [Default values](#) from Settings.

See the [Plate-associated information](#) section for details on how to change the plate type or other plate attributes. **Additional volume** must be set correctly; it is the volume added to each well of the plate not using the dispenser and is used for concentration and normalization calculations (see [Appendix D—Concentration calculations](#)).

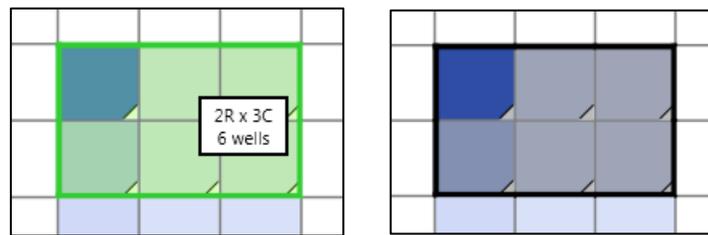
Well selection and de-selection

Prior to using well tools and functions, wells must be selected in the well plate grid.

Selecting wells via rectangle

Select a large group of adjacent wells by dragging a rectangle.

- Move your cursor to the desired area of the well plate grid.
- Click the left button once and keep it pressed, then drag over the target area. Wells will be highlighted as you drag.
- Once you release the left button **all selected wells** are shaded and outlined.



Row/column selection

Entire rows or columns can be selected by clicking on a row or column header to start selection. Only rows or columns from a single plate may be selected until the mouse button is released.

Entire well selection

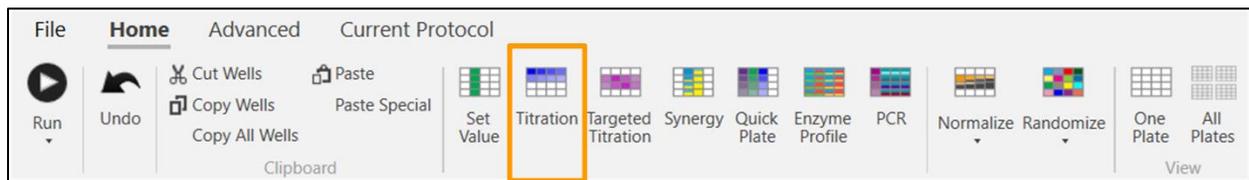
All wells in a plate can be selected by double clicking on any well.

Additional selection/de-selection

Once wells have been highlighted as selected, hold down the **CTRL** key while clicking wells to:

- Add unselected wells by holding the **CTRL** key while clicking an **unselected** well.
- Remove wells from the selection by holding the **CTRL** key while clicking a **selected** well. To de-select all wells, just click outside the plate.

Basic titration example



Fluids can be added to wells in three basic ways: as a fixed concentration or volume ([Set value](#)), as an evenly spaced titration series ([Titration](#)), or as a targeted titration series ([Targeted titration](#)).

The example in this section describes how to set up an evenly-spaced titration series consisting of 8-point triplicates using two different fluids.

- Select 24 wells in the well plate for titration of the first fluid.
- On the Home tab, select **Titration**. An editing panel is then displayed.

- Select a fluid from the drop-down list.
- Specify titration using highest and lowest concentrations.
- Assign **Highest concentration** (or volume) parameter.
- Assign **Lowest concentration** (or volume) parameter.

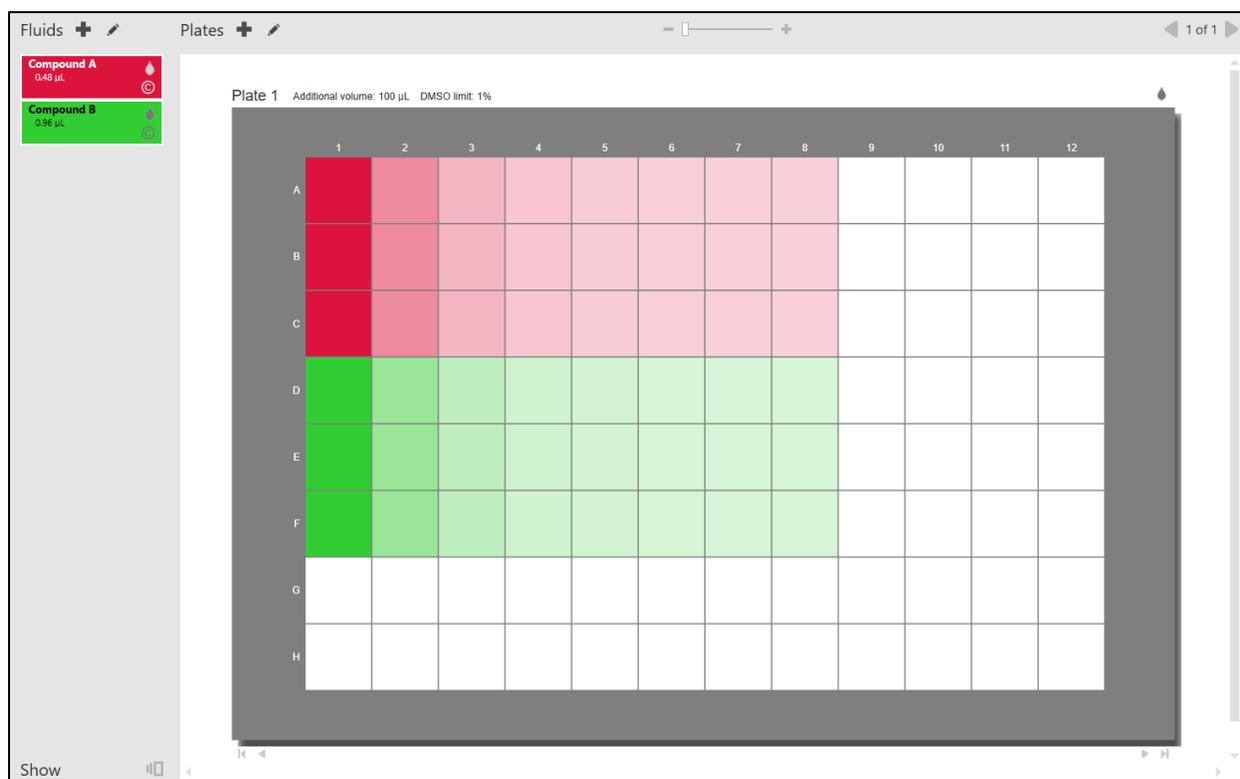


TIP: Use the **paste function** in the lower-right area of the panel to auto-fill the highest and lowest concentration with the valid range limits. The high value is determined from fluid concentration and DMSO limit or recommended maximum aqueous dispense volume. The low value is limited by minimum dispense volume.



- Change **Replicates per level** to 3 and leave the distribution and titration pattern unchanged.
- Once you have completed your edits, click **Apply**. “Apply” applies the function without closing the dialog box, while, “Done” closes the dialog box.
- Now, with the titration dialog box still open, select a new set of 24 wells into which to apply the second fluid.
- In the titration dialog box, select a different fluid from the dialog box and click **Apply**. This will apply the second fluid to the wells with the same settings for concentration/volume as the first fluid.

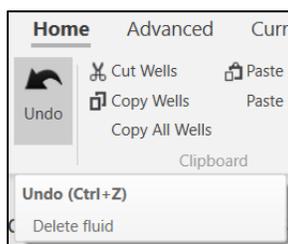
The well plate grid now shows the fluids to be titrated in the selected wells of the well plate. If the titration pattern is not what was intended, or the concentration/volume needs to be adjusted, simply re-select the wells and enter the titration dialog box again, to over-write the wells with the new parameters.



Additional detail about titration can be found in the [Titration](#) section.

Undo

To undo an action after you perform it, click the Undo icon in the Home tab of the ribbon or use the keyboard shortcut (CTRL-Z).



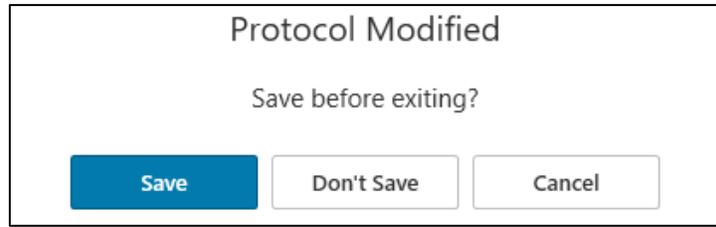
Saving the protocol file

Make sure you save your work during and after each protocol design session. On the File Menu, click **Save** at any time to save your work.

Note: If you are saving a protocol for the first time, a browsing window will be displayed where you can name and relocate the file.

Closing the application and saving your work

In the Main Operating Window, clicking the X closes the current protocol and exits the program. If you modified the protocol and haven't saved, you are asked if you want to save the modifications.



TIP: Even if you have saved changes in one or more of the tools and functions described above, if you click **Don't Save**, *all of your modifications to the protocol will be discarded.*

Additional protocol features

The [Building a simple protocol](#) chapter detailed the basics of creating a protocol. This chapter discusses all the tools and functions you can use to build protocols.

Multi-plate protocols and plate navigation



The **Plate Navigation** area of the Main Operating Window allows you to

- Create a protocol with more than one plate,
- Navigate through each of the plates in a multi-plate protocol, and
- Zoom the well plate grid.

To **add a new, empty plate** to the current protocol, click the **+** sign. When you do this, the empty plate is displayed and becomes the one you are working on.



NOTE: To minimize evaporation, it is recommended that protocols using aqueous fluid classes be limited to 1536 wells or four plates.

The **Plate N of X** label indicates the current plate number you are working on (N) and the number of plates in the protocol (X). To navigate through multiple plates in a protocol, use the **back and forward arrows** on either side of the label. The keys “Page Up” and “Page Down” can also be used to navigate plates.

If you are working on a multi-plate protocol, refer to this area to see which plate is the “current” plate (plate N of X).

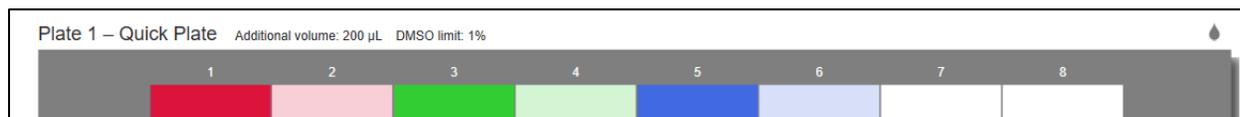
Zoom

To zoom the well plate grid:

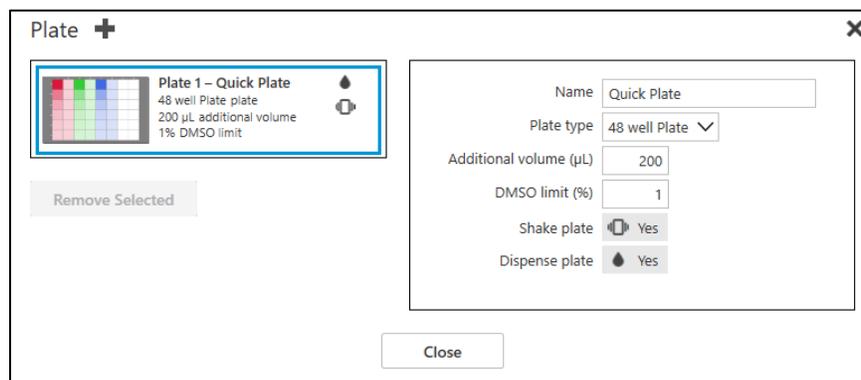
- Move the bar towards zoom out  to see more real estate in the current plate and additional plates in the protocol.
- Move the bar towards zoom in  to focus in on the plate wells.
- Alternately, you can click the icons to zoom in or out by a small amount. The mouse wheel can also be used to zoom in or out while holding the **CTRL** key.

Plate-associated information

Plate-associated information is displayed above each plate.



This information can be changed by clicking  on the menu bar, double-clicking the information area, or by selecting **Edit Plate** from the right-click context menu.



- You can **Name** the current plate. Use this name as your own reference to help you determine which plate to load during runtime.
- Modify the **Plate type** via an updated choice from the drop-down menu.
Note that the plates in a multi-plate protocol do not need to be of the same plate type, nor do they need to have the same additional volume or DMSO limits.
- Change **Additional volume** via manual entry. Additional volume is the volume added to each well of the plate not using the dispenser and is used for concentration and normalization calculations (see [Appendix D—Concentration calculations](#)).
- Change **DMSO limit** via manual entry.
- Click **Shake plate** to keep this plate from shaking even when shaking is enabled for the protocol. Don't shake is always set for 1536-well plates and cannot be changed.
- Click **Dispense plate** to keep this plate from being dispensed during a multi-plate run.

The Plate Edit dialog box allows you to go directly to edit other plates without exiting by selecting a plate icon on the left side of the editor.

- You can also add a new plate directly from the dialog box by clicking . Note that this method also copies the edit plate type, additional volume, and DMSO limit to the new plate.

Saving/canceling your edits:

- To save your edits, click **Apply** or move to another plate.
- To cancel your edits, click **Cancel**. Note that this will only cancel edits to the current plate, but it will not cancel saved changes to previously edited plates. Once you click either of these options, the dialog box closes, and you are brought back to the Main Operating Window.

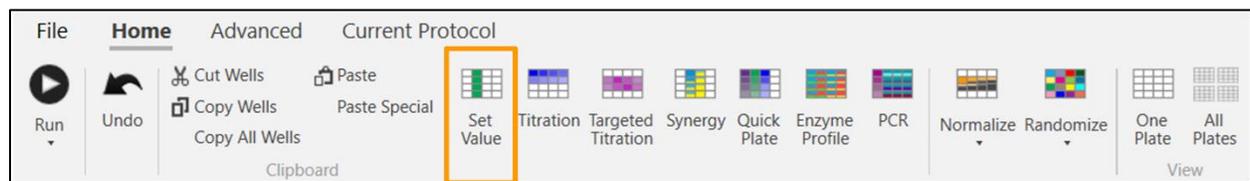
Don't dispense a plate

At the far right of the plate-associated information is a **Droplet** icon  that controls whether a plate is dispensed.



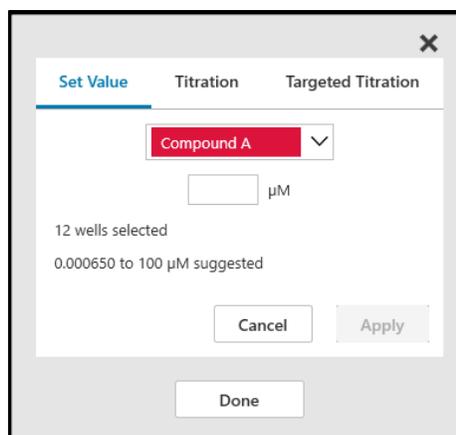
Click the **Droplet**  to initiate **Don't dispense plate** to skip the current plate when running the protocol. Note that this setting is temporary; it is not saved with the protocol.

Set value



If you want to add fluid to wells at a single concentration/volume, as opposed to as a titration series, use the **Set Value** function.

- Select the wells to which you wish to add the fluid.
- On the Home tab, select **Set Value**.



- Select a **fluid** from the drop-down and specify the **concentration** or **volume** of the fluid depending on the dispense mode for the fluid. Ensure that the value is within the valid range. Note that the range minimum is a “hard” limit based on dispenser drop size, but the upper limit is a suggestion to keep the DMSO limit or recommended maximum aqueous dispense volume from being exceeded.

Once you have updated the parameters, click **Apply** (applies the function) and **Done** (closes the panel).



NOTE: A diluted fluid may be suggested if the requested value with the stock fluid concentration would be significantly below the minimum dispense volume. See the [Fluid shift or extend](#) section for more information.

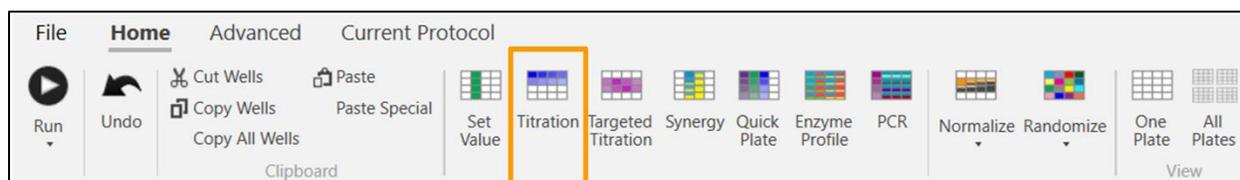
NOTE: For the functions of: **Set Value, Titration, and Targeted Titration:**



After you click **Apply**, if the fluid was previously assigned to any of the selected wells, you will receive a warning message to Replace or Cancel. If you choose Replace, the newly defined values will overwrite the previous values for that fluid in the selected wells. If different fluids were previously assigned to any of the selected wells, the new fluid will be added to, and not replace, the existing fluids.



Titration



This section describes how to set up an evenly-spaced titration series in more detail than the basic titration example section did.

- Select the wells in the plate for titration.
- On the Home tab, select **Titration**. An editing panel is then displayed.

Highest and lowest concentration

Use this selection to specify both the high and low limits for the titration and let the software fill in the remaining points, equally spaced according to the selected distribution.

- Select a **Fluid** from the drop-down list.
- Assign the **Highest concentration** (or volume) parameter.
- Assign the **Lowest concentration** (or volume) parameter.



TIP: Use the **paste function** in the lower-right area of the panel to auto-fill the highest and lowest concentration with the valid range limits. The high value is determined from fluid concentration and DMSO limit or recommended maximum aqueous dispense volume. The low value is limited by minimum dispense volume.



- Select the **Distribution** via the drop-down list - Logarithmic or Linear.
- Assign **Replicates** per level. Each titration level will be replicated the specified number of times. The number of selected wells must be a multiple of Replicates. For instance, if 12 wells are selected, 3 replicates will result in 4 levels.
- Select the **Titration pattern** from the drop-down menu. The graphics illustrate where the Highest and Lowest levels will be placed in the selected area as well as whether the titration is distributed by row or column.

Once you have completed your edits, click **Apply** (applies the function) then **Done** (closes the panel).



NOTE: A fluid shift or extension may be suggested if the requested titration cannot be achieved with the stock fluid concentration specified. See the [Fluid shift or extend](#) section for more information.

Highest concentration or lowest concentration

Use this selection when you wish to specify either the highest or the lowest concentration and apply a specific distribution, such as a 1:4 dilution, to the remaining points.

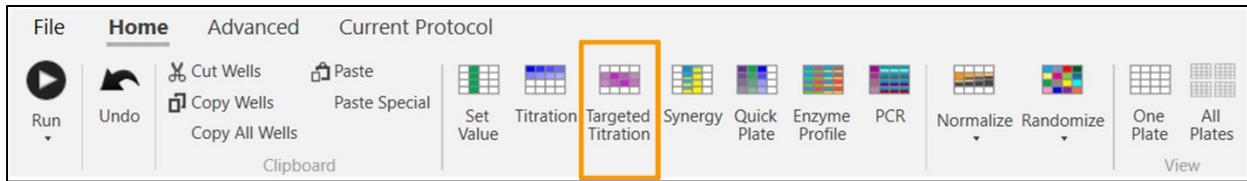
The screenshot shows a dialog box titled "Titration" with three tabs: "Set Value", "Titration", and "Targeted Titration". The "Titration" tab is selected. The dialog contains the following elements:

- Section: "Specify titration using..."
 - Radio buttons: "Highest and lowest concentrations", "Highest concentration" (selected), "Lowest concentration".
- Fluid: "Compound B" (dropdown menu)
- Highest concentration: [] μM
- Lowest concentration: *Insufficient data to calculate*
- Distribution: "1/2 Log" (dropdown menu)
- Replicates per level: "1/2 Log" (dropdown menu)
- Titration pattern: "1:2 (50%)" (dropdown menu)
- 24 wells selected
- Paste range is 0.000325
- Buttons: "Cancel", "Apply", "Done"

- Select a **Fluid** from the drop-down list.
- Assign **Highest concentration** (or volume) parameter if you are using highest or assign **Lowest concentration** (or volume) parameter if you are using lowest.
- Select the desired **Distribution** via the drop-down menu – **1/2 Log**, **1/3 -Log**, **1/4-Log**, **1:2**, **1:3**, **1:4**, **1:N**, or **Linear**. For **1:N**, N is a numerical value greater than 1 (does not need to be an integer). For **Linear**, the value increment is also specified.
- Assign **Replicates** per level. Each titration level will be replicated the specified number of times. The number of selected wells must be a multiple of Replicates. For instance, if 12 wells are selected, 3 replicates will result in 4 levels.
- Select the **Titration pattern** from the drop-down menu. The graphics illustrate where the Highest and Lowest levels will be placed in the selected area as well as whether the titration is distributed by row or column.

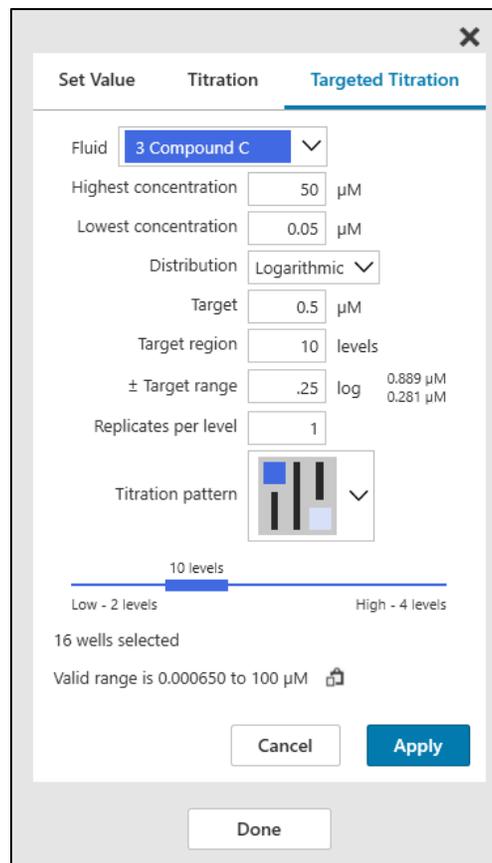
Once you have completed your edits, click **Apply** (applies the function) then **Done** (closes the panel).

Targeted titration



In a targeted titration, the levels are not distributed evenly. The targeted titration includes a user-defined zone where the levels are densely spaced around a value of interest. Targeting the titration around an expected IC50 value, for instance, can improve the IC50 estimate or make the same quality of estimate with fewer wells.

Many of the fields required for Targeted Titration are the same as the fields for Titration. A targeted titration can only be programmed by specifying both the highest and lowest concentrations. The additional fields for targeted titration are related to specifying the target range.

A screenshot of a configuration dialog box titled 'Targeted Titration'. The dialog has three tabs: 'Set Value', 'Titration', and 'Targeted Titration', with the latter being the active tab. The configuration includes:

- Fluid: 3 Compound C (dropdown)
- Highest concentration: 50 μM
- Lowest concentration: 0.05 μM
- Distribution: Logarithmic (dropdown)
- Target: 0.5 μM
- Target region: 10 levels
- \pm Target range: .25 log, with sub-values 0.889 μM and 0.281 μM
- Replicates per level: 1
- Titration pattern: A visual representation of a titration curve with a highlighted target region (dropdown)

At the bottom, there is a horizontal scale for '10 levels' with 'Low - 2 levels' on the left and 'High - 4 levels' on the right. Below the scale, it states '16 wells selected' and 'Valid range is 0.000650 to 100 μM '. At the very bottom are 'Cancel', 'Apply', and 'Done' buttons.

- Assign the **Target** concentration value – This value will become the center of the densely spaced region and must lie within the Highest and Lowest concentrations. This value might be, for example, the expected IC50.
- Assign the number of wells in the **Target region**. These wells will lie on either side of the target concentration value.
- Assign the **+/- Target range**. This defines the concentration or volume range above and below the target value that will be covered by the specified number of wells.

When the panel contains sufficient information, the diagram will indicate how many wells are planned for each “tail” region above and below the target region and will show where the target region falls within the overall titration. When calculating the targeted titration, the software will always apply one well to each end concentration (the highest and lowest) and then any wells remaining after being distributed to the target region will be evenly spaced within each tail region.

Fluid shift or extend

If the titration specified in Titration, Targeted Titration, Synergy, Quick Plate, Enzyme Profile or PCR would result in well volumes significantly below the minimum well dispense volume, a shifted or extended fluid will be suggested. The suggested fluid will be an integral dilution of the selected fluid: 1:2 through 1:9, 1:10 through 1:90, etc. and less than 1:10,000 dilution.

A diluted fluid may also be suggested from Set Value, PCR, etc., if the well volume would be significantly below the minimum well dispense volume.

If the diluted fluid is accepted, a new fluid will be automatically added to the fluid list and the titration will be completed using that fluid. In this case, you must create a new stock fluid based on the original stock fluid, following the recommended dilution. For a 1:20 dilution, it is expected that you will add 1 part of the original fluid to 19 parts diluent, to create a 5% solution of the original stock fluid.

If the titration can be accomplished with one fluid spanning the full concentration range requested, you will be offered a Fluid Shift. If, however, the concentration range is greater than that which can be achieved with a single fluid, you will be offered a Fluid Extension and the titration will be comprised of both the original stock fluid and the diluted extended fluid.

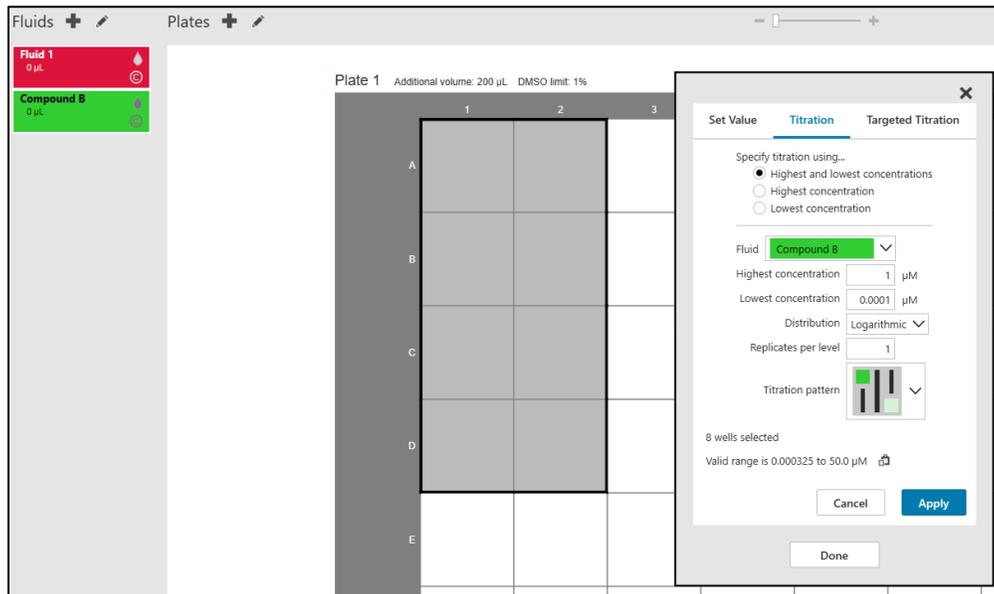
If the diluted fluid is not accepted, the titration will be completed using the original stock fluid, and the wells that were requested with concentrations/volumes outside the accessible range will be flagged as being below the minimum dispense volume.



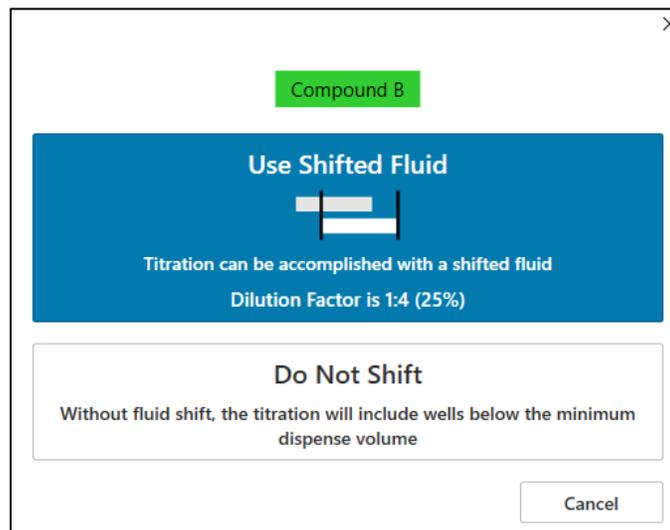
NOTE: Diluted fluids are permanently associated with the original stock fluid. If the original fluid is modified (name, concentration, etc.) the diluted fluid will also be changed.

Fluid shift

For example, if the following titration is specified:



You will be prompted to create a shifted fluid, diluted 1:4 (25%).



This creates a new fluid and the titration will use the new fluid.



Fluid extend

If the limits of the titration are modified as shown:

Fluid	Compound B	▼
Highest concentration	5	µM
Lowest concentration	0.00001	µM

The titration cannot be accomplished with a shifted fluid and an extended fluid is offered.

Compound B

Use Extended Fluid



Titration can be accomplished using an extended fluid in addition to the selected fluid

Dilution Factor is 1:40

Do Not Extend

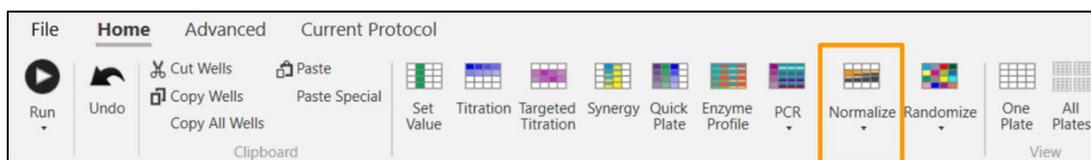
Without fluid extension, the titration will include wells below the minimum dispense volume

Cancel

This creates a new fluid and the titration will use both the original and new fluids. If possible, the maximum DMSO % of the extended fluid will not exceed that of the original fluid in the titration. And, the maximum DMSO % of the extended fluid will never exceed that specified in the plate parameters.



Normalize



For some assays, it may be advantageous to normalize the DMSO or aqueous surfactant content of the wells. Because of the digital nature of dispensing from the dispenser, without normalization, not all wells will contain the same DMSO or surfactant content. It may also be desired to normalize the total volume of all fluids in a well, regardless of fluid class.

It is typical for the DMSO content in dispensed wells to range from 0.00001%-1%. For many assays it is preferred to minimize the DMSO content, but for others, normalizing the DMSO content to the highest level or some intermediate level may be desired.

It may also be advantageous to normalize the dispensed surfactant content for certain assays.



TIP: The dispense range for Normalization Fluid is specified in [Appendix E—Dispense specifications](#). Normalization cannot dispense a quantity lower than 1nL per well. If normalizing a well requires dispensing less than 1nL of fluid, no fluid will be added to that well.

- Select the wells for to be normalized with a specific fluid class. Wells may be selected from multiple plates. Note that Normalize is not accessible unless you have selected wells.
- In the Home tab, select **Normalize**.
- To normalize a class of fluids, select one of the available normalization options via a radio button on the Normalize Fluid Class tab.

Select the fluid class to normalize from the list. A well can be included in multiple normalization groups, but there can be only one group for each fluid class.

The screenshot shows a dialog box titled "Normalize Selection" with a close button (X) in the top right corner. It has two tabs: "Normalize Fluid Class" (which is selected and underlined) and "Normalize Total Volume". Under the "Normalize Fluid Class" tab, there is a "Fluid class" dropdown menu currently set to "DMSO". Below this are three radio button options: "Normalize to highest class volume." (which is selected), "Normalize to specified volume.", and "Normalize to % of total volume.". At the bottom of the dialog are two buttons: "Cancel" and "Normalize".

Normalize to highest class volume: This will add normalization fluid to selected wells such that the normalization fluid + titration fluid volumes all equal the highest volume well in the selection. It will not add normalization fluid to the highest volume well.

Normalize to specified volume: This will add normalization fluid to selected wells so that the normalization fluid + titration fluid volume is equal to the specified volume, unless the titration fluid volume already equals or exceeds the specified volume, in which case no normalization fluid will be added.

Normalize to specified % of total volume: This will add normalization fluid to selected wells such that the normalization fluid + titration fluids all equal the specified % of total well volume. It will not add normalization fluid to wells that contain titration fluid equal to or in excess of the specified % of total volume.



NOTE: If normalizing to **% of total volume**, each well is normalized to the volume appropriate for the well. This includes additional volume that may be different for each plate in the normalization group.

- To normalize the total volume of a well – all fluids regardless of fluid class, and including **Additional volume** (see [Plate-associated information](#)) – select one of the normalization options on the Normalize Total Volume tab.

The fluid class to use for normalization must be selected. A well can be included in multiple normalization groups, but there can be only one group for each fluid class.

The screenshot shows the same "Normalize Selection" dialog box, but with the "Normalize Total Volume" tab selected and underlined. Under this tab, there are two radio button options: "Normalize to highest total total volume." (which is selected) and "Normalize to specified total volume.". Below these is a "Normalize using fluid class" dropdown menu currently set to "Aqueous+Tween 20". At the bottom are "Cancel" and "Normalize" buttons.

Normalize to highest fluid volume: This will add normalization fluid to selected wells such that the normalization fluid + titration fluids + additional volume equals the highest total volume well in the selection. It will not add normalization fluid to the highest total volume well.

Normalize to specified volume: This will add normalization fluid to selected wells so that the normalization fluid + titration fluids + additional volume is equal to the specified volume, unless the additional volume + titration fluid volumes already equals or exceeds the specified volume, in which case no normalization fluid will be added.

- Click **Normalize** to normalize the selected wells in the protocol or **Cancel** to abort normalization.



TIP: Wells that are normalized together using the same options are referred to as a **normalization group**.

The normalized wells are subsequently displayed with a **black tick mark** in the lower-left corner of the well, indicating well normalization.

When you move your cursor over a normalized well, you see the normalization details in the displayed information:

Plate 1 Well B9
DMSO Volume: 186 nL (0.911 %)
Aqueous Volume: 187 nL
Additional Volume: 20.0 μ L
Total Volume: 20.4 μ L

■ **Compound A** **DMSO**
Concentration: 24.5 μ M
Volume: 49.9 nL

■ **Compound B** **a+Tw**
Concentration: 138 ng/mL
Volume: 187 nL

■ **DMSO Normalization**
Volume: 136 nL

[Normalize DMSO to highest volume: 186 nL](#)

The Normalization display in the Main Operating Window below the Fluid list shows up if the plate has a normalization scheme. When the mouse is moved over the  icon, the total normalization volume for each fluid class is displayed.

Total volume		Dispense order
DMSO	12.3 μ L	Before fluids <input type="button" value="v"/>
a+Tw	2.2 μ L	

- Set the **Dispense order** for the normalization fluids. Normalization fluids can be dispensed strictly before or after the fluids or they can be dispensed with the fluids.



NOTE: When dispensing a protocol using high-volume cassettes, the Dispense order may cause dispensing to alternate between cassette types.

- The **dispense icon**  allows you to not dispense all normalization fluids.

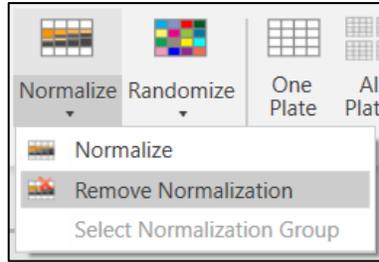
Show normalization

See the [Show Normalization](#) section in [Show](#).

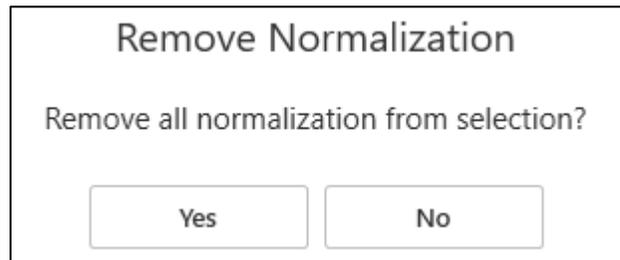
Remove normalization

Use this function to remove from the **selected wells** any normalization you previously set up. If there are no normalized wells in the selected wells, this function is grayed-out and not accessible.

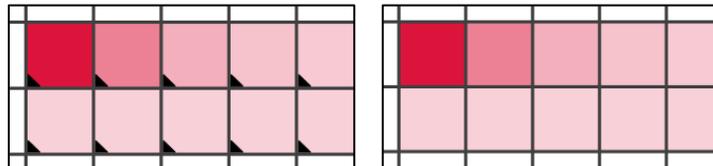
- Select the wells from which you want to remove normalization.
- Click **Remove Normalization**. This function is also available from the right-click contextual menu.



- A dialog box is then displayed, asking you for confirmation to remove all normalization from selected wells.



- Click **Yes** to confirm the removal of normalization or **No** to abort the function. If you cancel, the normalization is left in place (see below, left). If you give confirmation, the cells marked for normalization lose their tick marks, and the normalization is removed (see below, right).



Select normalization group

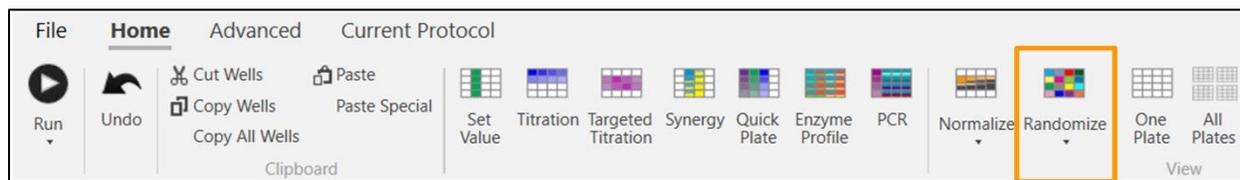
This function shows you which wells are normalized together following the same options. For instance, if you used “normalize to highest fluid volume” this function will show you which wells are included in this calculation. This is important if you specify and then later wish to view multiple separate groups that are all normalizing to unique highest values.

- Select a well that is normalized.

- Click Select Normalization Group.
- If the well is included in multiple normalization groups (different fluid classes), indicate which group to select.

The wells that are normalized together become selected.

Randomize



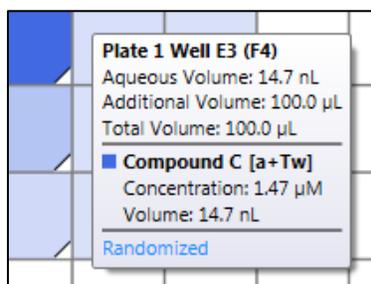
Once a plate has been configured, the wells can be reordered randomly on the plate. This is a useful way to minimize the impact of systematic assay effects on the output data.



NOTES: Randomization is a “per plate” feature. It is not possible to create separately randomized regions within a single well plate. The selected wells on the plate will be randomized. If you later wish to add more randomized wells to the plate, they will be added to the rest of the randomized wells with all randomized together as one group.

- Select the wells desired for randomization.
- Click **Randomize** to randomize the selected wells.

The randomized wells are subsequently displayed with a **white tick mark** in the lower-right corner of the well indicating well randomization. In addition, when you move your cursor over a normalized well, you see the “**Randomized**” tag in the displayed information:



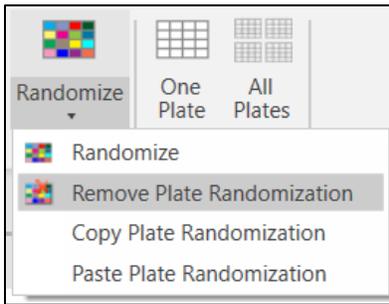
Show plate randomization

See the [Show Randomized](#) section in [Show](#). Note, if you are viewing wells as randomized, some other functionality will be limited. Change the Show option to unselect Show Randomized to enable full control.

Remove plate randomization

For the current plate in a protocol, use this function to **remove all randomization from the current plate**. If there are no randomized wells in the current plate, this function is grayed-out and not accessible.

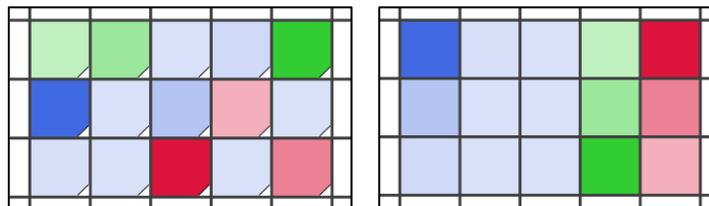
- Click Remove Plate Randomization.



- A dialog box is then displayed, asking you for confirmation to remove the randomization scheme.



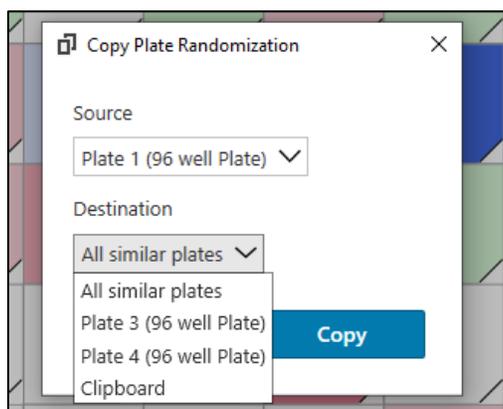
- Click **Yes** to confirm the removal of the randomization scheme or **No** to abort the function. If you give confirmation, the cells marked for randomization lose their tick marks, and the randomization scheme is removed.



Copy plate randomization

Use this function to copy the randomization scheme from one plate to another within the same protocol. You can only copy randomization to plates with the same dimensions as the source plate.

- Click **Copy Plate Randomization**.
- A dialog box is then displayed, asking you for the source plate and the target plate(s). Use the drop-down menu to specify source and destination. Note that you have the option to copy the randomization scheme from the source plate to all similar plates in the protocol.

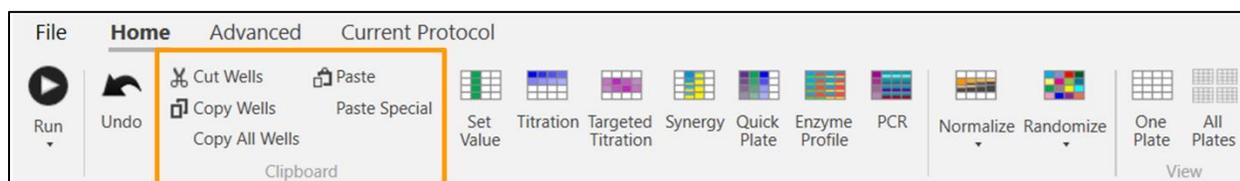


- Click **Copy** to perform the copy or **Cancel** to abort it. If you perform the copy to **All similar plates**, all plates with the same dimensions as the source plate will now have the same randomization scheme. You can also copy to the **Clipboard** and paste the randomization scheme to a plate in another protocol.

Paste plate randomization

Use this function to paste a plate randomization scheme that was copied to the Clipboard from another plate or protocol.

Copy & paste basics



You can copy and paste wells to duplicate them within a plate or copy them to another plate. You can also copy and paste all wells to duplicate the well contents in the current protocol or move them to another protocol.



NOTE: Pasting wells overwrites any current information in the target wells or plates.

Copy Wells/Paste Wells

- Select the desired well(s) on the plate, using the techniques discussed previously.
- Click **Copy Wells** or **Cut Wells** on the Home tab. “Cutting wells” makes the selected wells revert to empty.
- If the wells are being pasted into another plate, navigate to that plate to make it the current, target plate.
- Set the **Paste target** by clicking the upper-left-most well of the target area.

- Click **Paste** to bring the copied wells into the target area. The pasted wells will overwrite, without warning, any other well contents. If there is not enough contiguous space in the target area, you will get a paste warning message about truncation; at that point, you can cancel the Paste or continue with the truncation.



NOTE: All fluids and normalization are included in copy, cut, and paste. When copying to an external application (e.g. Excel), however, normalization is not included.

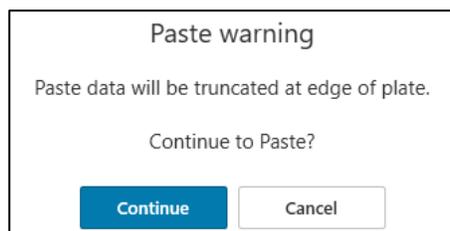
Copy All Wells

Use this function to copy all well fluid contents from one plate to another. The plate types do not need to be the same. If a less dense plate is copied to a higher density plate, all the wells will be copied. Conversely, if a higher density plate is copied to a lower density plate, the wells will be truncated, with a warning. When the contents of a plate are copied and pasted into another plate, all fluids and normalization are copied but randomization is not copied. Additionally, target plate parameters such as additional volume and DMSO limit, are not copied.

- Navigate to the plate that you want to copy.
- Click **Copy All Wells**.
- Navigate to the target plate. If you want to set up a new plate with an existing one as a source, click the  sign to create the new plate.
- Click **Paste** to paste the copied wells into the target plate.

If the target plate is not empty you will be warned before pasting. If you choose not to replace all fluids on the plate, the copied fluids will be added to existing plate fluids, unless the same fluid occurred in both the source and target plates, in which case that fluid will be replaced.

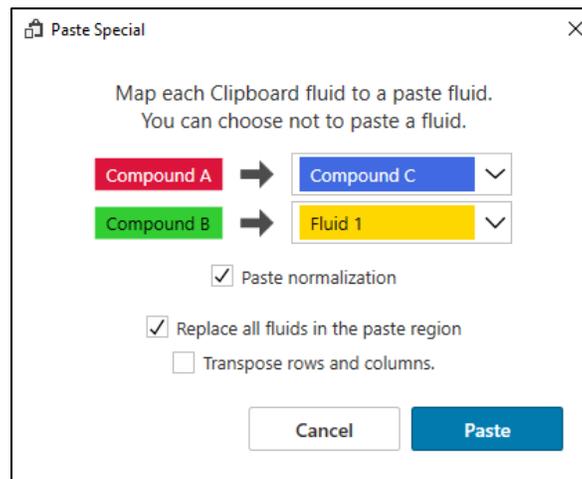
If there is a mismatch in the plate size being copied vs. the target plate, you will receive a warning message. At that point, you may cancel or continue the paste. The result may be that the wells all fit and become one quadrant of the new plate, or the wells will be truncated.





Paste Special

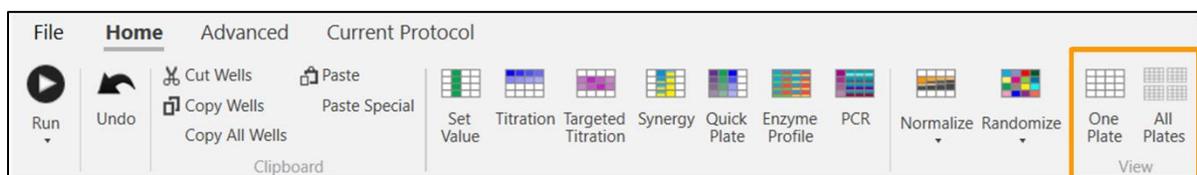
Paste Special is similar to Paste, but with the option of mapping each fluid to a different fluid. You can also choose not to paste individual fluids if you wish to leave selected wells of a specific fluid blank in the pasted region.



- If the source wells include normalization, you can also choose to **Paste normalization**.
- If the **Replace all fluids in the paste region** box is not checked, the pasted fluids will be added to fluids that already exist in the target plate, unless the same fluid occurred in both the source and target plates, in which case that fluid will be replaced.
- It is optional to **Transpose rows and columns**.

You cannot map fluids that have different unit types (molarity vs. mass/volume) or dispense modes (by concentration/volume).

View



This **Home tab** function is a complement to [Zoom](#).

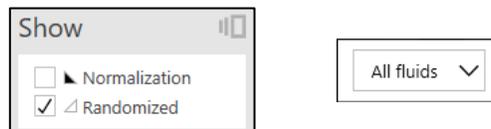
- Click **One Plate** to size and center the current plate in view.

- Click **All Plates** to bring all plates into view.

 **TIP:** Double-click a plate border to size and center that plate in view.

Show

The **Show** area of the Main Operating Window allows you to filter what is displayed in the Well plate. Move the mouse over the  icon to show the fluid filter selection list.



Show Normalization

If normalization is used in the protocol, the checkbox allows you to view or not view the normalization fluid in the well plate.

Show Randomized

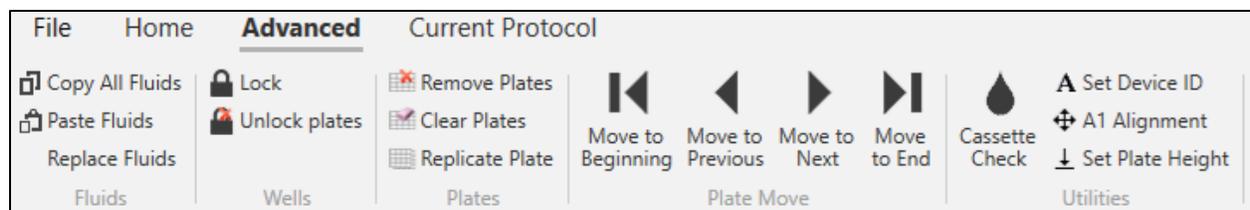
If randomization is used in the protocol, the checkbox allows you to view or not view the plates randomized.

Show Fluids

Click and select from the drop-down menu to indicate which fluids in the protocol's fluid list should be displayed on the plate.

- All fluids
- Dispensed fluids
- Only the fluid selected from the list

Plate functions



Remove Plate

- Navigate to the plate that you want to remove.
- Click **Remove Plate** on the **Advanced** ribbon tab. A dialog box is displayed, asking you to confirm the removal or cancel it.

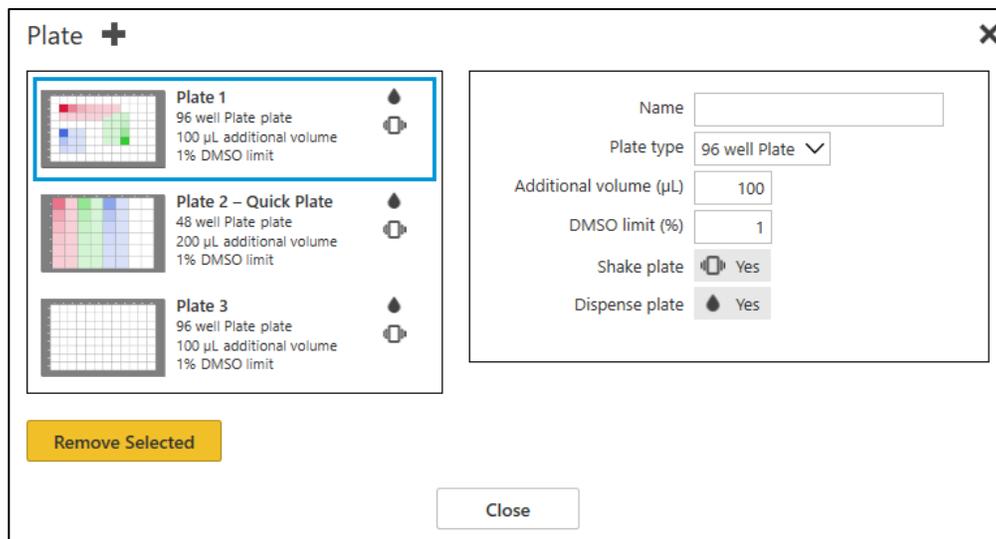


- Click **Remove** to remove the current plate. Click **Cancel** to stop the removal.

Removing plates with the plate editor

You can remove plates by selecting the  icon next to **Plates**.

- The current plate is always selected.
- You can toggle plate selection for an individual plate by selecting the icon of a given plate.
- Plates can be removed by clicking the plate icon and clicking **Remove Selected**.



- Confirm plate removal by selecting **Remove** or **Cancel** to go back to the plate editor.



Removing multiple plates

You can remove multiple plates by holding down the **CTRL** key, selecting multiple plates by clicking the plate name (e.g. Plate 1, Plate 2, etc.), right clicking the last plate selected, and selecting **Remove Plates**.



Clear Plate

- Navigate to the plate from which you want to clear all fluids. You can select multiple plates by holding down the **CTRL** key and selecting wells of interest on multiple plates.
- Click **Clear Plate** on the **Advanced** ribbon tab. A dialog box is displayed, asking you to confirm the clear or cancel it.



- Click **Clear** to clear the current plate of all fluids, turning it into an "empty" plate. Click **Cancel** to stop the clear. Note that clearing the plate does not remove fluids from the Fluids List.

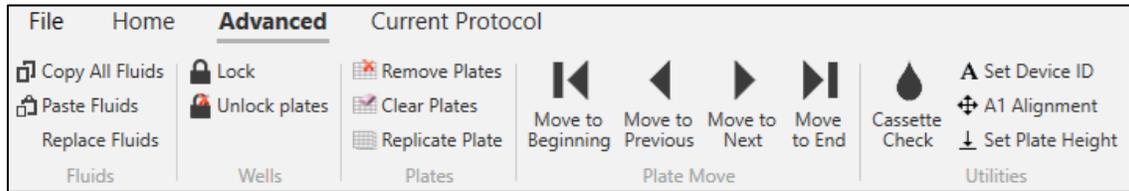
Replicate Plate

This function is handy for an automatic, one-step creation of a new plate with the exact characteristics of the current plate, including Normalization and Randomization. In contrast, the [Copy All Wells](#) function described previously allows you to copy the well fluid contents from one plate to another plate without copying randomization and without constraints on the plate types.

- Navigate to the plate that you want to replicate.

- Click **Replicate Plate** on the **Advanced** ribbon tab. A new plate with the exact characteristics is then created and appended to the end of the plate series; the new plate becomes the current plate.

Ribbon—Advanced tab



The Advanced ribbon tab contains features in addition to the [Plate functions](#).

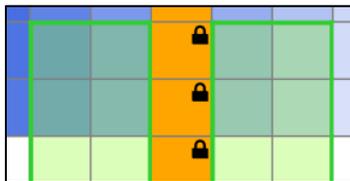
Copy/Paste Fluids from the fluid list

This function allows you to copy a Fluid List from one protocol to another or from a separate list of fluids, such as from a Microsoft Excel file (refer to [Appendix B—Moving data between the dispensing software and other programs](#) for information about this function). This function is handy when you are creating a new protocol that is similar to an existing one, and you do not want to enter a new Fluid List from scratch.

- Open the source protocol, and then click **Copy All Fluids**. This copies the current Fluid List to the clipboard.
- Bring up the target protocol and click **Paste Fluids**. This brings in the entire Fluid List from the source, modifying any changed parameters of existing fluids in the target protocol (matched by name) and appending new fluids to the Fluid List.
- **Replace Fluids** will replace each existing fluid with pasted fluids in the order they occur. All fluid parameters are changed to match the source. This allows you to use the same protocol repeatedly, but with fluids of different properties (name, stock concentration, source location names). Certain fluid properties cannot be changed (unit type).

Lock and unlock wells

Selected wells can be locked to prevent inadvertent changes to well fluids. Locked wells show a lock icon and cannot be selected.



If a Paste operation would overwrite a locked well, an error is displayed.



Plate moves

Use the ribbon buttons to move the current plate relative to other plates in the protocol. These features are also available from the menu [Plate-associated information](#) bar.

Utilities

You must have a device connected to use these features.

- You can check the dispense head(s) on a cassette by pressing the **Cassette Check** button. From the dialog box press **Check** to test a cassette and display results.
- **Set Device ID** allows you to give a unique name to the device which is included in reports run on that device.
- A1 Alignment is used to adjust the A1 position. Refer to the dispenser **Operating Manual** by navigating to the **File** → **Documentation** menu of the software for more detail.

Ribbon—Current Protocol tab

The screenshot shows the 'Current Protocol' ribbon tab with the following controls:

- Concentration (molarity): μM (dropdown)
- Concentration (mass): ng/mL (dropdown)
- Volume: nL (dropdown)
- Units: (text label)
- Shake during dispense:
- Shake after each dispensehead:
- Volume Threshold (nL): 100 (input field)
- Duration (sec): 1 (input field)
- Label: (text input field)
- Last run summary: (button)
- View: (button)
- Dispense fluids in successful single cell wells:
- Cell Dispensing: (text label)

The functions on this tab apply to the current protocol, and any changes are saved in the current protocol file.

Concentration and Volume units

Well fluids are displayed using the specified units of concentration and volume.

Shake during dispense



TIP: It is highly recommended to **shake during dispense** for cellular assays when dispensing DMSO, particularly for larger volumes, to prevent DMSO from sinking to the well bottom and creating a local excess of DMSO and compound.

To aid in mixing the dispensed fluids you can program the dispenser to shake the well plate while it is dispensing fluids. The dispenser shaking intensity parameters were optimized by plate type and are not user selectable. The duration of the shaking for each dispense event depends on the amount of dispensing occurring; wells with large amounts of fluid will be shaken longer because it takes more time to dispense them. This type of shaking will occur for every well dispensed for all fluids, as long as the dispensed volume exceeds the programmed Volume Threshold. Manually enter a **Volume Threshold** (in nL) and check the **Shake during dispense** box. When dispensing DMSO onto plated cells, it is recommended that shake during dispensing is turned on for any volumes greater than 100nL.



NOTE: Shake during dispense is automatically enabled when a low-density plate (48 wells or fewer) is created, however, it can be disabled.



NOTE: Shaking is always disabled for 1536-well plates, regardless of the protocol setting.

Shake after each dispense head



TIP: If not mixing off-line after dispensing, in addition to **shake during dispense** it is also recommended to **shake after each dispense head** on each plate for DMSO dispensing. Depending on the dispensed volumes, additional volume, and plate type you should determine the shake duration during assay development or by visually testing the mixing of dye/DMSO in your well plate. Shake after each dispense head is not recommended when dispensing aqueous fluids.

To aid in mixing the dispensed fluids, you can program the dispenser to shake the well plate after each dispense head has completed dispensing on the current plate. The shaking intensity parameters were optimized by plate type and are not user selectable.

Manually enter a shake **Duration** (in seconds) and check the Shake after each dispense head box. The duration is typically 3 seconds for a 384-well plate or 20 seconds for a 24-well plate.



NOTE: Shaking is always disabled for 1536-well plates, regardless of the protocol setting.

Dispense fluids in successful single cell wells

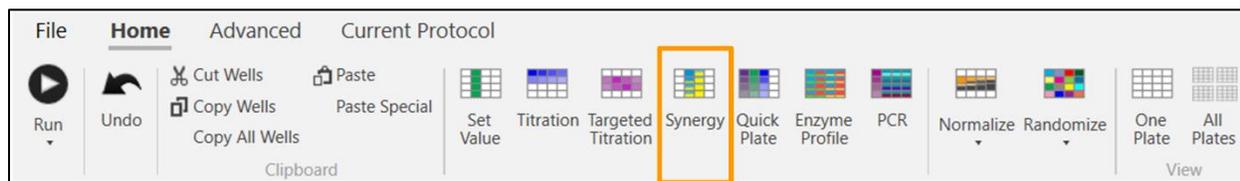
Check the **Dispense fluids in successful single cell wells** box to dispense fluids only in wells where a single cell was successfully isolated. To dispense fluids regardless of single cell status in a given well, leave this box unchecked.



NOTE: Wells determined to not contain a single cell contain an undefined number of cells: $n=0$ or $n>1$

Special features

Synergy



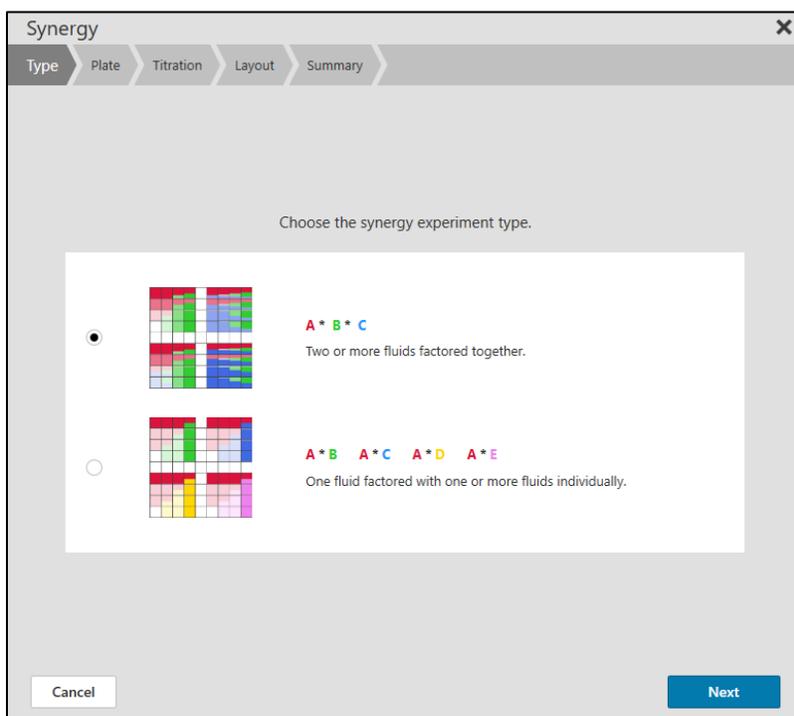
The **Synergy** function allows you to create a plate design for a drug-drug interaction or synergy experiment where multiple compounds are placed in the same well for many wells in a set of orthogonal titration sequences. The dispenser makes it especially easy to create synergy sequences because of the non-contact digital dispensing. Each compound is added to every well independently, and multiple compounds can be added to the same wells with no additional user effort. Synergy wells can also be normalized and randomized, if desired.

On the Home Tab, click **Synergy** to create a synergy sequence. You then navigate and enter parameters in a series of pages to create the sequence. Once you have completed the procedure, a new plate or plates with the synergy layout is added to the protocol.



NOTE: When using the Synergy function, it is not necessary to pre-select the wells to which you wish to apply fluids. The Synergy function automatically creates new plates with the programmed well content and ignores any wells that you may have already selected in an existing plate.

Type page



There are two types of Synergy experiments available. The first type factors all of the fluids together. The second type of experiment factors a single fluid against a group of other fluids one at a time.

- Select the synergy experiment type.

Click **Next** to go onto the next step or **Cancel** to abort the in-process layout procedure.

Plate page

Synergy

Type Plate Titration Layout Summary

Plate type: 96 well

Additional volume (µL): 100

DMSO limit (%): 1

Click row and column headers to exclude wells from the synergy layout.

	1	2	3	4	5	6	7	8	9	10	11	12
A												
B												
C												
D												
E												
F												
G												
H												

Cancel Back Next

Synergy

Type Plate Titration Layout Summary

Plate type: 96 well

Additional volume (µL): 100

DMSO limit (%): 1

Click row and column headers to exclude wells from the synergy layout.

	1	2	3	4	5	6	7	8	9	10	11	12
A												
B												
C												
D												
E												
F												
G												
H												

Primary synergy grid must fit in each plate region; regions are formed by excluded rows and columns.

Cancel Back Next

- Assign the **Plate type**, **Additional volume**, and **DMSO limit** in the fields provided at the top of the dialog box – the current plate's parameters are the defaults.



NOTE: **DMSO limit** is only displayed when DMSO fluids are in the protocol.

- If desired, click **the row letter(s) and/or column number(s)** to select the rows and columns of wells that should be **excluded** from the synergy layout. In the figure above, the user excluded rows A, H and columns 6, 7 from the layout. To de-select a row or column, just click the row letter or column number again. This can be used to leave rows or columns empty for use with controls.
- If rows or columns have been excluded, you have the option to keep the primary Synergy grid from crossing excluded wells (if possible).

Click **Next** to go onto the next step or **Cancel** to abort the in-process layout procedure.

Titration page – A*B*C type

The titration page is different depending on the type of synergy experiment selected on the Type page.

The screenshot shows the 'Synergy' application window with the 'Titration' tab selected. The interface is titled 'Specify titrations for synergy fluids.' and indicates '8 rows available per plate' and '12 columns available per plate'. There are two configuration panels side-by-side. The left panel is for 'Compound A' and the right panel is for 'Compound B'. Each panel has a 'Set Value' tab, a 'Titration' tab (which is active), and a 'Targeted Titration' tab. Under the 'Titration' tab, there is a 'Titration levels' input field. Below that, a section 'Specify titration using...' has three radio button options: 'Highest and lowest concentrations' (selected), 'Highest concentration', and 'Lowest concentration'. A 'Fluid' dropdown menu shows the selected compound. Below are input fields for 'Highest concentration' and 'Lowest concentration' in μM . There is a checked checkbox for 'Include "0" value', a 'Distribution' dropdown set to 'Logarithmic', and an 'Order' dropdown. The left panel's order is 'High to Low' and its valid range is '0.00130 to 100 μM '. The right panel's order is 'Low to High' and its paste range is '0.000650 to 50.0 μM '. A plus sign button is visible on the right edge of the configuration area. At the bottom are 'Cancel', 'Back', and 'Next' buttons.

The first two synergy fluids will form the **primary synergy grid**. The first fluid will flow down the non-excluded rows and the second fluid will flow across the non-excluded columns. The number of available rows and columns is displayed for reference.

Use the plus  button on the right edge to add additional fluids from the fluids list to the synergy.

This screenshot shows the 'Synergy' application window with the 'Titration' tab selected, now showing three configuration panels. The first two panels are identical to the previous screenshot, for 'Compound A' and 'Compound B'. The third panel on the right is for 'Compound C'. It has the same layout as the other panels, with 'Highest and lowest concentrations' selected, 'Logarithmic' distribution, and 'Low to High' order. Its valid range is '0.00130 to 50.0 μM '. A plus sign button is visible on the right edge of the configuration area. At the bottom are 'Cancel', 'Back', and 'Next' buttons.

For each synergy fluid, choose Titration, Targeted Titration, or Set Value.

- Specify the **number of Titration levels** via manual entry.
- Fill out all other fields according to instructions in the [Titration](#), [Targeted titration](#), or [Set value](#) sections.
- You can **Include “0” value** by clicking the check box.

Click **Next** to go onto the next step or **Cancel** to abort the in-process layout procedure.



NOTE: A fluid shift or extension may be suggested. See the [Fluid shift or extend](#) section for more information.

Titration page – A*B A*C A*D A*E type

The titration page is different depending on the type of synergy experiment selected on the Type page.

The primary titration fluid is used for non-excluded rows of the synergy grid. A separate grid will be created for each secondary fluid. The secondary fluid will flow across non-excluded columns.

- **Select the secondary fluid(s)** to be used by clicking the checkbox next to the fluid name (below left). You can de-select an already-selected fluid by re-clicking the checkbox to clear it. Additionally, you can select **all fluids** by clicking the **Fluids** checkbox at the top of the list.



NOTE: You cannot select secondary fluids with different unit types (molarity vs. mass/volume) or dispense modes (by concentration/volume) for the same Synergy experiment.

For both the primary and secondary synergy fluids, choose **Titration**, **Targeted Titration**, or **Set Concentration** (or **Volume**).

- Specify the **number of Titration levels** via manual entry.
- Fill out all other fields according to instructions in the [Titration](#), [Targeted titration](#), or [Set value](#) sections.
- You can include “0” value by clicking the check box.

Click **Next** to go onto the next step or **Cancel** to abort the in-process layout procedure.



NOTE: A fluid shift or extension may be suggested. See the [Fluid shift or extend](#) section for more information.

Layout page

Choose how to arrange synergy titrations on plates.

 Layout does not leave empty wells.

 Titration levels determine the number of rows and columns in a layout grid.
Each plate contains a single grid.

 Titration levels determine the number of rows and columns in a layout grid.
Each plate contains multiple grids.

Layout grids by rows.

Layout grids by columns.

Cancel Back Next

Choose **how to arrange the synergy titrations** on the plate. Select a layout option by clicking the associated button(s):

- **Layout does not leave empty wells:** This option fills in all the wells along the columns, starting in A1, and does not leave empty rows or columns. It might not result in neatly ordered titration levels. Wells are filled by row or column as specified.

For the remaining options, titration levels of the first 2 fluids determine the number of rows and columns in the layout grid. They start by creating a primary synergy grid from the first two fluids, repeating that grid combined with the remaining fluids as needed.

- **Each plate contains a single grid:** Only a single synergy grid will be placed on each plate.
- **Each plate contains multiple grids:** If multiple copies of the primary synergy grid can be placed on a plate, they will be placed by row or column as specified.
- **Each plate contains multiple grids, with plate breaks:** Plate breaks will be made to keep titration sequences together. For example:

If 9 primary synergy grids fit on each plate.

If 3rd fluid has **5 levels** and 4th fluid has 3 levels, **5 grids** will be placed on each plate.

If 3rd fluid has **2 levels**, 4th fluid has **4 levels**, and 5th fluid has 3 levels, **8 grids** will be placed on each plate.



NOTE: Not all of the layout options may be available, depending on the synergy Type, Titration options and Plate options.

Summary page

32 Wells
1 Plate

Press Finish to create the synergy titrations.

Normalize Wells

Randomize each plate

Cancel Back Finish

In the final page you are given a summary of the number of wells and plates needed and an illustration of what the well contents in the Synergy plate(s) will look like. The mouse can be moved over individual wells to display contents.

Normalize

If you want to normalize all of the wells in the synergy experiment, click the **Normalize wells** button and specify the normalization (see [Normalize](#)).

Randomize

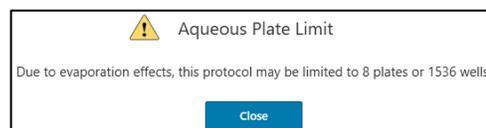
If you want to randomize all of the wells in the synergy experiment, click the **Randomize each plate** checkbox. Each plate will be randomized separately.

Click **Finish** to create the synergy titrations. **Back** and **Cancel** are also available.

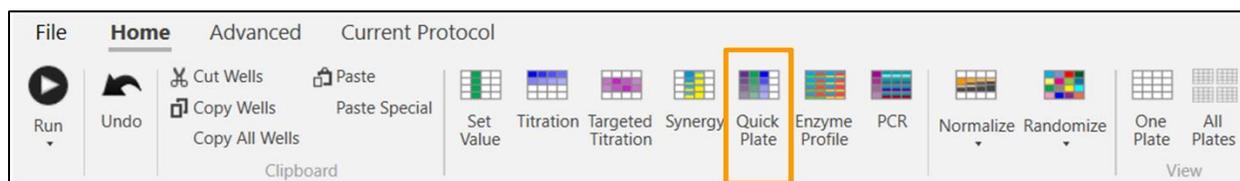
Once you click **Finish**, the new Synergy plate (or plates) is added to the protocol and becomes the current plate.



NOTE: If Aqueous fluids are in use for the Synergy wizard and more than 8 plates are generated a warning message is displayed.



Quick plate



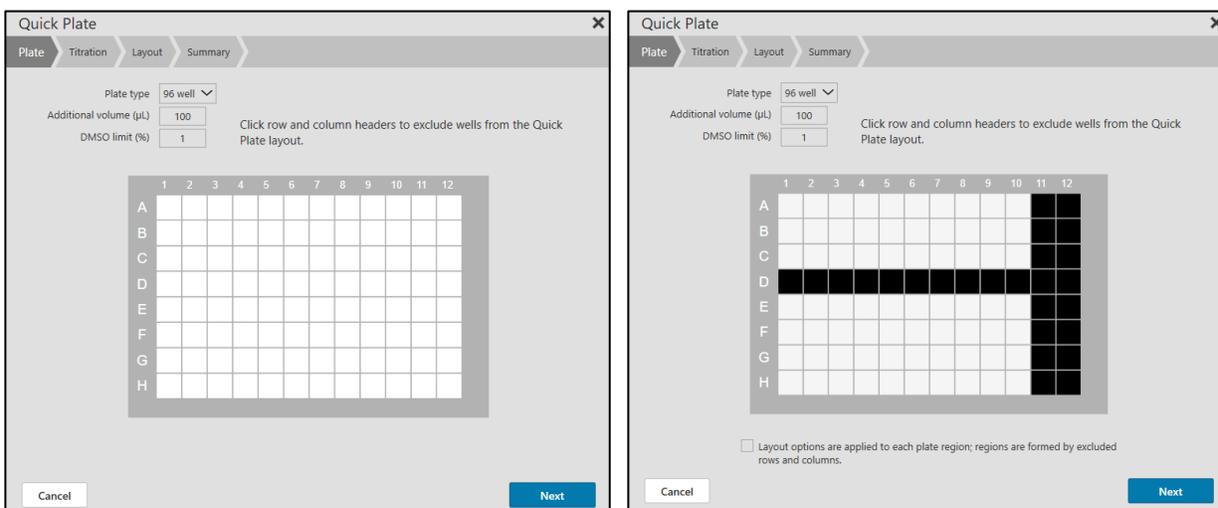
The **Quick Plate** function allows you to create a group of identical titrations using multiple different fluids. This is useful when you want to make identical titrations with many fluids in the fluid list in only a few steps. The titrations can span multiple plates but must have the same concentrations and number of replicates.

On the Home Tab, click **Quick Plate**. Similar to [Synergy](#), you then navigate and enter parameters in a series of dialog boxes to create the plates. Once you have completed the procedure, the new plates are added to the protocol.



NOTE: When using the Quick Plate function, it is not necessary to pre-select the wells to which you wish to apply fluids. The Quick Plate function automatically creates new plates with the programmed well content and ignores any wells that you may have already selected in an existing plate.

Plate page



- Assign the **Plate type**, **Additional volume**, and **DMSO limit** in the fields provided at the top of the dialog box – the current plate's parameters are the defaults.



NOTE: **DMSO limit** is only displayed when DMSO fluids are in the protocol.

- If desired, click **the row letter(s) and/or column number(s)** to select the rows and columns of wells that should be **excluded** from the Quick Plate layout. In the figure above (right), the user excluded row D and columns 11 and 12 from the layout. To de-select a row or column, just click the row letter or column number again.
- If rows or columns have been excluded, you have the option of apply fluid Layout to each region separately. The default is to skip over excluded rows and columns during layout.

Click **Next** to go onto the next step or **Cancel** to abort the in-process layout procedure and discard all modifications.

Fluid & titration page

- **Select the fluid(s)** to be used by clicking the checkbox next to the fluid name (below left). You can de-select an already-selected fluid by re-clicking the checkbox to clear it. Additionally, you can select **all fluids** by clicking the **Fluids** checkbox at the top of the list.



NOTE: You cannot select fluids with different unit types (molarity vs. mass/volume) or dispense modes (by concentration/volume) for the same Quick Plate Titration.

To set up Quick Plate Titration, you must enter information similar to that needed for Titration or Targeted Titration:

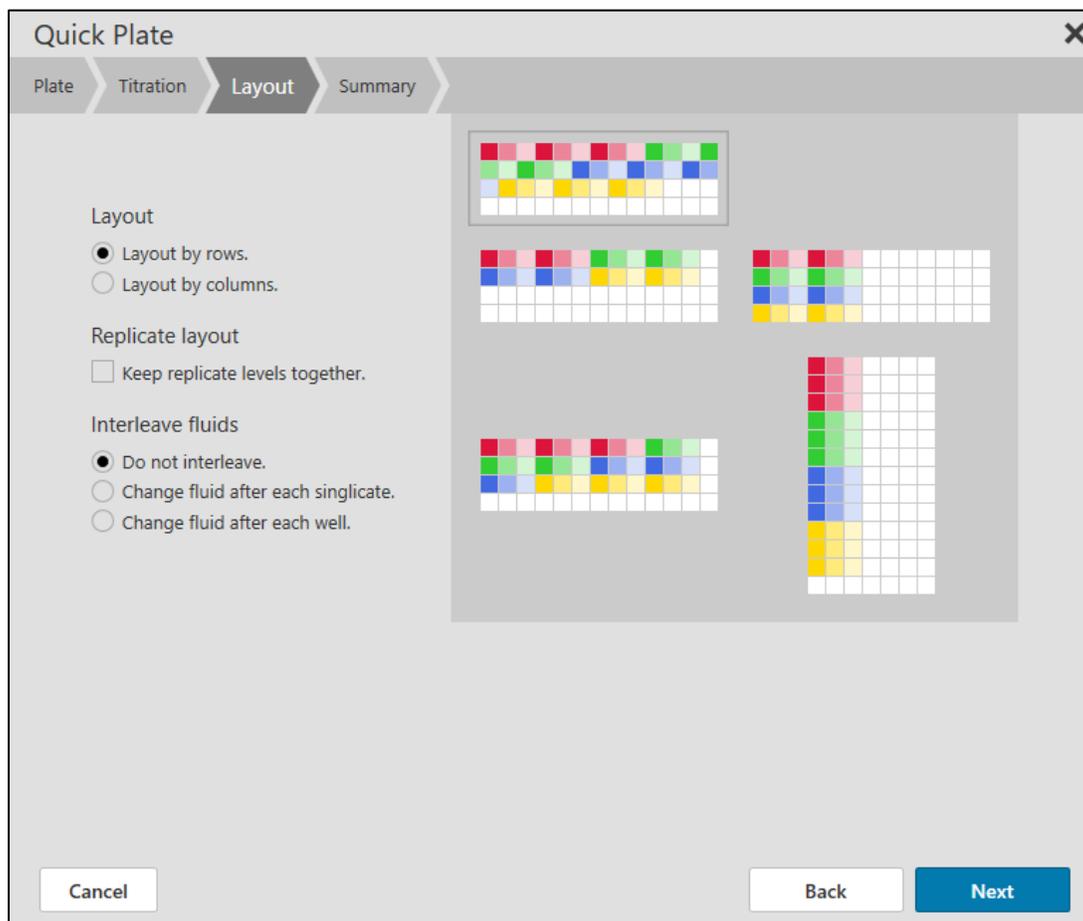
- Specify the number of titration levels desired for each fluid.
- Choose Titration or Targeted titration and then follow the instructions in the [Titration](#) or [Targeted titration](#) sections.

Once all parameters have been entered, click **Next** to go on to the next Quick Plate step. **Back** and **Cancel** are also available.



NOTE: A fluid shift or extension may be suggested. See the [Fluid shift or extend](#) section for more information.

Layout page



The Quick Plate Layout step allows you to decide how the titration series are laid out in the plate. You can arrange the titration points in many ways, including with the different fluids interleaved if desired. With your desired layout in mind, adjust the Quick Plate outcome by experimenting with various layout settings. As you change the primary layout settings, you can see the effects in the example layout options.



NOTE: Several of the setting options will change based on previous selections.

Layout

- **Layout by rows:** The Well plate grid shows a layout with each fluid's replicate levels progressing across a row.
- **Layout by columns:** The Well plate grid shows a layout with each fluid's replicate levels progressing down a column.

Replicate layout

Choose to keep replicate levels together or not. Keeping replicate levels together will keep all the concentration replicates of each fluid together. Not keeping replicate levels together will instead keep the fluid's titration series, of decreasing concentration, together. The remaining choices will adjust to the selection here.

Interleave fluids

- **Do not interleave:** The wells for each fluid are maintained together, according to the other layout selections.
- **Change fluid after each level replicate** (option available when keep replicate levels together is selected): After all the replicated for a given fluid level are applied, the next fluid will then be placed into the wells, until all of the fluids are placed. Then, the next set of level replicates for the first fluid will be applied, etc.
- **Change fluid after each singlicate** (option available when keep replicate levels together is not selected): All the titration levels for a single fluid are applied to the wells, then the titration levels of the next fluid are applied, until all the non-replicated fluid titration levels are applied. Then the replicates for all the fluids are applied in the same manner.
- **Change fluid after each well:** If replicate levels are kept together, this option places the first titration level of each fluid into the wells, changing fluid for every well. Then the replicates for all the fluids are applied in the same manner, followed by the other titration levels, until all the levels and replicates have been applied.

However, if replicate levels are not being kept together, this option places the first titration level of each fluid into the wells, changing fluid for every well. Then it places the next titration level for each fluid in the same manner until all the singlicates for all the fluids have been placed. Then the replicates for all the fluids are applied in the same manner.

Example layout options

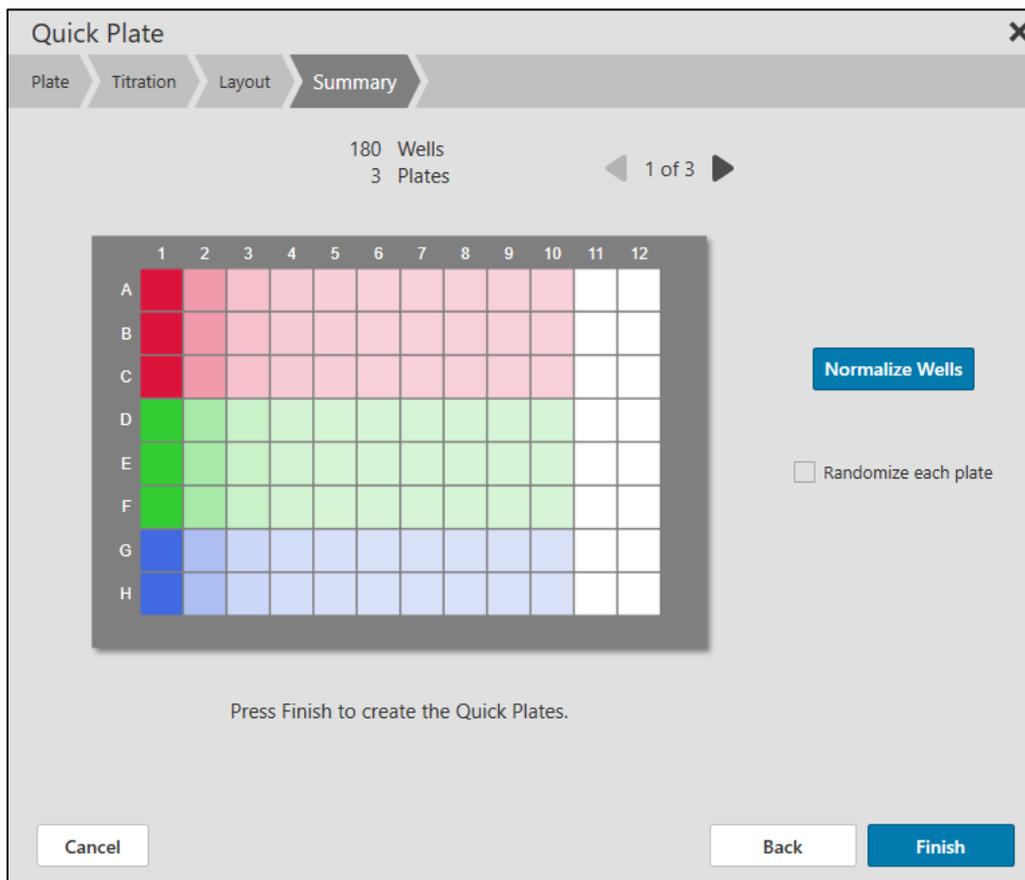
The final layout selection affects the flow of the layout. Example layouts are shown for each option. These options vary significantly based on previous selections. Move the mouse over an option to display a brief description of the layout behavior.

Once all parameters have been entered, click **Next** to go on to the next Quick Plate step. **Back** and **Cancel** are also available.

Summary page

The Summary gives you an illustration of how the new plate will look. It also indicates the number of wells and plates involved in the scheme.

The mouse can be moved over individual wells to display contents.



Click **Finish** to create the new plate(s). The new plate will then become the current plate. **Back** and **Cancel** are also available.



NOTE: If Aqueous fluids are in use for the Quick plate wizard and more than 8 plates are generated a warning message is displayed.



Aqueous Plate Limit

Due to evaporation effects, this protocol may be limited to 8 plates or 1536 wells.

Close

Quick plate examples

In this section, several examples will be provided for **Quick Plate** layouts that increase in complexity. These examples are intended to provide initial guidance for how to use the different layout options in **Quick Plate**. They are not intended to be exhaustive, and it is recommended that users experiment with the various layout options to achieve the desired plate layout.

Example 1: One 96-well plate with 8-point triplicates of four compounds

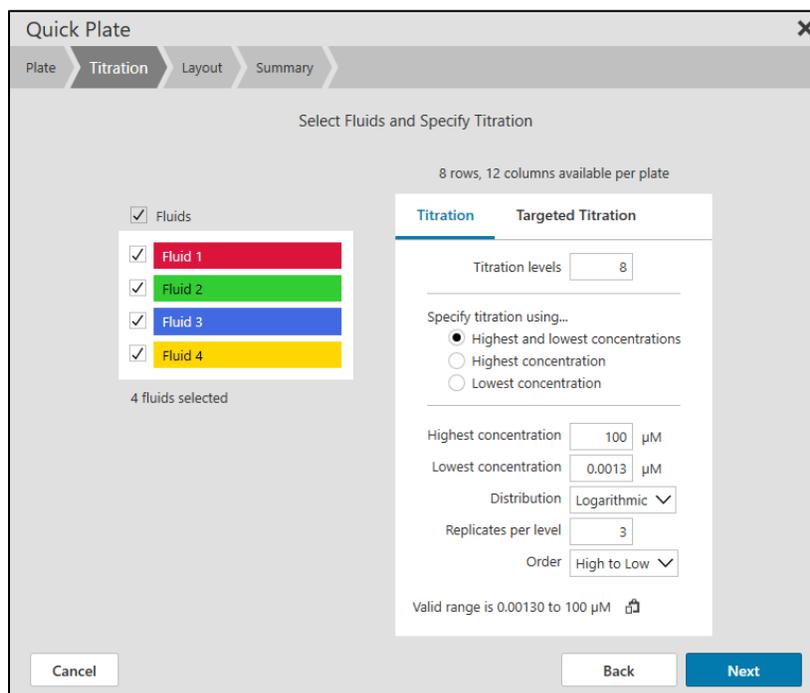
This example uses all the wells of the 96-well plate, with no columns or rows used for controls/left blank.

The titrations are laid out in columns, with the replicate series placed in adjacent columns.

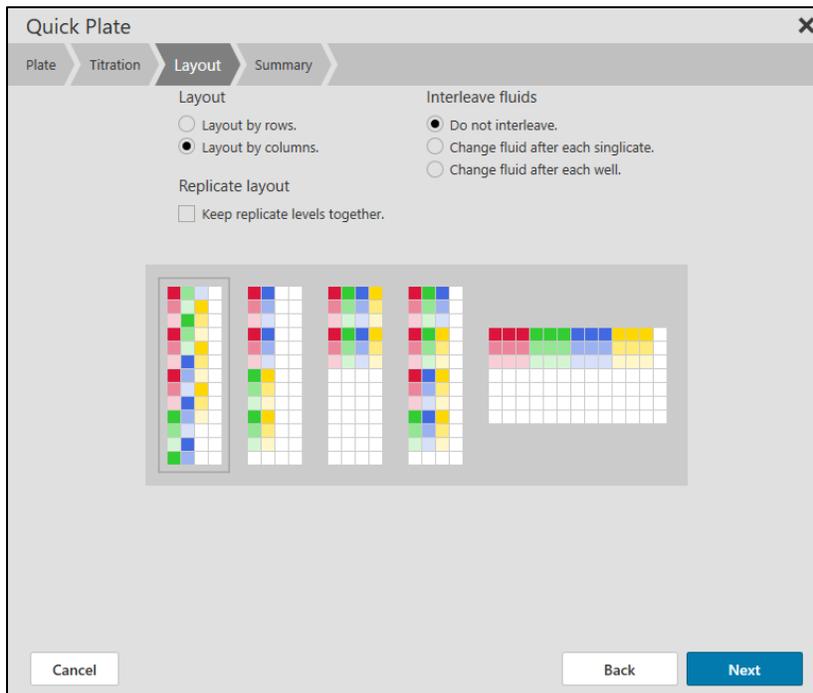
The desired resulting plate is shown, along with the titration and layout page selections made to generate the plate.



In the Fluids & Titration page, select the fluids to apply to the newly created quick plates and specify the Titration or Targeted titration details, including number of levels, number of replicates, and concentration range and distribution.



In the Layout page, specify how to arrange the desired wells in the newly created plate/s. The example provides visual guidance regarding the impact of the different selections, but it does not represent the actual plate that will result. The resulting plate/s are shown after **Next** is clicked, at which point you may still click **Back** within the **Quick Plate** setup.

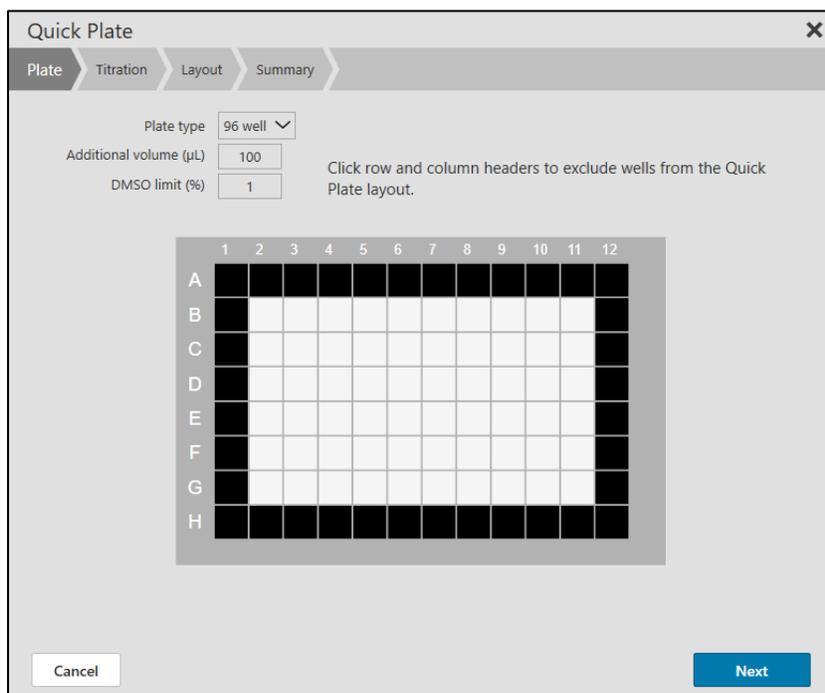


Example 2: Three 96-well plates with 10-point triplicates of six compounds, replicates in different plates

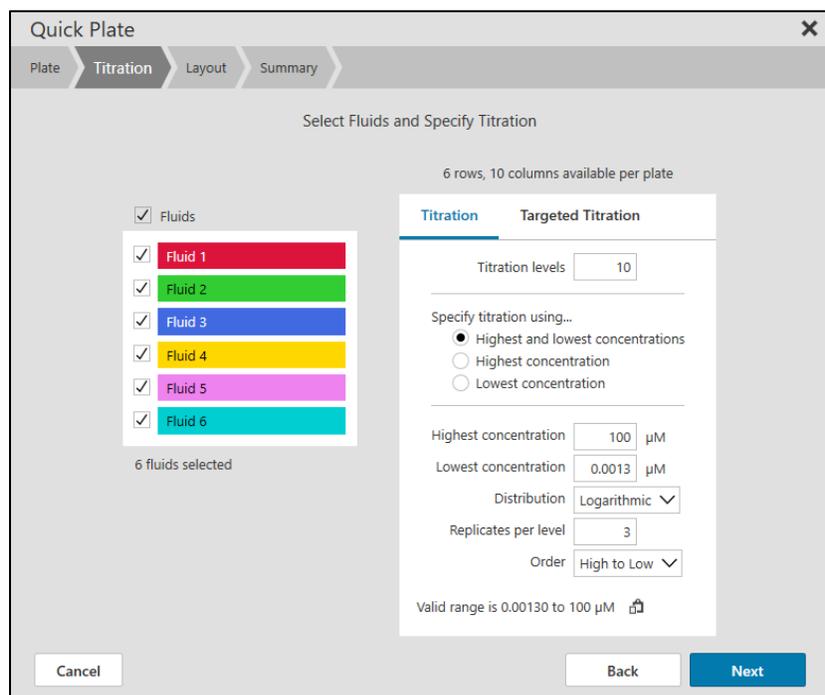
In the second example, more compounds are applied to multiple 96-well plates using the Quick Plate function. The six fluids are titrated as 10-point triplicates, with unused rows and columns at the plate edges. In this example, the replicates are applied to different plates, so the fluid singlicates are interleaved in the plate layout. Shown below are the desired plates and the Quick Plate pages used to generate them.



In the Plate page, select any rows and/or columns to exclude from the layout. In this case, the plate perimeter was excluded.



In the Fluids & titrations page, select the fluids to apply to the newly created quick plates and specify the Titration or Targeted titration details, including number of levels, number of replicates, and concentration range and distribution.

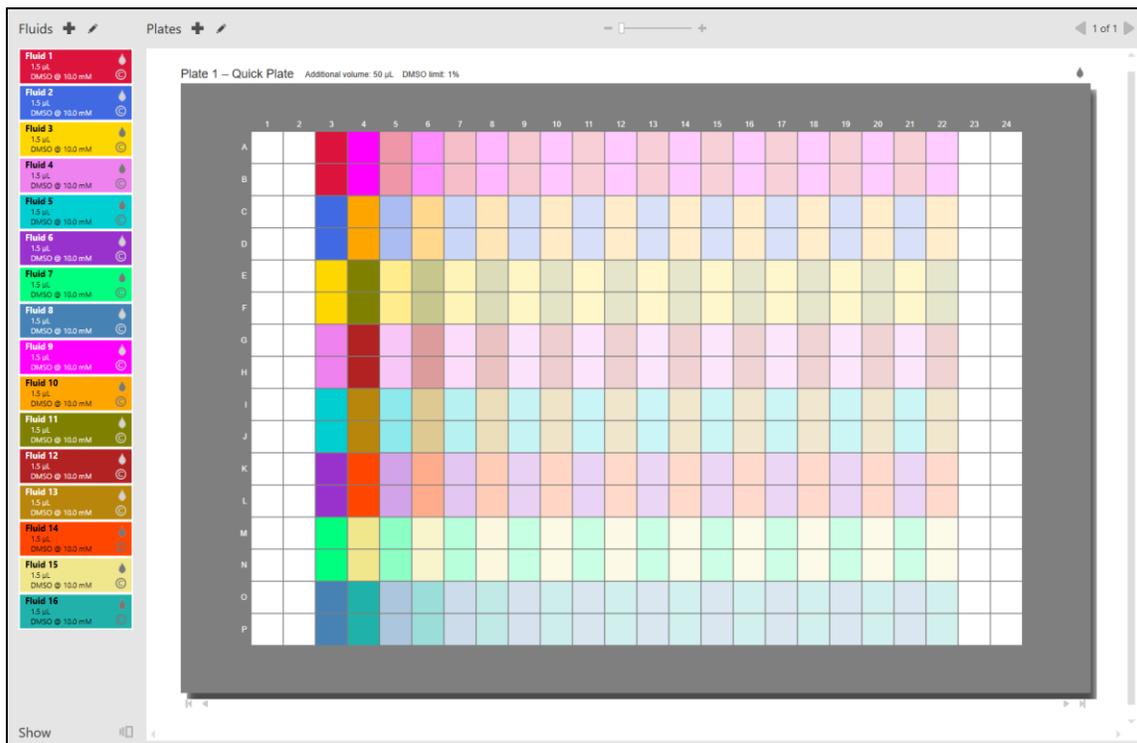


In the Layout page, specify how to arrange the desired wells in the newly created plate(s). For this example, to place the replicates into different plates, we select the option **interleave fluids – change fluid after each singlicate**.



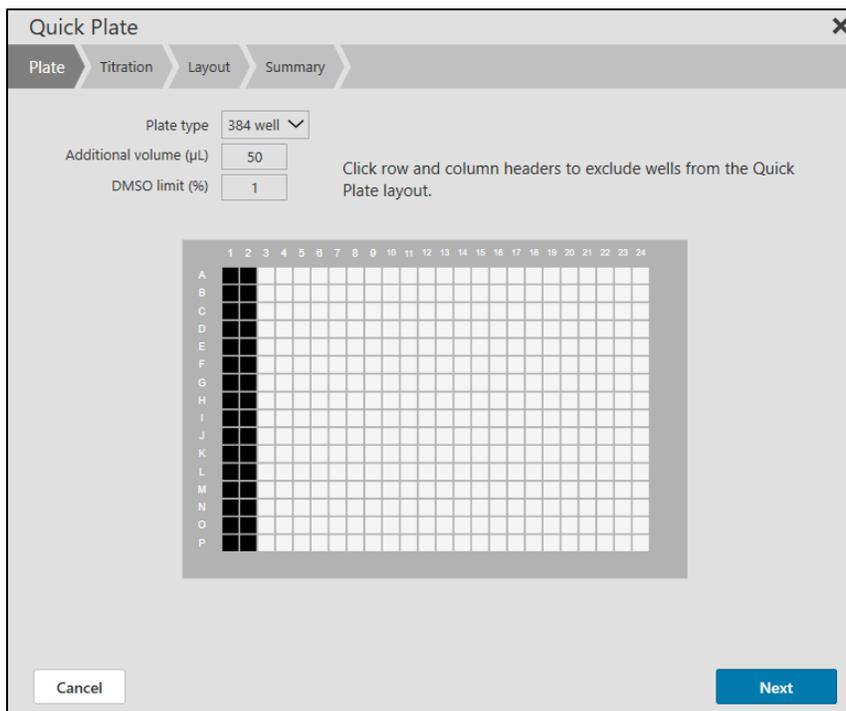
Example 3: One 384-well plate with 10-point duplicates of 16 compounds, interleaved

In the third example, a plate with multiple fluids that alternate, or are interleaved, on a well-by-well basis will be created. This plate layout might be applicable if you need to match existing plates that have been generated by transferring wells from a 96-well to a 384-well plate. The two outermost columns on each side of the plate

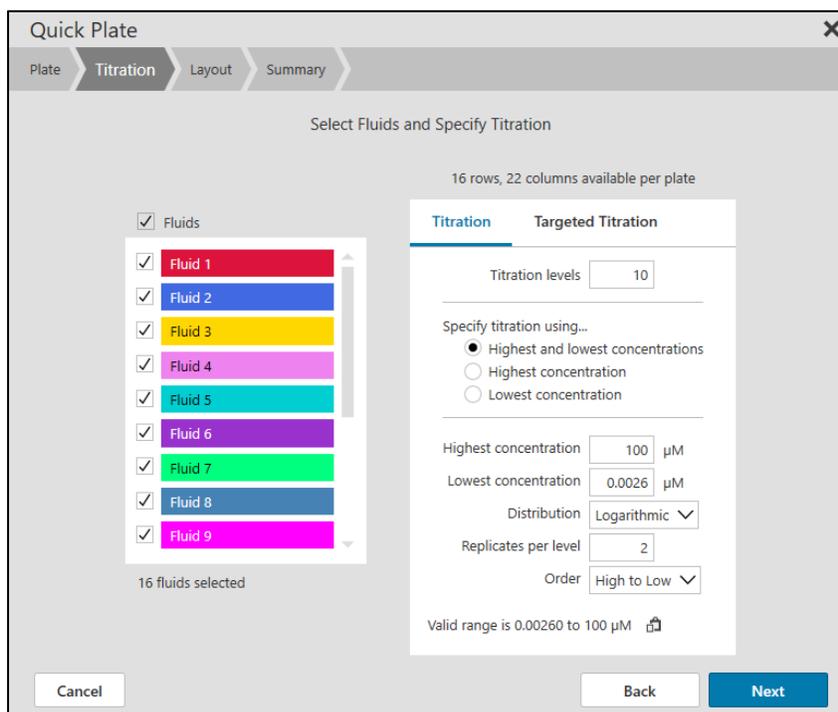


are excluded from this layout. The desired plate layout, using 16 fluids titrated in 10-point replicates, is shown, along with the titration and plate layout pages used to generate the plate.

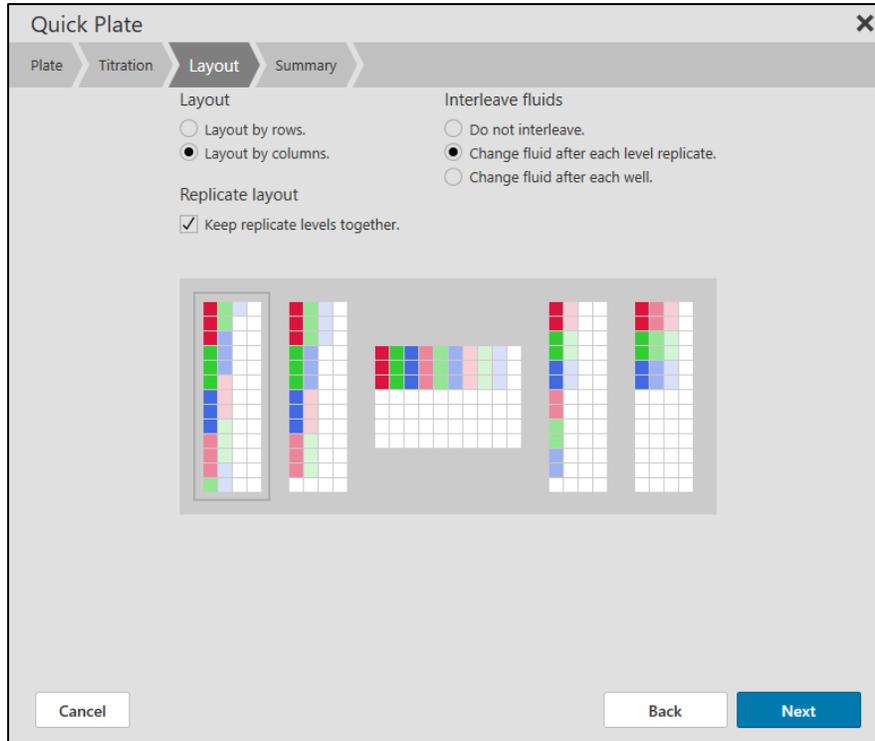
In the Plate page, select the plate type and any rows and/or columns to exclude from the layout. In this case, a 384-well plate was selected and two columns at the left edge excluded.



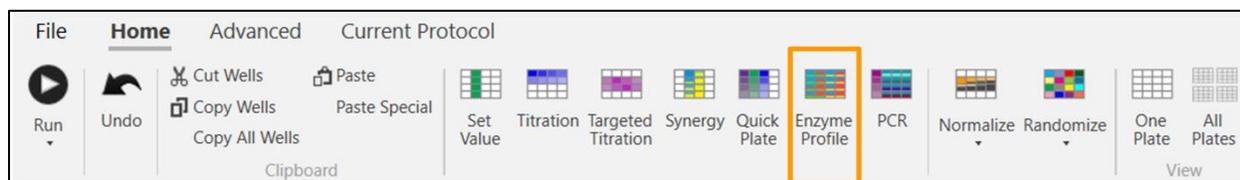
In the Fluids & titrations page, select the fluids to apply to the newly created quick plates and specify the Titration or Targeted titration details, including number of levels, number of replicates, and concentration range and distribution.



In the Layout page, specify how to arrange the desired wells in the newly created plate/s. For this example, to achieve the desired layout, we have to lay out the fluids by columns and keep replicate levels together. Then, to achieve the interleaved pattern, we change fluid after each level replicate and keep the levels together, starting each level on a new column.



Enzyme Profile



The **Enzyme Profile** function allows you to automate the process of enzymatic assay development, validation, and generation of Michaelis-Menten kinetic analyses and Lineweaver-Burk plots (for example).

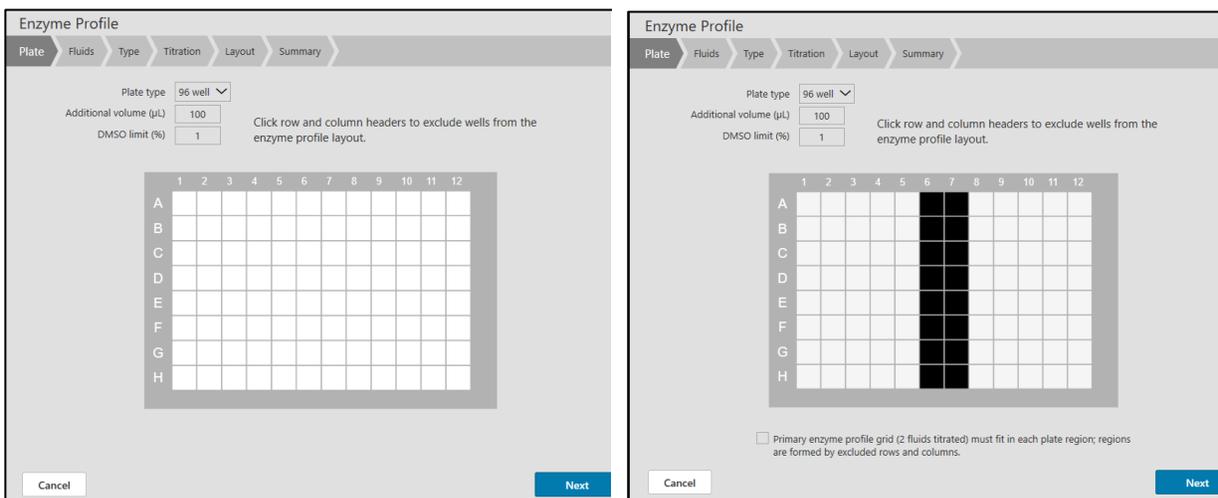
Once the enzymes, substrates/cofactors, and inhibitors are set up as fluids, you can design well layouts where enzymes, substrates/cofactors, or inhibitors are dispensed in constant (fixed) or variable (titrated) amounts. This makes it easy to determine the proper enzyme and substrate/cofactor concentrations, and inhibition mechanisms for enzymatic assays.

On the Home Tab, click **Enzyme Profile**. You then navigate and enter parameters in a series of dialog boxes to create the plates. Once you have completed the procedure, the new plates are added to the protocol.



NOTE: When using the Enzyme Profile function, it is not necessary to pre-select the wells to which you wish to apply fluids. The Enzyme Profile function automatically creates new plates with the programmed well content and ignores any wells that you may have already selected in an existing plate.

Plate page



- Assign the **Plate type**, **Additional volume**, and **DMSO limit** in the fields provided at the top of the dialog box – the current plate's parameters are the defaults.



NOTE: **DMSO limit** is only displayed when DMSO fluids are in the protocol.

- If desired, click **the row letter(s) and/or column number(s)** to select the rows and columns of wells that should be **excluded** from the Enzyme Profile layout. In the figure above (right), the user excluded columns 6 and 7 from the layout. To de-select a row or column, just click the row letter or column number again.

- If rows or columns have been excluded, you have the option of apply fluid Layout to each region separately. The default is to skip over excluded rows and columns during layout.

Click **Next** to go onto the next step or **Cancel** to abort the in-process layout procedure.

Fluids page

The image shows two screenshots of the 'Enzyme Profile' dialog box, which is used for selecting fluids for an enzyme profile. The dialog has a title bar with a close button (X) and a breadcrumb trail: Plate > Fluids > Type > Titration > Layout > Summary.

The first screenshot shows the 'Fluids' step. The instructions are: 'Select up to 2 Enzymes. Select up to 2 Substrates/Cofactors. Select Inhibitors.' There are three columns of fluid selection options: 'Enzymes', 'Substrates/Cofactors', and 'Inhibitors'. Each column contains a list of fluid types with checkboxes: Enzyme 1, Co-factor 1, Substrate A, Inhibitor 1, Inhibitor 2, and Inhibitor 3. In this screenshot, all checkboxes are unchecked. At the bottom, there are 'Cancel', 'Back', and 'Next' buttons.

The second screenshot shows the same dialog after selections have been made. Under 'Enzymes', 'Enzyme 1' is selected (checkbox checked). Under 'Substrates/Cofactors', 'Co-factor 1' and 'Substrate A' are selected. Under 'Inhibitors', 'Inhibitor 1', 'Inhibitor 2', and 'Inhibitor 3' are selected. The counts below each column are: '1 Enzyme', '2 Substrates/Cofactors', and '3 Inhibitors'. The 'Next' button is now highlighted in blue.

Select fluids to be used as enzymes, substrates/cofactors, and inhibitors.

- You must select one or two enzymes.
- You may select one or two substrates/cofactors, but you do not need to select any.
- You may select one or more inhibitors, but you do not need to select any. All inhibitors will be titrated using the same settings.

Click **Next** to go onto the next step or **Cancel** to abort the in-process layout procedure.



NOTE: You cannot select fluids with different unit types (molarity vs. mass/volume) or dispense modes (by concentration/volume) in a list.

Type page

Enzyme Profile

Plate Fluids **Type** Titration Layout Summary

Select 1 or 2 fluids to titrate.
Remaining fluids will have fixed values.

Enzyme: **Enzyme 1**
 Titrated

Substrates/Cofactors: **Co-factor 1** **Substrate A**
 Titrated Titrated
Down rows ▾

3 Inhibitors: **Inhibitor 1** **Inhibitor 2** **Inhibitor 3**
 Titrated (all Inhibitors)
Across columns ▾

Cancel Back Next

- Select one or two fluids for titration. Fluids not selected for titration will be dispensed at a fixed volume or concentration.
- If two fluids are selected for titration, you can also choose which fluid will be titrated down rows and which will be titrated across columns.
- Click **Next** to go onto the next step or **Cancel** to abort the in-process layout procedure.

Titration page

For each titrated fluid, choose Titration, or Targeted Titration.

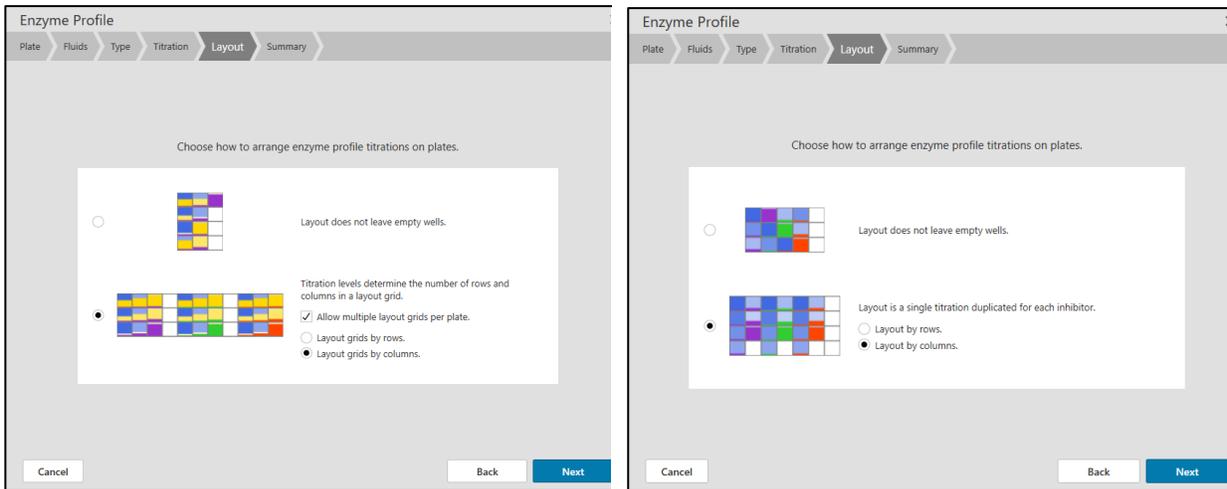
- Specify the number of Titration levels via manual entry.
- Fill out all other fields according to instructions in the [Titration](#) or [Targeted titration](#) sections.
- You can Include “0” value by clicking the check box.
- Distribution is pre-selected as Linear or Logarithmic depending on the type of fluid, but you can change the selection.

If there are two titration fluids, they will form a titration grid, flowing down the non-excluded rows and across the non-excluded columns. The number of available rows and columns is displayed for reference.

- For each non-titrated fluids, specify a fixed volume or concentration.

Click **Next** to go onto the next step or **Cancel** to abort the in-process layout procedure.

Layout page



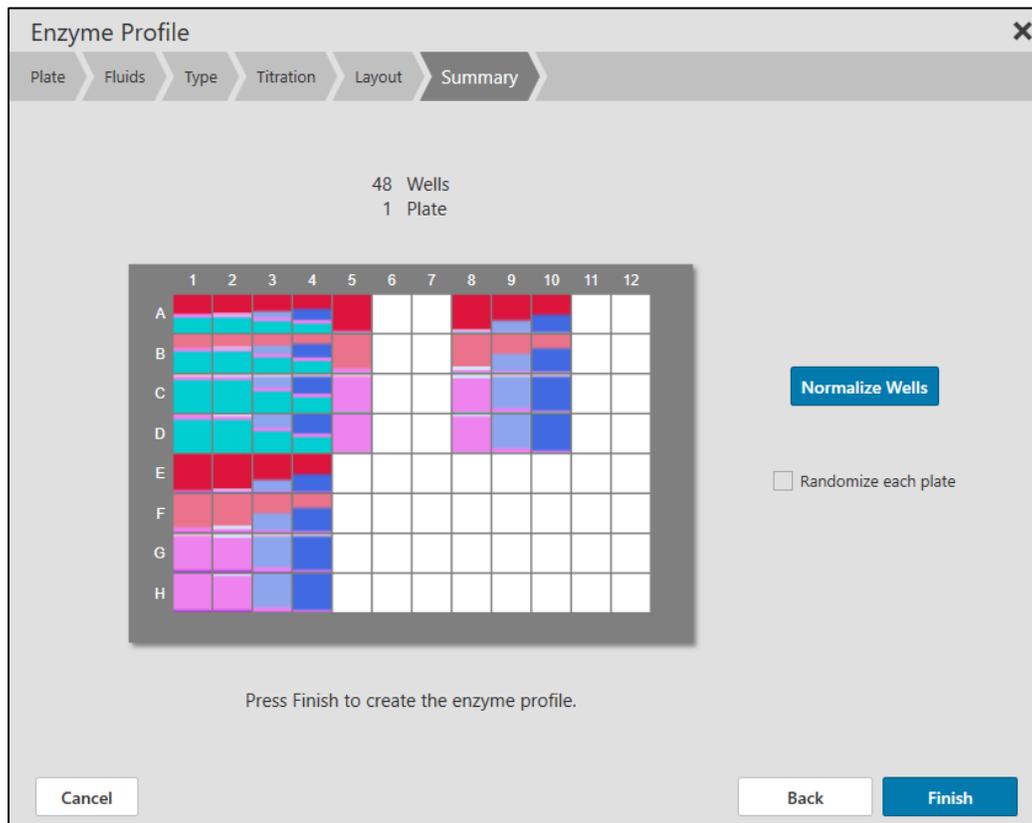
Layout options will vary depending on the number of titrated and fixed fluids, plate settings, etc.

Select a primary layout option and then detailed options for that layout. In the example above, checking or clearing the “Allow multiple layout grids per plate” checkbox will change the example graphics and detailed options available.

Summary page

The Summary gives you an illustration of how the new plate will look. It also indicates the number of wells and plates involved in the scheme.

The mouse can be moved over individual wells to display contents.



Click **Finish** to create the new plate(s). The new plate will then become the current plate. **Back** and **Cancel** are also available.



NOTE: If Aqueous fluids are in use for the Enzyme profile wizard and more than 8 plates are generated a warning message is displayed.

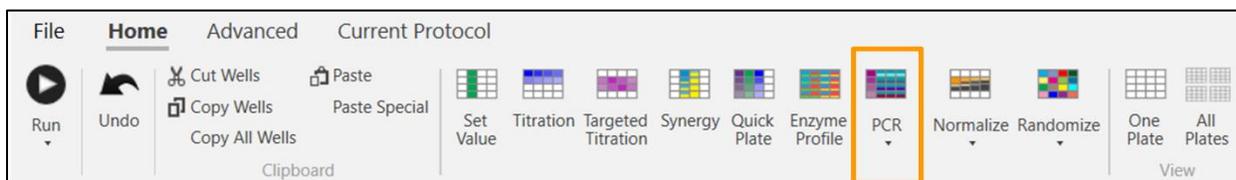


Aqueous Plate Limit

Due to evaporation effects, this protocol may be limited to 8 plates or 1536 wells.

Close

PCR



The PCR function allows you to automate the process of designing well plates for PCR experiments. This involves setting up dispense of master mixes, primers (also known as targets or probes) and samples (also known as templates) using the dispenser. Master mixes that are added using another dispense technology should be specified as “Additional Volume” on the Plate Page below.

Before using the PCR function, all fluids must be defined. If master mixes are premixed with primers, enter each mixture as a single fluid. Master mixes must be set to dispense by volume.

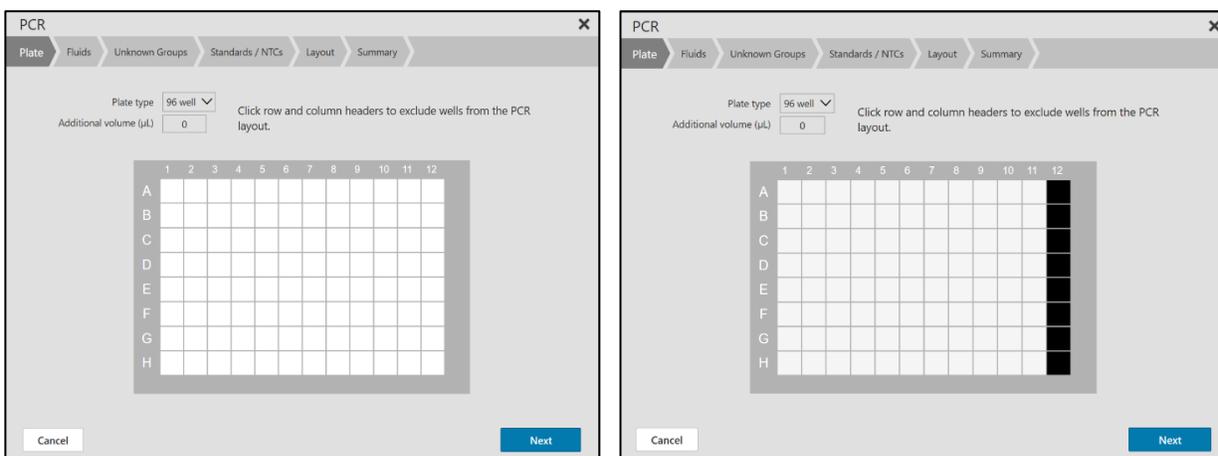
After fluids to be dispensed are defined, protocols can be designed where all wells have the same master mix volumes. All master mixes will be dispensed in every well of the layout. For each primer/sample combination, the concentrations or volumes as well as the number of replicates can be specified. Optionally, standard curves and an NTC (no template control) region can be included; wells can be normalized to have the same total volume.

On the Home Tab, click **PCR**. You then navigate and enter parameters in a series of dialog boxes to create the plates. Once you have completed the procedure, the new plates are added to the protocol.



NOTE: When using the PCR function, it is not necessary to pre-select the wells to which you wish to apply fluids. The PCR function automatically creates new plates with the programmed well content and ignores any wells that you may have already selected in an existing plate.

Plate page



- Assign the **Plate type**, **Additional volume**, and **DMSO limit** in the fields provided at the top of the dialog box. The current plate's type and DMSO limit will be the default choices; however, **additional volume** will always default to 0 (zero) for PCR, meaning that all well volume is added by the dispenser. As noted above, if master mix is to be dispensed by another method, specify that master mix volume in the additional volume field.



NOTE: DMSO limit is only displayed when DMSO fluids are in the protocol.

- If desired, click the **row letter(s) and/or column number(s)** to select the rows and columns of wells that should be **excluded** from the PCR layout. In the figure above (right), the user excluded column 12 from the layout. To de-select a row or column, just click the row letter or column number again.

Click **Next** to go onto the next step or **Cancel** to abort the in-process PCR layout procedure.

Fluids page

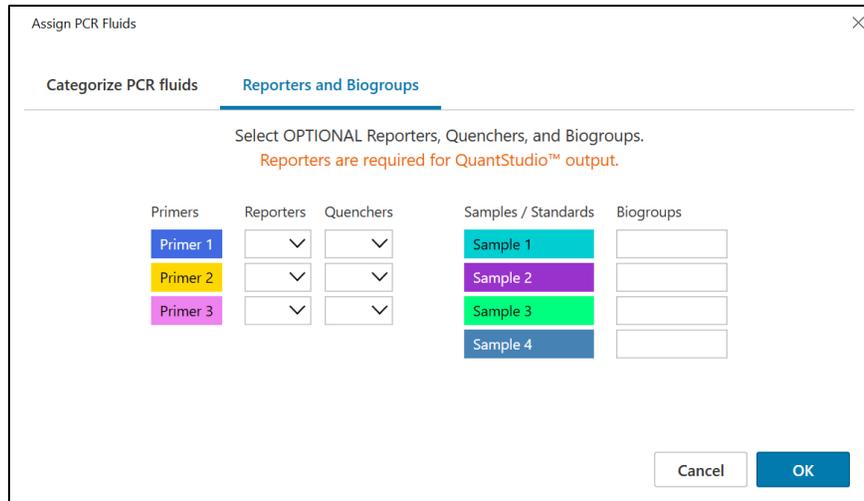
Select **Master Mix**, **Primer**, and **Sample** fluids. Only fluids dispensed by volume are included in the Master Mix selection list.

- You must select at least one primer.
- You must select at least one sample.
- You do not need to select any master mixes if your primers include master mix or if master mixes will be dispensed by another method.
- You do not need to categorize all fluids.



TIP: If master mix is not to be dispensed by the dispenser, do not select any master mixes on the Fluids page, but include their volume as Additional Volume in the Plate tab (see [Plate-associated information](#)).

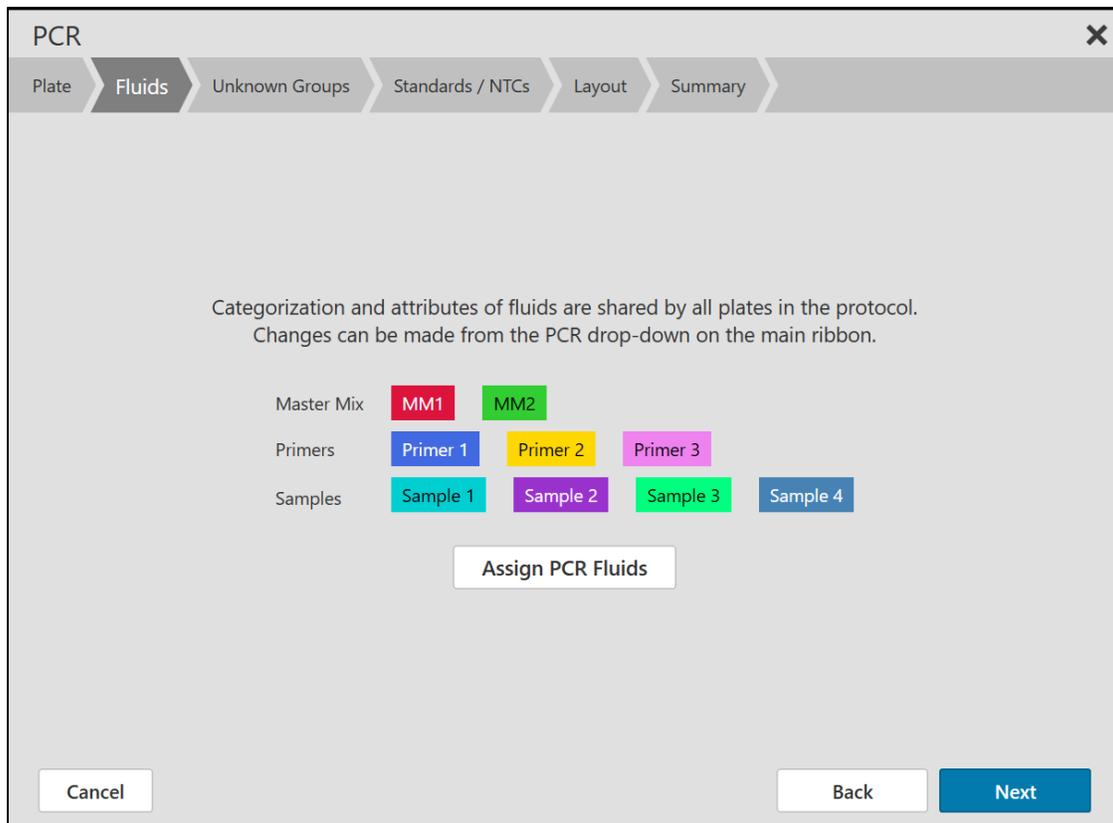
To add more information to your experimental groups including information about reporters, quenchers, and Biogroups. Reporter information is required when loading your dispense information report into Quantstudio™ as described in [Appendix A-Reports](#) under **QuantStudio Compatible format (*.DA.csv)**



The 'Assign PCR Fluids' dialog box is shown with the 'Reporters and Biogroups' tab selected. It contains a table for assigning optional reporters, quenchers, and biogroups to primers and samples. A note states 'Reporters are required for QuantStudio™ output.' The table has columns for Primers, Reporters, Quenchers, Samples / Standards, and Biogroups. There are three primers and four samples listed.

Primers	Reporters	Quenchers	Samples / Standards	Biogroups
Primer 1	▼	▼	Sample 1	
Primer 2	▼	▼	Sample 2	
Primer 3	▼	▼	Sample 3	
			Sample 4	

If protocol fluids have previously been assigned PCR tasks, a summary is displayed. Press **Assign PCR Fluids** to add or modify PCR fluid tasks.



The 'PCR' summary screen shows a breadcrumb trail: Plate > Fluids > Unknown Groups > Standards / NTCs > Layout > Summary. A message states: 'Categorization and attributes of fluids are shared by all plates in the protocol. Changes can be made from the PCR drop-down on the main ribbon.' Below this, a summary of assigned fluids is shown: Master Mix (MM1, MM2), Primers (Primer 1, Primer 2, Primer 3), and Samples (Sample 1, Sample 2, Sample 3, Sample 4). An 'Assign PCR Fluids' button is centered below the summary. At the bottom are 'Cancel', 'Back', and 'Next' buttons.

Click **Next** to go onto the next step or **Cancel** to abort the in-process PCR layout procedure.

Unknown Groups page

PCR

Plate Fluids Unknown Groups Standards / NTCs Layout Summary

Specify Master Mix and one or more Unknown groups. All Primers will be combined with each Sample in a group.

2 Master Mixes

MM1 nL
0.0130 nL minimum

MM2 nL
0.0130 nL minimum

Unknown Group 1

Replicates per sample

Primers

Primer 1

Primer 2

Primer 3

Samples

Sample 1

Sample 2

Sample 3

Sample 4

+

Cancel Back Next

This dispense volume for each **Master Mix** is specified separately and will be dispensed in all Unknown, Standard, and NTC wells.

Multiple PCR Unknown groups can be specified. Use the plus **+** button on the right edge to add additional PCR groups.

PCR

Plate Fluids Unknown Groups Standards / NTCs Layout Summary

Specify Master Mix and one or more Unknown groups. All Primers will be combined with each Sample in a group.

2 Master Mixes

MM1 nL
0.0130 nL minimum

MM2 nL
0.0130 nL minimum

Unknown Group 1

Replicates per sample

Primers

Primer 1

Primer 2

Primer 3

2 Primers

μM
0.000433 to 33.3 μM suggested

Samples

Sample 1

Sample 2

Sample 3

Sample 4

4 Samples

ng/mL
0.433 to 33.333 ng/mL suggested

Unknown Group 2

Replicates per sample

Primers

Primer 1

Primer 2

Primer 3

1 Primer

μM
0.000433 to 33.3 μM suggested

Samples

Sample 1

Sample 2

Sample 3

Sample 4

2 Samples

ng/mL
0.433 to 33.333 ng/mL suggested

+

Cancel Back Next

For each PCR Unknown group:

- Specify the number of **Replicates** that will be created for each **Primer/Sample** combination.
- All **Primers** in the group share a single dispense value.

- All **Samples** in the group share a single dispense value.
- Each **Sample** will be dispensed separately with all **Primers** in the group.

In this example, there are two Unknown groups. The first group uses 4 samples with 8 replicates, for a total of 32 wells with two multiplex primers in each well. The second group uses 2 samples with 6 replicates, for a total of 12 wells.

Click **Next** to go onto the next step or **Cancel** to abort the in-process PCR layout procedure.



NOTE: You cannot select fluids with different unit types (molarity vs. mass/volume) or dispense modes (by concentration/volume) in a list (column).

Standards / NTCs page

Optional Standards and NTC (No Template Controls) can be specified.

Specify optional standards to be dispensed on one or all PCR plates.

- If **Standards** is not checked, no standards will be planned.
- Select one or more **Primers** and specify the fixed dispense volume or concentration for the standard.
- Select one or more **Samples (standards)** and specify the titration to use for each standard.
- The replicates setting in the titration panel will determine how many replicate standards are created for each primer/sample combination
- Each primer will be combined separately with each sample. This is different from how the Unknown groups are planned where primers can be multiplexed.

Specify optional NTC (No Template Control) region.

- If **NTCs** is not checked, no NTCs will be planned.
- Specify the **Replicates** to be used for each primer.
- Select one or more **Primers** and specify the fixed dispense volume or concentration.
- Each primer is dispensed separately.

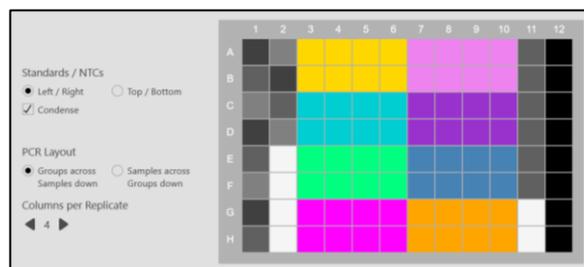
If **Standards/NTCs on each plate** is not checked, standards and NTCs will only be dispensed on the first PCR plate.

Click **Next** to go onto the next step or **Cancel** to abort the in-process PCR layout procedure.



NOTE: You cannot select fluids with different unit types (molarity vs. mass/volume) or dispense modes (by concentration/volume) in a list (column).

Layout page



If Standards or NTCs were specified, there are additional layout options.

- Standards/NTCs can be located down the left and right edges or along the top and bottom edges of the plate.
- Standards can be condensed so the next replicate will occur immediately following the previous replicate even if that results in curves being broken across columns or rows. The NTC region is always condensed.

Changing the columns/combination setting will cause the PCR regions to change shape in the layout.

Since all replicated combinations will be planned on the same plate, multiple plates may be required for all of the primer/sample combinations.

You can also create a layout that **does not leave any empty wells** by selecting the checkbox at the top of the page.

Click **Next** to go onto the next step or **Cancel** to abort the in-process PCR layout procedure.

Summary page

The Summary gives you an illustration of how the new plate will look. It also indicates the number of wells and plates involved in the scheme.

The mouse can be moved over individual wells to display contents. Unknown groups, Standard, and NTC areas are indicated below.

It is optional to [Normalize](#) PCR wells. It may be desirable to normalize to a constant total assay volume if samples are different concentrations. This would typically be accomplished by normalizing with water and surfactant using an aqueous fluid class.

Click **Finish** to create the new plate(s). The new plate will then become the current plate. **Back** and **Cancel** are also available.

Well tooltips show PCR information for all PCR fluids (Master Mix, Primer, and Sample) and wells (Unknown, Standard, and NTC).

Plate 1 Well B7
Unknown Group 2
 Aqueous Volume: 3,630 nL
 Additional Volume: 0 µL
 Total Volume: 3.6 µL

- **MM1** Master Mix
Volume: 2,500 nL
- **MM 2** Master Mix
Volume: 1,000 nL
- **Primer 3** Primer
Concentration: 1.50 µM
Volume: 54.4 nL
- **Sample 2** Sample
Concentration: 2,500 ng/mL
Volume: 75.6 nL

Plate 1 Well B2
Standard
 Aqueous Volume: 3,596 nL
 Additional Volume: 0 µL
 Total Volume: 3.6 µL

- **MM1** Master Mix
Volume: 2,500 nL
- **MM 2** Master Mix
Volume: 1,000 nL
- **Primer 1** Primer
Concentration: 1.00 µM
Volume: 35.9 nL
- **Sample 2** Sample
Concentration: 2,000 ng/mL
Volume: 59.9 nL

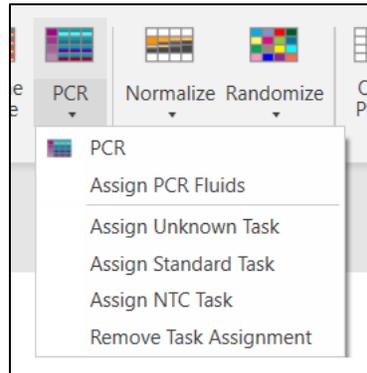
Plate 1 Well A11
NTC
 Aqueous Volume: 3,535 nL
 Additional Volume: 0 µL
 Total Volume: 3.5 µL

- **MM1** Master Mix
Volume: 2,500 nL
- **MM 2** Master Mix
Volume: 1,000 nL
- **Primer 1** Primer
Concentration: 1.00 µM
Volume: 35.4 nL

Manual PCR specification

It is possible to specify PCR information outside of the automated process. It may be that you need to create Standards with specific values that are simple to create manually, or to use multiplexed primers for Standards or NTCs. The automated PCR process could continue to be used to create the Unknown groups.

PCR Fluid and Task assignments can be made from the PCR ribbon drop-down menu. Task assignments can also be made from the right-click well menu.



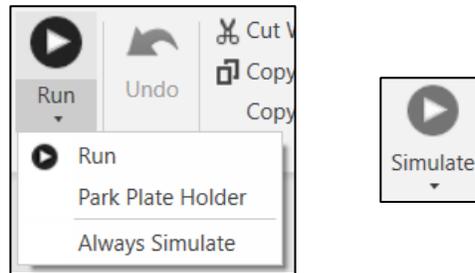
To assign a group of wells to a PCR task, select the wells and then select the task from the PCR menu. You can also remove any existing task assignments.

Running a fluid dispensing protocol

The dispensing software provides the environment for you to set up a protocol and run that protocol, either for dispensing fluids or isolating single cells. The runs can be:

- an **actual** run with your system connected to a dispenser and requiring physical loading of plates and cassettes, or
- a **simulated** run that gives you a preview of how a protocol will play out, with all the features of an actual run.

Run menu



Park Plate Holder will retract the plate holder.

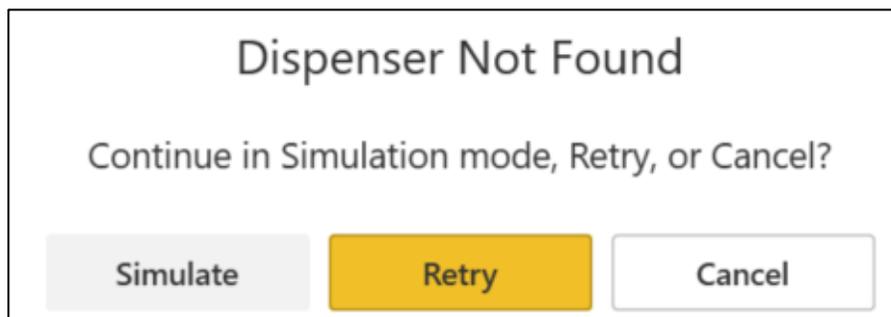
Always Simulate will simulate protocol runs, even if a device is connected. When Always Simulate is active, the Run button changes to Simulate.

Initial display and dialog

Load your protocol into the Main Operating Window. Then click the **Run** button on the Home Tab.



The dispensing software checks for a connection to a dispenser. If the software fails to detect a device, a dialog box is displayed, asking if you want to **Simulate** a run, **Retry** the connection for an actual run, or **Cancel** the run altogether.



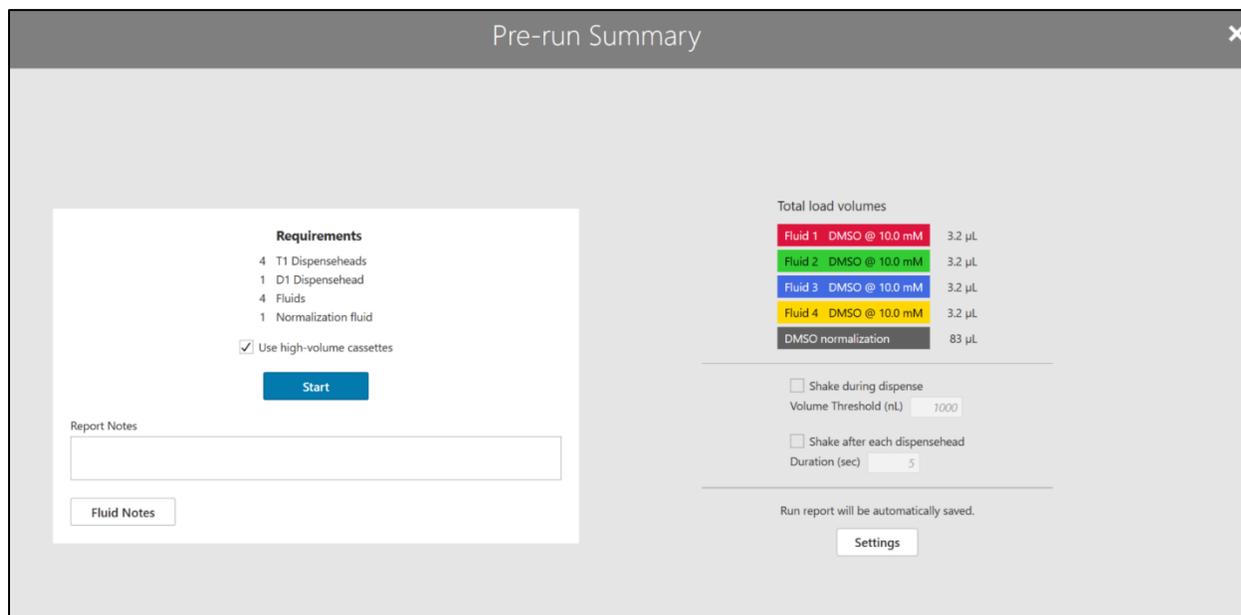


NOTE: To minimize evaporation, a protocol run dispensing aqueous fluid classes will be limited to 1536 wells or four plates.



NOTE: To ensure optimum performance, it is recommended that all plates include a well with volume of 30nL or greater for each fluid. Refer to [Appendix F—Best practices for dispensing](#) for more information.

The initial **Run** Window is then displayed.



The left area indicates the number of fluids and dispense heads that will be needed for the run. There is also a **Start** button to initiate the dispensing preparation and the subsequent run. Note that if the plate has normalized wells, the protocol calls for **normalization fluids**.

If high-volume dispense heads are appropriate for this protocol, the number of dispense heads will be listed for each cassette type. You can override the high-volume recommendation with the Use high-volume cassettes checkbox if you do not want to use high-volume cassettes. The default state of the checkbox is set in [Run Settings](#).

If text is entered into the Report Notes area, those notes will be written to the report file created after the run. Notes can also be added for each fluid, by clicking the **Fluid Notes** button.

The total load volume for each fluid is listed in the right portion of the screen along with a display of the **current shake parameters for wells and plates**. Changing these parameters changes the protocol. From this area it is also possible to enter [Settings](#) and modify report options.



NOTE: The protocol shake parameters are indicated, but individual plates may have Don't Shake selected.

Click **Start** to move onto the next phase of the run.

Running a protocol on the dispenser

Refer to the dispenser [Operating Manual](#) for detailed information about running a protocol on the dispenser.

Run complete

The screenshot shows a 'Run completed' dialog box with a green header bar. The main content is divided into two sections: 'General Dispense Summary' and 'Dispensed Fluids'. The 'General Dispense Summary' section contains the following text: 'Protocol saved to file', 'Run by user AUTH\blakem', 'Local timezone is (UTC-08:00) Pacific Time (US & Canada)', 'Simulation started at 11/11/2022 5:32:36 PM', 'Simulation Completed at 11/11/2022 5:33:01 PM', and 'Total runtime was 24 seconds'. The 'Dispensed Fluids' section lists five items: 'Fluid 1 (DMSO @ 10.0 mM)' with 3.2 µL loaded and 2.5 µL dispensed; 'Fluid 2 (DMSO @ 10.0 mM)' with 3.2 µL loaded and 2.5 µL dispensed; 'Fluid 3 (DMSO @ 10.0 mM)' with 3.2 µL loaded and 2.5 µL dispensed; 'Fluid 4 (DMSO @ 10.0 mM)' with 3.2 µL loaded and 2.5 µL dispensed; and 'DMSO normalization' with 83 µL loaded and 55 µL dispensed. Below the 'Dispensed Fluids' section, there is a message 'Report saved to (new) (7).xml' and a blue button labeled 'Open Report File Location'. At the bottom of the dialog, there are two buttons: 'Park Plate Holder and Close' and 'Close'.

Fluid Name	Loaded (µL)	Dispensed (µL)
Fluid 1 (DMSO @ 10.0 mM)	3.2	2.5
Fluid 2 (DMSO @ 10.0 mM)	3.2	2.5
Fluid 3 (DMSO @ 10.0 mM)	3.2	2.5
Fluid 4 (DMSO @ 10.0 mM)	3.2	2.5
DMSO normalization	83	55

If report settings are set to generate a report on every run, a message appears to let you know that the report is being generated. For information on what to look for in your report, see [Appendix A-Reports](#).

If a report is not automatically saved, it can later be saved using Save Report from the [Run menu](#).

Click **Close** to return to the Operating Window. Click **Park plate holder and Close** to retract the plate holder before closing. You can always retract the plate holder from the [Run menu](#).

Running a single cell dispensing protocol

Single cell dispensing and isolation into well plates up to 384 wells can be accomplished with the HP D100 Digital Dispenser. The D300e Digital Dispenser cannot be used for this purpose at this time due to electronic sensing limitations. On the D100, single cell cassettes are required for this workflow, please see [Appendix E—Dispense specifications](#) for more information about these cassettes.



NOTE: To ensure optimum performance, it is recommended that specific procedures be followed for cell stock generation and single cell protocols. Refer to [Appendix F—Best practices for dispensing](#) for more information.

Cell dispense fluids

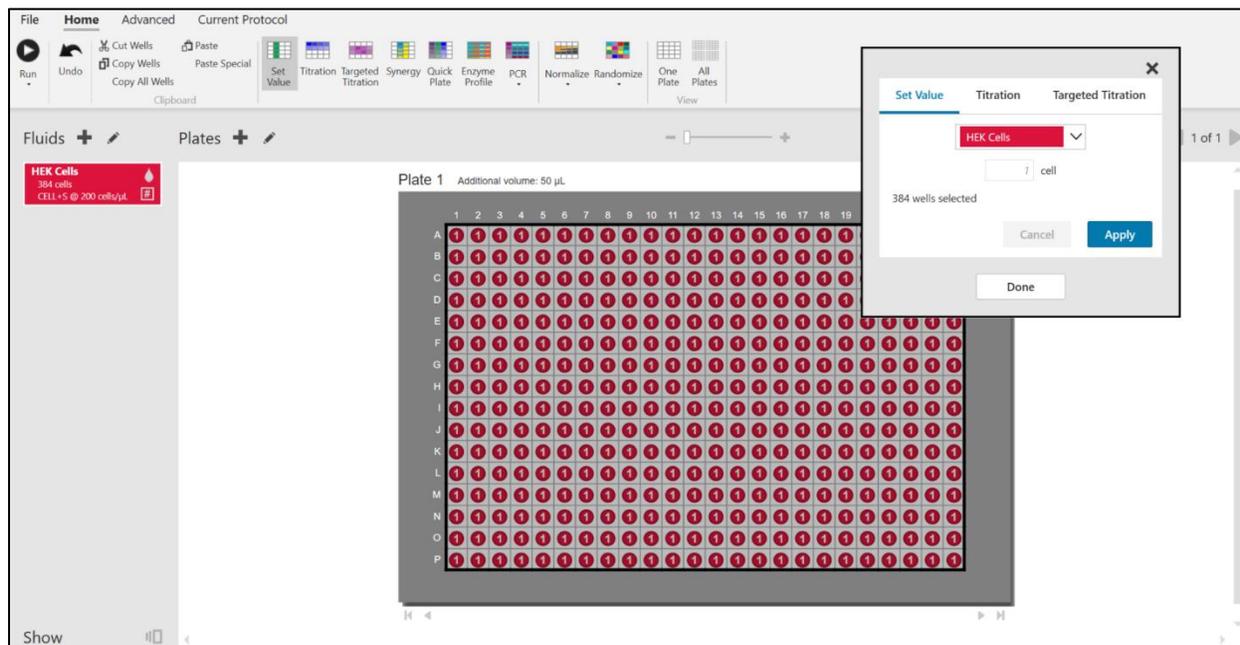
Add or modify a cell fluid group and class. Select **Cells** or **Sticky Cells** in the **Fluid Group** depending on your cells. Sticky cells require more clearing volume to dispense through the microfluidics. Use the Fluid class that corresponds to your average cell diameter: Small (10-14 μ m), Medium (15-17 μ m), Large (18-20 μ m) or Extra-Large (21-25 μ m). A concentration of 200 cells/ μ L is recommended for optimum performance.

The screenshot shows a 'Fluid' configuration window with the following details:

- Fluid List:** HEK Cells, 0 cells, CELL+S 200 cells/ μ L. Buttons: Remove Selected, Create Diluted Fluids.
- Configuration Fields:**
 - Name: HEK Cells
 - Fluid group: Cells
 - Fluid class: Small (10-14 μ m)
 - Concentration: 200 cells/ μ L
 - Dispense by: # (selected), Count
 - Source plate: (empty)
 - Source well: (empty)
 - Color: 1
 - Dispense: Yes
- Buttons:** Close

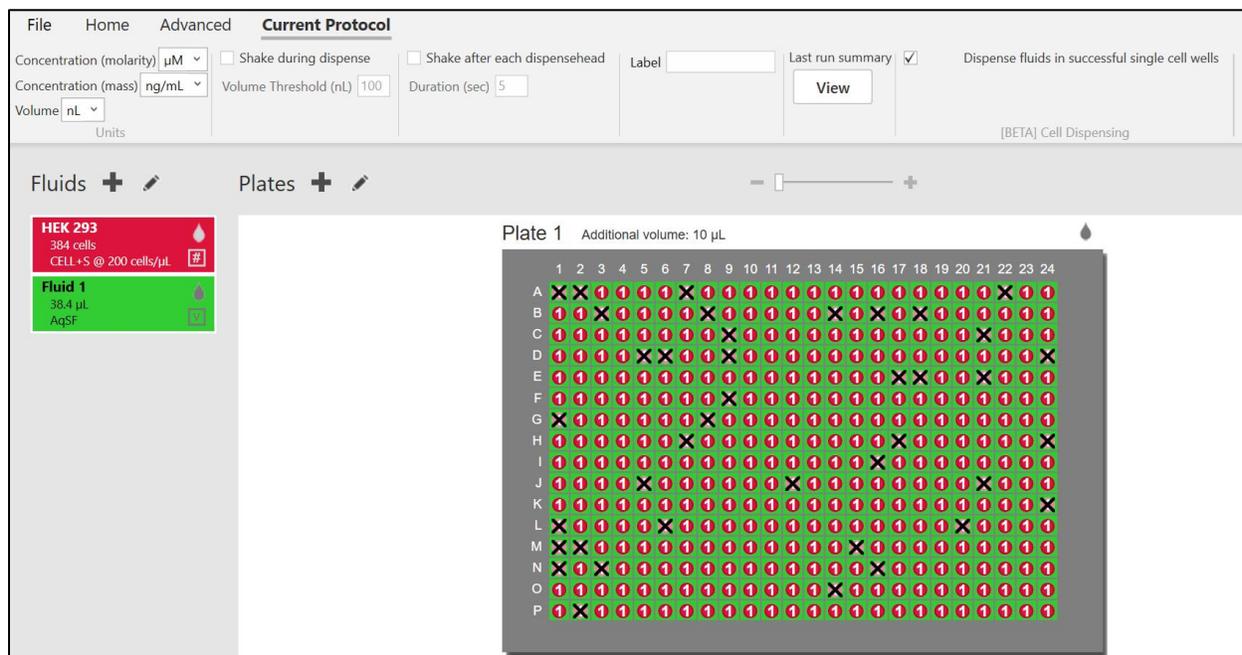
Cell dispense wells

Add the cell fluid to the wells you wish to contain isolated single cells by selecting the wells and using **Set Value** from the Home tab tool bar:



If you want to add reagents to isolated single cell wells in this protocol, there are two options. The figure below shows the main display after a single cell isolation run is performed.

In the **Current Protocol** tab of the Ribbon, there is a check box to **Dispense fluids in successful single cell wells**. If the box is checked prior to the run, the subsequent non-cell fluids will only be dispensed into selected wells that contain a single cell. If the box is unchecked, subsequent non-cell fluids will be dispensed in all selected wells including the non-single cell wells (marked with an X).

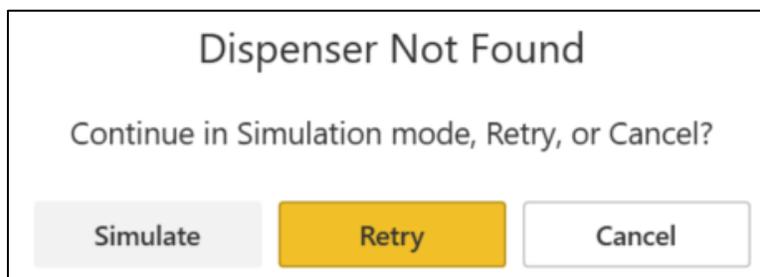


Running a single cell dispense protocol

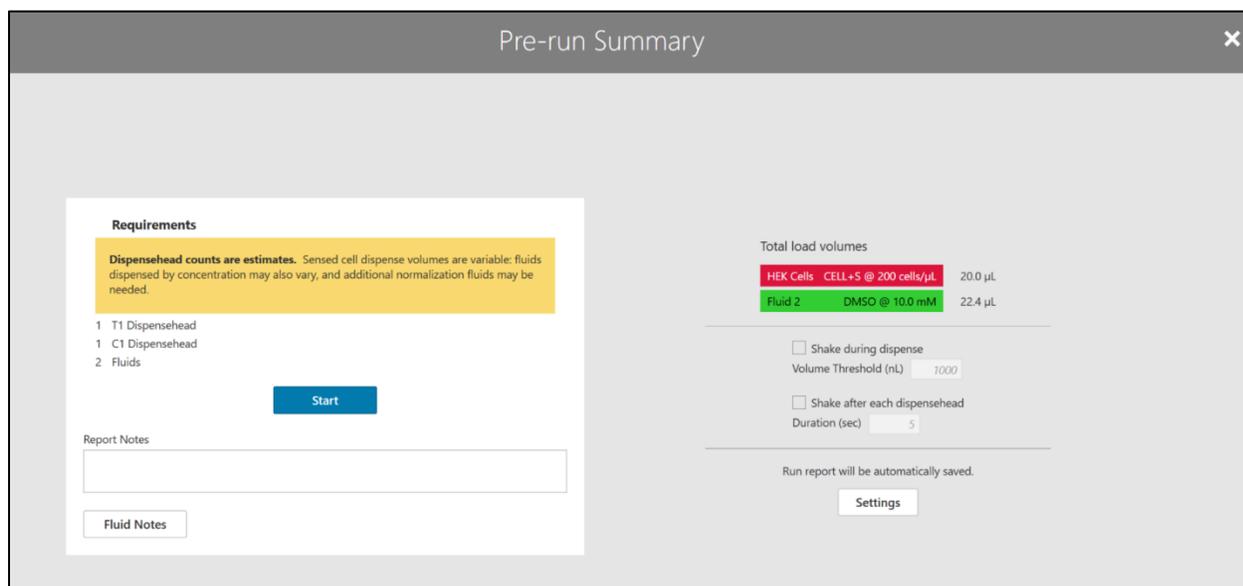
Then click the **Run** button on the Home Tab.



The dispensing software checks for a connection to a dispenser. If the software fails to detect a device, a dialog box is displayed, asking if you want to **Simulate** a run, **Retry** the connection for an actual run, or **Cancel** the run altogether.



The initial **Run** Window is then displayed.



The left area indicates the number of fluids and dispense heads that will be needed for the run. The right side indicates the concentration and volume of cell solution required. The **Start** button initiates the dispensing preparation and the subsequent run.

If text is entered into the Report Notes area, those notes will be written to the report file created after the run. Notes can also be added for each fluid, by clicking the **Fluid Notes** button.

Click **Start** to begin the dispense.



NOTE: To ensure optimum performance, it is recommended that specific procedures be followed for cell stock preparation. Refer to [Appendix F—Best practices for dispensing](#) for more information.

Running a single cell dispense protocol

Refer to the dispenser **Operating Manual** by navigating to the **File** → **Documentation** menu for detailed information about running a single cell dispensing protocol.

Run complete

Run completed

General Dispense Summary

Protocol saved to file
Run by user: AUTH\blakem
Local timezone is (UTC-08:00) Pacific Time (US & Canada)
Simulation started at 11/11/2022 5:45:58 PM
Simulation Completed at 11/11/2022 5:46:33 PM
Total runtime was 35 seconds

Counted Cell Dispense Summary

353 wells successfully dispensed of 384 wells requested (91% successful)

Dispensed Fluids

HEK Cells (CELL+ S @ 200 cells/µL)
20.0 µL loaded
2.7 µL dispensed

Fluid 2 (DMSO @ 10.0 mM)
20.6 µL loaded
17.7 µL dispensed

Plates Used

Plate 1 (384 well Plate)
50 µL additional volume
1% DMSO limit

Report saved to (new) (8).xml
[Open Report File Location](#)

[Park Plate Holder and Close](#) [Close](#)

If report settings are set to generate a report on every run, a message appears to let you know that the report is being generated. For information on what to look for in your report, see [Appendix A—Reports](#).

Note that the Counted Cell Dispense Summary for the plate is shown above in the Run completed screen as well as in the report. Specific wells reported to contain single cells are also shown in the right hand plate graphic.

If a report is not automatically saved, it can later be saved using Save Report from the [Run menu](#).

Click **Close** to return to the Operating Window. Click **Park plate holder and Close** to retract the plate holder before closing. You can always retract the plate holder from the [Run menu](#).

Settings

In the Main Operating Window **File Menu**, click **Settings** to assign **application-wide** parameters and values.

1. Each section is opened by clicking on the section title region. Some sections will scroll to access all settings.
2. Edit the settings via specified drop-down menu choices, manual entry, check box/radio button, or browsing.
3. Once you have updated the settings, click **OK** to save them or **Cancel** to ignore all changes made in the current Settings session.
4. Note that you can override many application settings with protocol-specific settings described in previous chapters.

Default values

There are a number of values which are automatically filled in with defaults that may be changed. For instance, when you create a new protocol or add a fluid or plate, the default settings are used. However, any of the values can be changed when you first set up a protocol or later on in the protocol preparation process.

Default values ⌵

Default protocol values

Molarity concentration units ⌵

Mass concentration units ⌵

Volume units ⌵

Enable shake during dispense

Volume threshold (nL)

Enable shake after each dispensehead

Duration (sec)

Default fluid values

Class ⌵

Concentration ⌵

Dispense by Concentration

Default plate values

Plate type ⌵

DMSO limit (%)

Normalization

Default dispense order ⌵

Additional volumes

Default additional volume (μL) ⌵

Item	Measure	Choices
Default protocol values	Molarity concentration units for new protocols	Drop-down menu
	Mass concentration units for new protocols	Drop-down menu
	Volume units for new protocols	Drop-down menu
	Enable shake during dispense	If checked, shake during dispense will be enabled
	Volume threshold for shake during dispense	Manual entry
	Enable shake after each dispense head	If checked, shake after each dispense head will be enabled
	Shake duration after each dispense head	Manual entry
Default fluid values	Fluid class for new fluids	Drop-down menu
	Concentration and units for new fluids	Manual entry and drop-down menu.
	Dispense mode for new fluids	Toggle button
Default plate values	Plate type for new protocols	Drop-down menu
	DMSO limit for new protocols	Manual entry
Normalization	Default dispense order	Drop-down menu
Additional volumes	Default additional volume for each plate type	Drop-down menu with manual entry. For each well plate type , you can assign a default additional volume.

Display

You can set up the default software parameters for how numbers, concentrations and volumes, will be displayed. Changing these settings does not impact how dispensing occurs, just how numbers are displayed.

Display ⌵

Wells

Concentration significant digits (minimum)

Volume significant digits (minimum)

Fluids

Concentration decimal places

Volume decimal places

Show more information in fluid list

Item	Measure	Choices
Wells	Concentration significant digits (minimum)	Manual entry – 2 to 8
	Volume significant digits (minimum)	Manual entry – 2 to 8
Fluids	Concentration decimal places	Manual entry – maximum of 4
	Volume decimal places	Manual entry – maximum of 4
	Show more information in fluid list	Check to show more fluid information in the fluid list on the main operating window.

Reports

The reports menu allows you to set system defaults for report generation. More information about Reports can be found in [Appendix A—Reports](#).

Reports

Auto-save reports after protocol run.

Do not save aborted protocol runs.

Report name

Name is the protocol name combined with date and time.

Fixed name:

Report formats (select one or more)

Excel compatible XML (*.xml)

Comma separated values (*.csv)

PDF (*.pdf)

Data XML (*.DATA.xml)

Report options

Significant digits

Show all digits.

Round to significant digits.

Include "Don't Dispense" fluids

Include empty wells in "Tabular detail"

Include revision history.

PDF report options

Page size

Include signature lines.

Digitally sign reports ⓘ

Certificate has been selected

Item	Value
Auto-save reports after protocol run	<p>If the box is un-checked, reports must be manually saved after each protocol run (Save Report in Run menu).</p> <p>If the box is checked, you can set the auto-save report parameters and you will not be prompted for report save after each run.</p> <ul style="list-style-type: none"> • Do not save aborted protocol runs: Check this box so that reports are not saved after an aborted run. • Report file name: <ul style="list-style-type: none"> ○ Name is the protocol name combined with date and time. Click this to produce unique report names after each run. ○ Fixed name. Selecting this option will give the same name to each protocol report. Manually enter the desired report filename. Select whether a new file is saved with an appended name – e.g., ReportName(1) or if older files are overwritten. <div data-bbox="500 464 989 646" style="border: 1px solid black; padding: 5px; margin: 10px 0;"> <p>Report name</p> <p><input type="radio"/> Name is the protocol name combined with date and time.</p> <p><input checked="" type="radio"/> Fixed name: <input type="text" value="MyReport"/></p> <p>Report file exists</p> <p><input checked="" type="radio"/> Append to file name to make it unique.</p> <p><input type="radio"/> Overwrite existing report file.</p> </div> • If report file already exists: Select additional naming conventions for reports with the same name. • Report formats: Select one or more output formats.
Report options	<ul style="list-style-type: none"> • To round report data, click the Round to radio button and enter the number of significant digits. <div data-bbox="444 770 748 869" style="border: 1px solid black; padding: 5px; margin: 10px 0;"> <p>Significant digits</p> <p><input type="radio"/> Show all digits.</p> <p><input checked="" type="radio"/> Round to <input type="text" value="4"/> significant digits.</p> </div> <ul style="list-style-type: none"> • Check to include "Don't Dispense" fluid tables in reports. • Check to include empty well lines in the "Tabular detail" table of reports. • Check to include revision history in reports.
PDF report options	<ul style="list-style-type: none"> • PDF reports can be written for Letter or A4 page size. • Check to include signature lines in PDF. • Check to digitally sign reports using a certificate. If checked, you can select a certificate to use for signing.

File locations

The file locations menu allows you to set system defaults for protocol and report folders. This is also useful to recall where reports and protocols are saved on your computer.

Item	Value
Protocol folder	Use the system-assigned default or click the Browse button to search and assign the folder in which protocols are stored.
Report folder	Use the system-assigned default or click the Browse button to search and assign the folder in which reports should be written.

Run

Item	Choices
Default to use high-volume cassettes	The Use high-volume cassettes checkbox on the protocol Run screen will default to this setting.
Show "remove plate lid" reminder	If checked, a reminder will display each time a plate is loaded.
Require and validate plate ID	If checked, a plate ID is required and must be re-entered (and validated) each time the plate is loaded.
Do not show Help panel	If checked, the help panel will not be displayed.

Appendix A—Reports

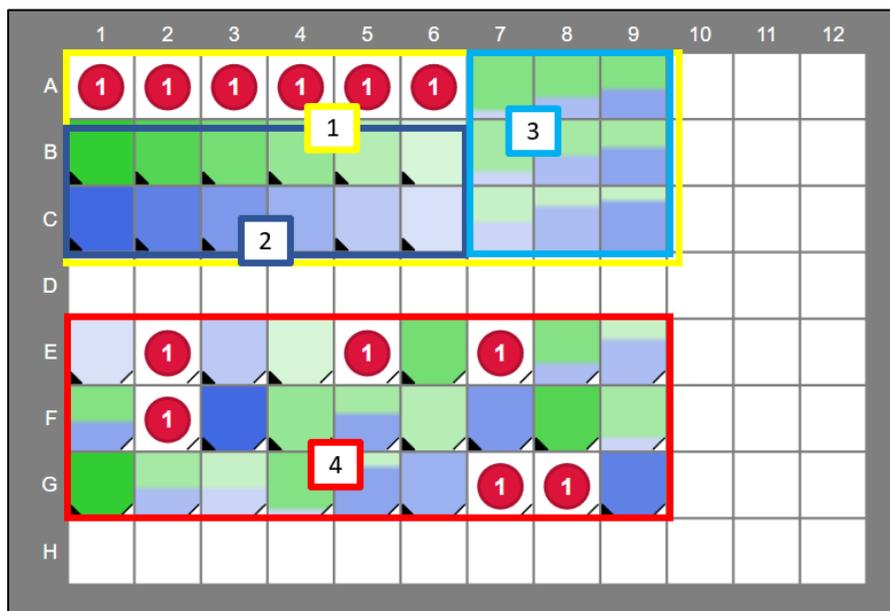
Report file formats

The dispensing software outputs report files in five formats:

- **Excel Compatible XML format (*.xml):** This Excel-specific XML (eXtensible Markup Language) file contains useful details from running a protocol in a file format designed to be opened and viewed in Microsoft Excel. This file includes formatting (page tabs and cell formatting) to improve direct reading by the user. The tabs within this format are described extensively within this appendix.
- **Comma Separated Values (*.csv):** This contains the same information as the Excel XML format, but in a comma delimited format. This format may be more useful for importing protocol information into a LIMS system.
- **Adobe Acrobat PDF format (*.pdf):** This file is intended to be less readily edited and can be generated with blank signature lines if desired for filing. Additionally, these reports can be digitally signed by selecting a certificate.
- **QuantStudio Compatible format (*.DA.csv):** This QuantStudio specific CSV file is formatted to enable importing qPCR plate designs into QuantStudio DA and DA2 software packages to reduce errors associated with recreating plate layouts in separate software packages. qPCR plates are easy to set up in the Digital Dispensing software as described elsewhere in this manual.
- **Data XML (*.DATA.xml):** This contains the same information as the Excel XML format but is in generic XML. This format may be more useful for importing protocol information into a LIMS system.

Example (Excel-compatible XML)

Below is an illustrative example of a protocol that contains multiple fluids, normalization, and randomization. This example is the basis for the report files used throughout the remainder of this Appendix. Descriptions of the regions of this protocol follow below:



1. **Non-Randomized Region:** This region of the protocol (A1-C9) is non-randomized.
2. **Normalized Region:** This region of the protocol (B1-C6) is normalized to the level of the highest volume well within the region

3. **Two-fluid synergy:** This region of the protocol (A7-C9) contains a two-fluid synergy experiment
4. **Randomized Region:** This region of the protocol (E1-G9) is a randomized replicate of the wells shown in region 1 (A1-C12)

*Cells are displayed as the number 1 surrounded by a round symbol (red for this fluid color).

Summary

The Summary tab shows summary information for the protocol. This includes the file location of the protocol, start and finish time for the protocol, and key protocol information, including the dispense order for the normalization fluid (if used), and the settings for plate shaking (if used).

Protocol User	HP Inc\Documents\HP Dispensing\Protocols\user guide report.hpdd	
Simulated	Friday, November 4, 2022	2:49:53 PM
Completed	Friday, November 4, 2022	2:52:17 PM
Conditions		
Software Version	3.5.3	
Time Zone	(UTC-08:00) Pacific Time (US & Canada)	
Normalization	Dispense order	No priority

Fluids

The Fluids tab shows details of the fluids used within the protocol. The top section contains information about all fluids defined within the protocol, including fluids marked as No Dispense. This section also includes the fluid load volume requested by the software, and the dispensed volume.

The second section contains information about the dispense heads and cassettes used within the protocol, including the mapping of fluid to dispense head number. In cases when a fluid is used in more than a single dispense head, the requested load volume and dispensed volume are shown by dispense head.

The bottom section with **Observed Cell Concentration** reports the calculation of the cell stock concentration based upon the number of sensed cells per known volume dispensed. This can be used to adjust cell stocks for improved performance as described in [Appendix F—Best practices for dispensing](#). This calculation will only be displayed if 20 or more single cells have been successfully dispensed into wells.

Fluid	Fluid Name	Stock Concentration	Units	Class	Source Plate	Source Well	No Dispense	Dispense by Volume	Small Molecule	Dispense Heads	Loaded Volume (µL)	Dispensed Volume (µL)
1	Cells	200	cells/µL	CELL+S						1	20	0.052
2	Fluid 1	10	mM	DMSO						2	13.8	11.5
3	Fluid 2	10	mM	DMSO						2	13.8	11.5
4	DMSO normalization			DMSO						1	14.2	12.1

Fluid	Fluid Name	Cassette Type	Cassette	Dispense Head	Fill Time	Loaded Volume (µL)	Dispensed Volume (µL)	Notes
1	Cells	C1	1	1	14:49:58.6	20	0.052	
2	Fluid 1	T1	2	1	14:52:02.0	6.9	5.7	
2	Fluid 1	T1	3	1	14:52:05.0	6.9	5.7	
3	Fluid 2	T1	4	1	14:52:08.0	6.9	5.7	
3	Fluid 2	T1	5	1	14:52:11.0	6.9	5.7	
4	DMSO normalization	T1	6	1	14:52:16.0	14.2	12.1	

Cassette	Cassette ID	Cassette Type	Observed Cell Concentration (cells/µL)
1		C1	190

Plates

The Plates tab shows information for all plates in the protocol, including plates marked as Don't Dispense. This includes plate number (which corresponds to the order of the plates within the protocol) and the defined assay volume of the plate. The **Plate name** is defined by the user within the protocol setup, while the **Plate ID** is defined by the user within the run process the first time the plate is loaded onto the instrument.

Tabular

The Tabular tab shows dispensed concentration and volume in a columnar format. Each **well** within a protocol has a single row in the tab, and each fluid within the protocol has a single concentration column and a single volume column. Composite concentration and volume information columns are also found within the tab. Composite information column headers are colored black, whereas individual-fluid column headers are colored with their corresponding fluid color. Well locations are shown in alphanumeric (“A5”, “C12”) as well as numeric row and column formats. Rows are sorted by plate number, then dispensed row, and then dispensed column. Descriptions of the regions of this report file tab follow below:

Plate ID	Plate	Dispensed Well	Dispensed Row	Dispensed Col	Non-random Well	Non-random Row	Non-random Col	Well Content	2		3		4		5		Total Well Volume (nL)	DMSO %
									Single Fluid Conc/Count	Units	Count (cell)	Conc. (µM)	Conc. (µM)	Dispensed Volume (nL)	Volume (nL)	Volume (nL)		
1	A01	1	1	A01	1	1	1	1	0	cell	0	0	0	23.2	23.2		100000	0.00%
1	A02	1	2	A02	1	2	2	2	0	cell	1	0	0	0.48	0.48		100000	0.00%
1	A03	1	3	A03	1	3	3	3	1	cell	1	0	0	0.48	0.48		100000	0.00%
1	A04	1	4	A04	1	4	4	4	Cells	1	cell	1	0	0.48	0.48		100000	0.00%
1	A05	1	5	A05	1	5	5	5	Cells	1	cell	1	0	0.48	0.48		100000	0.00%
1	A06	1	6	A06	1	6	6	6	Cells	1	cell	1	0	0.48	0.48		100000	0.00%
1	A07	1	7	A07	1	7	7	7	2 Fluids	Non-single cell	0	50	9.98	603.6	503	100	101000	0.00%
1	A08	1	8	A08	1	8	8	8	2 Fluids		0	50	30	806.4	504	302	101000	0.80%
1	A09	1	9	A09	1	9	9	9	2 Fluids		0	50	50	1009.6	505	505	101000	1.00%
1	A10	1	10	A10	1	10	10	10										
1	A11	1	11	A11	1	11	11	11										
1	A12	1	12	A12	1	12	12	12										
1	B01	2	1	B01	2	1	1	Fluid 1	100	µM	0	100	0	1010			101000	1.00%
1	B02	2	2	B02	2	2	2	Fluid 1	80	µM	0	80	0	1010	808		101000	1.00%
1	B03	2	3	B03	2	3	3	Fluid 1	50	µM	0	50	0	1010	606	202	101000	1.00%
1	B04	2	4	B04	2	4	4	Fluid 1	40	µM	0	40	0	1010	404	606	101000	1.00%
1	B05	2	5	B05	2	5	5	Fluid 1	20	µM	0	20	0	1010	202	808	101000	1.00%
1	B06	2	6	B06	2	6	6	Fluid 1	0.00129	µM	0	0.00129	0	1010.013	0.013	1010	101000	1.00%
1	B07	2	7	B07	2	7	7	2 Fluids			0	30	10	401.6	301	100	100000	0.40%
1	B08	2	8	B08	2	8	8	2 Fluids			0	30	30	603.2	302	302	101000	0.80%
1	B09	2	9	B09	2	9	9	2 Fluids			0	30	50	806.4	302	504	101000	0.80%
1	B10	2	10	B10	2	10	10											
1	B11	2	11	B11	2	11	11											
1	B12	2	12	B12	2	12	12											
1	C01	3	1	C01	3	1	1	Fluid 2	100	µM	0	0	100	1010			101000	1.00%
1	C02	3	2	C02	3	2	2	Fluid 2	80	µM	0	0	80	1010	808	202	101000	1.00%
1	C03	3	3	C03	3	3	3	Fluid 2	60	µM	0	0	60	1010	606	404	101000	1.00%
1	C04	3	4	C04	3	4	4	Fluid 2	40	µM	0	0	40	1010	404	606	101000	1.00%
1	C05	3	5	C05	3	5	5	Fluid 2	20	µM	0	0	20	1010	202	808	101000	1.00%
1	C06	3	6	C06	3	6	6	Fluid 2	0.00129	µM	0	0	0.00129	1010.013	0.013	1010	101000	1.00%
1	C07	3	7	C07	3	7	7	2 Fluids			0	9.98	9.98	200			100000	0.20%
1	C08	3	8	C08	3	8	8	2 Fluids			0	10	30	401.6	100	301	100000	0.40%
1	C09	3	9	C09	3	9	9	2 Fluids			0	9.98	50	603.6	100	503	101000	0.80%

- Well Contents** – When a well contains a single fluid with or without a normalization fluid, the composite ‘Fluid Name’ cell for that row will contain that fluid name and will be colored with the fluid color. When a well contains more than a single fluid (not including normalization fluid), the composite ‘Fluid Name’ cell for that row will be named ‘N fluids’, where N is the number of fluids within that well. The cell will be colored white.
- Concentration/count** – When a well contains a single fluid with or without a normalization fluid, the composite ‘Concentration/Count’ cell for that row will contain the concentration (or count if doing cell dispensing) of that single fluid and will be colored with the fluid color. When a well contains more than a single fluid (not including normalization fluid), the composite ‘Concentration/Count’ cell for that row will be blank and will be colored white.
- Count (Cells)** – When a well is reported to contain a single cell, the “Count” for that row is set to one, when it is reported to contain no single cells it reports a zero, and when that well is reported to contain more than one cell that row is set to “Non-single cell.”
- Total dispensed volume and total well volume** – When a well contains a single fluid without a normalization fluid, the “Total Dispensed Volume” cell for that row will contain the volume of that single fluid and will be colored with the fluid color. When a well contains more than a single fluid (including normalization fluid), the composite “Total Volume” cell for that row will include the total volume, including the volume of the normalization fluid, and will be colored white. The Total Well Volume cell for that row is the sum of the dispensed volume and the additional volume set in the protocol.
- DMSO %** - Only DMSO fluids and DMSO normalization are included in this calculation.

Tabular detail

The Tabular Detail tab shows dispensed concentration and volume in an alternative columnar format. Each **dispense** within a protocol has a single row in the tab, and there are single columns within the protocol for Dispensed Concentration and Dispensed Volume of each dispense. This tab may be more useful for pulling protocol dispense information into a LIMS system. Normalization fluids do not have a dispensed concentration. The fluid number, fluid name, cassette number, and dispense head for each dispense are colored with their corresponding fluid color.

Well locations are shown in alphanumeric (“A5”, “C12”) as well as numeric row and column formats. Rows are sorted by plate number, then dispensed row, and then dispensed column. In cases where there is a single fluid per well, each well will have a single row. In cases where there are multiple fluids per well, each well will have as many rows as it has fluids, including a separate row for the normalization fluid. In the example below, well B2 has two separate dispense events and two rows to correspond to fluid 1 and the normalization fluid.

Plate ID	Plate	Dispensed Well	Dispensed Row	Dispensed Col	Non-random Well	Non-random Row	Non-random Col	Fluid	Fluid Name	Cassette	Dispense Head	Dispensed Conc/Count	Units	Dispensed Volume	Units	Total Well Volume
1	1	A01	1	1	A01	1	1	1	Cells	1	1	0	cell	23.2	nL	100000
1	1	A02	1	2	A02	1	2	1	Cells	1	1	1	cell	0.48	nL	100000
1	1	A03	1	3	A03	1	3	1	Cells	1	1	1	cell	0.48	nL	100000
1	1	A04	1	4	A04	1	4	1	Cells	1	1	1	cell	0.48	nL	100000
1	1	A05	1	5	A05	1	5	1	Cells	1	1	1	cell	0.48	nL	100000
1	1	A06	1	6	A06	1	6	1	Cells	1	1	1	cell	0.48	nL	100000
1	1	A07	1	7	A07	1	7	2	Fluid 1	2	1	Non-single cell		9.64	nL	100000
1	1	A07	1	7	A07	1	7	3	Fluid 2	4	1	50	µM	503	nL	101000
1	1	A08	1	8	A08	1	8	2	Fluid 1	2	1	9.98	µM	100	nL	101000
1	1	A08	1	8	A08	1	8	3	Fluid 2	4	1	50	µM	504	nL	101000
1	1	A09	1	9	A09	1	9	2	Fluid 1	2	1	30	µM	302	nL	101000
1	1	A09	1	9	A09	1	9	3	Fluid 2	4	1	50	µM	505	nL	101000
1	1	A10	1	10	A10	1	10					50	µM	505	nL	101000
1	1	A11	1	11	A11	1	11									
1	1	A12	1	12	A12	1	12									
1	1	B01	2	1	B01	2	1	2	Fluid 1	2	1	100	µM	1010	nL	101000
1	1	B02	2	2	B02	2	2	2	Fluid 1	2	1	80	µM	808	nL	101000
1	1	B02	2	2	B02	2	2	4	DMSO norms	6	1			202	nL	101000
1	1	B03	2	3	B03	2	3	2	Fluid 1	2	1	60	µM	606	nL	101000
1	1	B03	2	3	B03	2	3	4	DMSO norms	6	1			404	nL	101000
1	1	B04	2	4	B04	2	4	2	Fluid 1	2	1	40	µM	404	nL	101000
1	1	B04	2	4	B04	2	4	4	DMSO norms	6	1			606	nL	101000
1	1	B05	2	5	B05	2	5	2	Fluid 1	2	1	20	µM	202	nL	101000
1	1	B05	2	5	B05	2	5	4	DMSO norms	6	1			808	nL	101000
1	1	B06	2	6	B06	2	6	2	Fluid 1	2	1	0.00129	µM	0.013	nL	101000

Figure 1 Tabular Detail tab example (row A and B)

Dispense head – When the amount of any protocol fluids exceeds the maximum amount of fluid that can be loaded into a single dispense head, dispensing of the fluid is split between two or more dispense heads. The software distributes the required amount of fluid between these two or more dispense heads approximately equally. Occasionally, this causes a transition between dispense heads to occur within a single well. In this case, the report file will report the word “Multiple” within the dispense head column.

The Tabular Detail tab contains a few other useful pieces of information highlighted below. Description follows:

Plate ID	Plate	Dispensed Well	Dispensed Row	Dispensed Col	Non-random Well	Non-random Row	Non-random Col	Fluid	Fluid Name	Cassette	Dispense Head	Dispensed Conc/Count	Units	Dispensed Volume	Units	Total Well Volume
1	1	E01	5	1	G06	7	6	3	Fluid 2	5	1	0.00129	µM	0.013	nL	101000
1	1	E01	5	1	G06	7	6	4	DMSO norms	6	1			1010	nL	101000
1	1	E02	5	2	E02	5	2	1	Cells	1	1	1	cell	0.48	nL	100000
1	1	E03	5	3	G05	7	5	3	Fluid 2	5	1	20	µM	202	nL	101000
1	1	E03	5	3	G05	7	5	4	DMSO norms	6	1			808	nL	101000
1	1	E04	5	4	F06	6	6	2	Fluid 1	3	1	0.00129	µM	0.013	nL	101000
1	1	E04	5	4	F06	6	6	4	DMSO norms	6	1			1010	nL	101000
1	1	E05	5	5	E03	5	3	1	Cells	1	1	1	cell	0.48	nL	100000
1	1	E06	5	6	F03	6	3	2	Fluid 1	3	1	60	µM	606	nL	101000
1	1	E06	5	6	F03	6	3	4	DMSO norms	6	1			404	nL	101000
1	1	E07	5	7	E06	5	6	1	Cells	1	1	1	cell	0.48	nL	100000
1	1	E08	5	8	E08	5	8	2	Fluid 1	3	1	50	µM	504	nL	101000
1	1	E08	5	8	E08	5	8	3	Fluid 2	5	1	30	µM	302	nL	101000
1	1	E09	5	9	G08	7	8	2	Fluid 1	3	1	10	µM	100	nL	100000
1	1	E09	5	9	G08	7	8	3	Fluid 2	5	1	30	µM	301	nL	100000
1	1	E10	5	10	E10	5	10									
1	1	E11	5	11	E11	5	11									
1	1	E12	5	12	E12	5	12									

Figure 2 Tabular Detail tab randomized example (rows E)

Non-random columns provide a tool for re-sorting the dispensing information from a randomized-region of a protocol back to the original non-random configuration. In the example protocol, the wells in region 1 (A1-C9) are duplicated in region 5 (E1-G9) and then randomized. Once plate-reader data has been matched to the “Dispensed Concentration” or the “Dispensed Volume” columns, then the data can be re-sorted using the “NonRandom rows” and “NonRandom col” columns.

It should be noted that when manually analyzing randomized data, it can often be just as useful to match the report file rows to plate reader output rows and then to sort first by fluid name and then by fluid concentration.

Dispensed Conc

The Dispensed Conc tab shows the dispensed concentration in each well in a spatially-representative plate format. This is done first in a composite section, and then in fluid-by-fluid sections. If a well in the composite section contains only a single fluid, that well is colored with the color code of that fluid. If a well in the composite section contains multiple fluids, the number of fluids dispensed in the well is indicated. If a well was requested to have a single cell but the system did not report a single cell in that well, that well does not contain an entry for the cell fluid as in wells A1 and A6 below.

If a protocol contains a single plate, then the tab will be named “1 Dispensed Conc.” If a protocol contains multiple plates, then subsequent plates in the protocol will be named with the corresponding plate number, i.e. “2 Dispensed Conc,” “3 Dispensed Conc,” etc.

(Composite)	1	2	3	4	5	6	7	8	9
A		1	1	1	1		2 Fluids	2 Fluids	2 Fluids
B	100	80	60	40	20	0.00129	2 Fluids	2 Fluids	2 Fluids
C	100	80	60	40	20	0.00129	2 Fluids	2 Fluids	2 Fluids
D									
E	0.00129	1	20	0.00129	1	60	1	2 Fluids	2 Fluids
F	2 Fluids	1	100	40	2 Fluids	20	60	80	2 Fluids
G	100	2 Fluids	2 Fluids	2 Fluids	2 Fluids	40	1	1	80
H									
Cells (cell)	1	2	3	4	5	6	7	8	9
A		1	1	1	1				
B									
C									
D									
E		1			1		1		
F		1							
G							1	1	
H									
Fluid 1 (µM)	1	2	3	4	5	6	7	8	9
A							50	50	50
B	100	80	60	40	20	0.00129	30	30	30
C							9.98	10	9.98
D									
E				0.00129		60		50	10
F	50			40	30	20		80	30
G	100	30	10	50	9.98				
H									

Dispensed Volume

The Dispensed Volume tab shows the dispensed volume in each well in a spatially-representative plate format. This is done first in a composite section and then in fluid-by-fluid sections. If a well in the composite section contains only a single fluid, that well is colored with the color code of that fluid. If a well in the composite

section contains multiple fluids, then that well is non-colored and contains the cumulative volume of all the fluids in the well. If a normalization fluid is used, the total volume reported in the composite section includes the volume of the normalization fluid.

If a protocol contains a single plate, then the tab will be named “1 Dispensed Volume.” If a protocol contains multiple plates, then subsequent plates in the protocol will be named with the corresponding plate number, i.e. “2 Dispensed Volume,” “3 Dispensed Volume,” etc.

(Composite)	1	2	3	4	5	6	7	8	9
A	23.2	0.48	0.48	0.48	0.48	9.64	604	806	1010
B	1010	1010	1010	1010	1010	1010	402	603	806
C	1010	1010	1010	1010	1010	1010	200	402	604
D									
E	1010	0.48	1010	1010	0.48	1010	0.48	806	402
F	1010	0.48	1010	1010	806	1010	1010	1010	402
G	1010	603	201	604	604	1010	0.48	0.48	1010
H									
Cells (nL)	1	2	3	4	5	6	7	8	9
A	23.2	0.48	0.48	0.48	0.48	9.64			
B									
C									
D									
E		0.48			0.48		0.48		
F		0.48							
G							0.48	0.48	
H									
Fluid 1 (nL)	1	2	3	4	5	6	7	8	9
A							503	504	505
B	1010	808	606	404	202	0.013	301	302	302
C							100	100	100
D									
E				0.013		606		504	100
F	505			404	302	202		808	301
G	1010	302	100	503	100				
H									

Start Time

The Start Time tab shows the dispense start time for each well in a spatially-representative plate format for each plate within a protocol. This is done first in a composite section, and then in fluid-by-fluid sections. If a well in the composite section contains only a single fluid, that well is colored with the color code of that fluid. If a well in the composite section contains multiple fluids, then that well is non-colored and contains the starting dispense time of the first fluid dispensed into that well.

If a protocol contains a single plate, then the tab will be named “1 Start Time.” If a protocol contains multiple plates, then subsequent plates in the protocol will be named with the corresponding plate number, i.e. “2 Start Time,” “3 Start Time,” etc.

(Composite)	1	2	3	4	5	6	7	8	9
A	15:08:29.1	15:08:29.1	15:08:29.2	15:08:29.2	15:08:29.2	15:08:29.3	15:08:32.9	15:08:32.9	15:08:33.0
B	15:08:32.9	15:08:32.9	15:08:33.2	15:08:33.1	15:08:33.1	15:08:33.1	15:08:33.1	15:08:33.0	15:08:33.0
C	15:08:38.7	15:08:38.8	15:08:38.9	15:08:39.0	15:08:39.0	15:08:39.0	15:08:33.2	15:08:33.2	15:08:33.2
D									
E	15:08:39.1	15:08:29.3	15:08:41.6	15:08:33.3	15:08:29.4	15:08:35.9	15:08:29.4	15:08:35.9	15:08:35.9
F	15:08:36.1	15:08:29.5	15:08:41.6	15:08:36.1	15:08:36.0	15:08:36.0	15:08:41.7	15:08:35.9	15:08:36.0
G	15:08:35.8	15:08:36.1	15:08:36.2	15:08:36.2	15:08:36.3	15:08:41.9	15:08:29.5	15:08:29.6	15:08:41.6
H									
Cells	1	2	3	4	5	6	7	8	9
A	15:08:29.1	15:08:29.1	15:08:29.2	15:08:29.2	15:08:29.2	15:08:29.3			
B									
C									
D									
E		15:08:29.3			15:08:29.4		15:08:29.4		
F		15:08:29.5							
G							15:08:29.5	15:08:29.6	
H									
Fluid 1	1	2	3	4	5	6	7	8	9
A							15:08:32.9	15:08:32.9	15:08:33.0
B	15:08:32.9	15:08:32.9	15:08:33.2	15:08:33.1	15:08:33.1	15:08:33.1	15:08:33.1	15:08:33.0	15:08:33.0
C							15:08:33.2	15:08:33.2	15:08:33.2
D									
E				15:08:33.3		15:08:35.9		15:08:35.9	15:08:35.9
F	15:08:36.1			15:08:36.1	15:08:36.0	15:08:36.0		15:08:35.9	15:08:36.0
G	15:08:35.8	15:08:36.1	15:08:36.2	15:08:36.2	15:08:36.3				
H									

Events

The Events tab shows key event information for the protocol. This includes, start and finish time for the protocol, when plates and cassettes were loaded, when dispense heads were filled with fluid, and when dispense heads started and finished dispensing. The starting dispense time per dispense head is shown in column A, while the ending dispense time is shown in column B.

	2:53:22 PM	Started
	2:53:23 PM	Loaded plate 1.
	2:53:23 PM	Loaded cassette 1.
	2:53:25 PM	Filled dispensehead 1 with 'Fluid 1'.
	2:53:25 PM	Filled dispensehead 2 with 'Fluid 2'.
	2:53:26 PM	Filled dispensehead 3 with 'Fluid 3'.
2:53:25 PM	2:53:27 PM	'Fluid 1' dispensed from head 1.
	2:53:27 PM	Filled dispensehead 4 with 'Fluid 3'.
	2:53:27 PM	Filled dispensehead 5 with 'DMSO normalization'.
2:53:27 PM	2:53:28 PM	'Fluid 2' dispensed from head 2.
2:53:28 PM	2:53:28 PM	'Fluid 3' dispensed from head 3.
2:53:28 PM	2:53:29 PM	'Fluid 3' dispensed from head 4.
2:53:29 PM	2:53:29 PM	'DMSO normalization' dispensed from head 5.
	2:53:29 PM	Completed

Appendix B—Moving data between the dispensing software and other programs

General copy and paste

Copying and pasting protocol data within the dispensing software and from the software to the clipboard is contextual, with two different modes depending on what is being copied and pasted:

- **Fluid list:** Copying and pasting fluids is available by right clicking on the Fluid list or through the Fluids section of the Advanced ribbon.
- **Well plate:** Copying and pasting wells is available by right clicking within the well plate, through the clipboard section of the Home ribbon, and through the global shortcuts (i.e. CTRL-C, CTRL-X, CTRL-V, etc.).

While this appendix illustrates moving protocol data between the dispensing software and Microsoft Excel, the techniques shown can be used to move between the software and other applications as well.



NOTE: When pasting from Excel, the display format may limit the significant digits for numbers.

Fluid list copy and paste



Below is an illustrative example of a Fluid list with three fluids. This figure uses the “Show More” feature of the fluid list (right-click on the list to change).

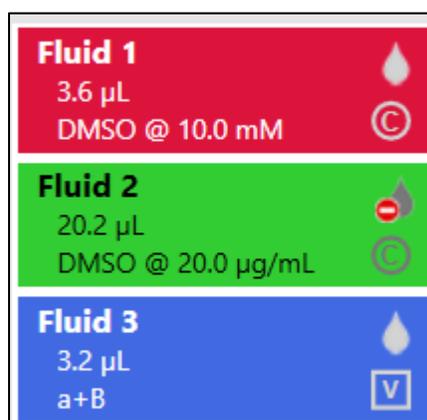


Figure 3 Example fluids list

- **Fluid Name:** For red fluid, this is “Fluid 1”
- **Class:** For the blue fluid, this is “Aqueous + Brij 35.”

- **Stock Concentration and Units:** For red fluid, this is 10 mM; for the green fluid, this is 20 µg/mL; for the blue fluid there is no stock concentration.
- **Source Plate:** For red fluid, this is “Source Plate1” (hover the cursor over fluid to see details). This is the optional, user provided, name of the stock compound source plate or vial.
- **Source Well:** For red fluid, this is “B17.” This is the optional, user provided, well location within a source plate.
- **Dispense by Concentration/Volume:** The red and green fluids are dispensed by concentration; the blue fluid is dispensed by volume (hover the cursor over the icon to see description).
- **Dispense/No Dispense:** For red fluid, this is “Dispense”; for the green fluid, this is “No Dispense” (hover the cursor over the icon to see description). A “Dispense” fluid will be dispensed during the running of the protocol; a “No Dispense” fluid will not be dispensed.

Fluid list copy and paste to external application

Clicking **Copy All Fluids** will copy all of the contents of the Fluid list to the clipboard. Pasting the information from the fluids list in [Figure 3](#) into Microsoft Excel yields the data shown in [Figure 4](#). Each row corresponds to one fluid in the Fluid list and the rows are sorted in the order displayed in the Fluid list. Headers are included for reference and should not be selected when copying from an external application.

Fluid name	Stock conc.	Stock conc. units	Class	Source plate	Source well	No Dispense	Dispense by Volume
Fluid 1	10	mM	DMSO	Source Plate1	B17		
Fluid 2	20	µg/mL	DMSO	Mother Plate	G11	No Dispense	
Fluid 3			a+B	Daughter Plate	E9		By Volume

Figure 4 Example fluids list pasted into Excel

Paste fluid information to modify the existing fluid list

Fluid information can be manipulated in another application and then pasted into the dispenser application using the **Paste Fluids** function on the **Advanced** ribbon tab. When a fluid name is unchanged, the software recognizes the pasted information as an existing fluid and other information associated with that fluid will be modified if necessary. For example, pasting the 3x7 array of cells shown in the left portion of [Figure 5](#) yields the updated fluids list shown in the right portion of [Figure 5](#). A few details follow:

- If the fluid names being pasted back into the Fluid list match the names already in the Fluid list, and the selection region includes other fields of the Fluid list, then these other fields are overwritten with the new values. If concentration (and units) or fluid class are blank, however, the existing fluid values will not be changed.
- Specify a fluid as “No dispense” by including any character (any non-blank value) in the No Dispense column.
- Specify a fluid as “Dispense by volume” by including any character (any non-blank value) in the Dispense by Volume column.

Fluid 1	10 mM	DMSO	Source Plate1	B17	x
Fluid 2	32 µg/mL	DMSO			
Fluid 3					

Fluid 1
3.6 µL
DMSO @ 10.0 mM

Fluid 2
12.5 µL
DMSO @ 32.0 µg/mL

Fluid 3
3.2 µL
a+B

Figure 5 Example Fluid list pasted back into dispensing software with fluid details modified

Since the fluid list was modified, the user is warned before Paste Fluids is completed.



TIP: Do not select column headings when pasting fluids from Excel.

Paste new fluids

When using the **Paste fluid** function, if a fluid has a name that is not already contained in the Fluid list, the software interprets the pasted information as belonging to a new fluid. In this case, the information associated with the new fluid is appended to the Fluid list. Pasting the 3x7 array of cells shown in the left portion of [Figure 6](#) yields the updated Fluid list shown in the right portion of [Figure 6](#).

circle	10 mM	DMSO	Source Plate1	B17	x
square	32 µg/mL	DMSO			
triangle					

Fluid 1
3.6 µL
DMSO @ 10.0 mM

Fluid 2
20.2 µL
DMSO @ 20.0 µg/mL

Fluid 3
3.2 µL
a+B

Circle
0 µL
DMSO @ 10.0 mM

Square
0 µL
DMSO @ 32.0 µg/mL

Triangle
0 µL
DMSO @ 10.0 mM

Figure 6 Example of new fluids pasted into the dispensing software, appended to the Fluid list

Replace existing fluids

When a protocol is repeatedly used as a template, and only the fluids are changing between protocol runs, the fluids can be changed using the **Replace fluids** function. This allows fluid information to be replaced without modifying any well fluids associated with the fluids already in the Fluid list. When starting with the Fluid list on the left portion of [Figure 7](#), copying the 2x1 array of cells shown in the center portion of [Figure 7](#) and using the **Replace fluids** function yields the Fluid list shown on the right portion of [Figure 7](#).

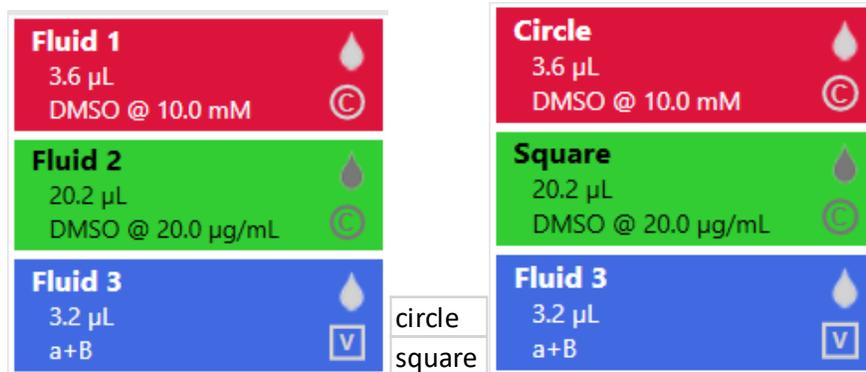
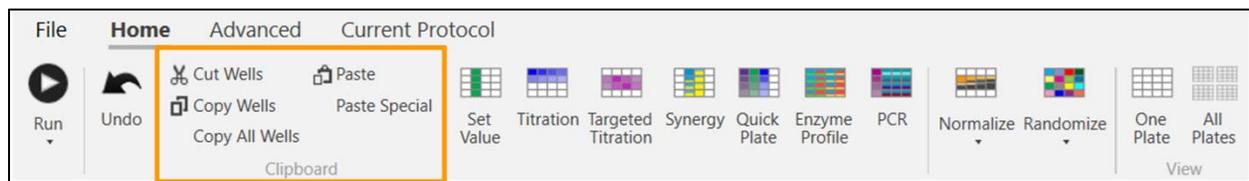
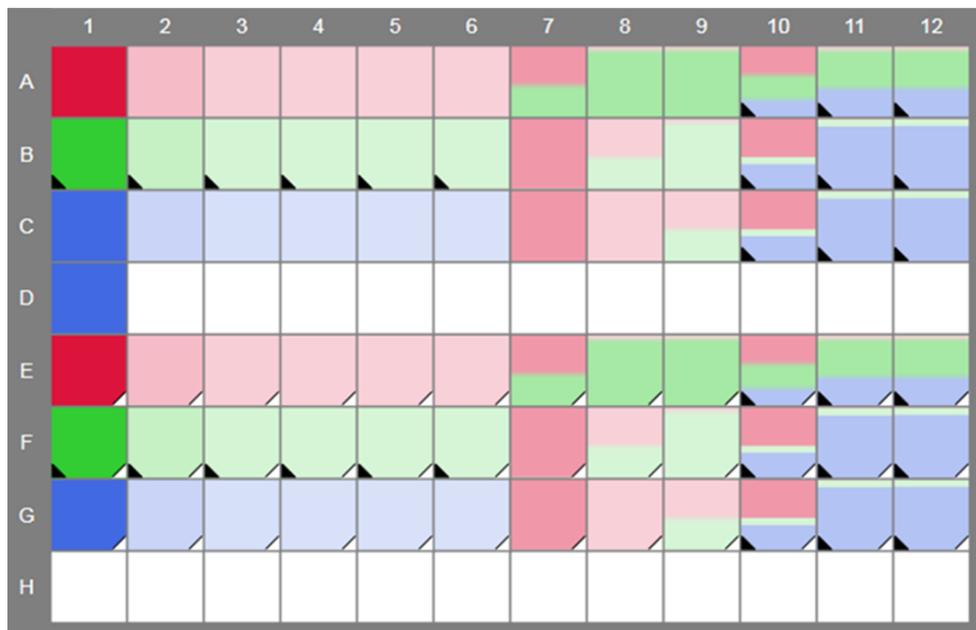


Figure 7 Original Fluid list [left], copy region with new fluid names [center], and resulting Fluid list after using Replace fluid function [right]

Well plate copy and paste



Below is an illustrative example of a well plate grid from a protocol. This protocol is used as a starting point for the following examples:



Copy wells to external application

Clicking **Copy Wells** will copy the contents of the selected wells to the clipboard. Pasting this information into Microsoft Excel yields data like that shown in [Figure 8](#). For each fluid in the plate, a representative well plate template with column and row headers is created. For fluids dispensed by volume, the values pasted in each cell within the template correspond to the volume of the respective fluid in current protocol units. For fluids dispensed by Concentration, the values pasted in each cell within the template correspond to the concentration of the respective fluid in current protocol units. In the row above each fluid template is information for that fluid, specifically (from left to right): Fluid name, Stock Concentration and Units, Class, Source Plate, Source Well, No Dispense, and Dispense Mode. A few additional details follow:

- **Copy Wells** does not function when viewing randomized.
- Normalization information will not be copied or pasted to an external application (e.g. Microsoft Excel).
- Randomization information is not copied to the clipboard.
- **Copy Wells** only copies the selected wells – it does not copy all the well contents of the plate (**Copy All Wells** copies all the well contents of the plate)

Fluid 1		10 mM	DMSO	First Sourc	E7								
μM	1	2	3	4	5	6	7	8	9	10	11	12	
A	100	10.53874	1.11065	0.117049	0.012335	0.0013	30	0.197484	0.0013	30	0.197484	0.0013	
B							30	0.197484	0.0013	30	0.197484	0.0013	
C							30	0.197484	0.0013	30	0.197484	0.0013	
D													
E	100	10.53874	1.11065	0.117049	0.012335	0.0013	30	0.197484	0.0013	30	0.197484	0.0013	
F							30	0.197484	0.0013	30	0.197484	0.0013	
G							30	0.197484	0.0013	30	0.197484	0.0013	
H													
Fluid 2		10 mM	DMSO	Second Sc	G11								
nL	1	2	3	4	5	6	7	8	9	10	11	12	
A							300	300	300	300	300	300	
B	1000	105.3874	11.1065	1.170485	0.123354	0.013	1.974842	1.974842	1.974842	1.974842	1.974842	1.974842	
C							0.013	0.013	0.013	0.013	0.013	0.013	
D													
E							300	300	300	300	300	300	
F	1000	105.3874	11.1065	1.170485	0.123354	0.013	1.974842	1.974842	1.974842	1.974842	1.974842	1.974842	
G							0.013	0.013	0.013	0.013	0.013	0.013	
H													
Fluid 3		20 μg/mL	DMSO										
ng/mL	1	2	3	4	5	6	7	8	9	10	11	12	
A										50	50	50	
B										50	50	50	
C	200	21.07748	2.221301	0.234097	0.024671	0.0026				50	50	50	
D	200												
E										50	50	50	
F										50	50	50	
G	200	21.07748	2.221301	0.234097	0.024671	0.0026				50	50	50	
H													

Figure 8 Output of **Copy Wells** function after pasting into Microsoft Excel

Paste structured well information

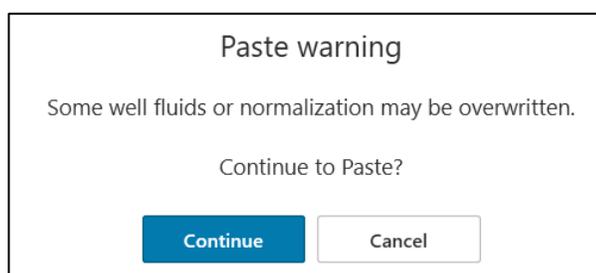
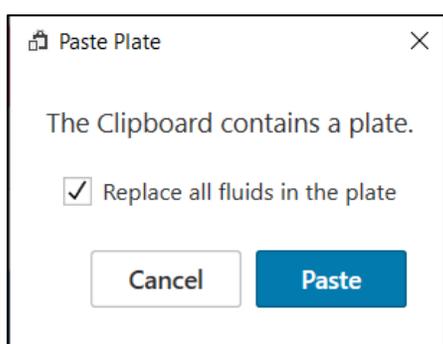
Well information can be manipulated in Microsoft Excel and then pasted into the dispensing software using the **Paste** function. Values in each cell are interpreted using the units in the upper-left corner of the fluid headers. The units may be concentration or volume units.



NOTE: To paste raw concentration or volume values, see the [Paste unstructured well information](#) section.

A few additional details:

- If the entire well plate template region is copied from Microsoft Excel onto the clipboard, including the fluid header row above each fluid region, then the **Paste** function in the software recognizes the template as being a plate and displays the following messages:



- If **Replace all fluids in the plate** is checked, then all existing well contents on the plate are deleted and replaced with the new well values
- If **Replace all fluids in the plate** is unchecked, then existing well contents on the plate for fluids in the clipboard are replaced, but existing well contents for fluids not on the clipboard remain. In this case, if a well already contains Fluid 1, for example, and the paste region for that well contains Fluid 8, the resulting well will contain both Fluid 1 and Fluid 8.
- If only a portion of a well plate is selected when copying from Microsoft Excel, then only the wells within the selected region will be written upon pasting into the software. Wells outside of the selected region will not be overwritten. For example, if only the upper 5x3 grid of wells from the example protocol shown in the left half of [Figure 9](#) are pasted back into the protocol onto a blank plate, then the message shown in the right half of [Figure 9](#) is shown. The final message shown in the bottom half of [Figure 9](#) warns users that existing fluids in the plate area will be overwritten.

Fluid 1	10 mM	DMSO	
μM	1	2	3
A	100	10.53874	1.11065
B			
C			
D			
E	100	10.53874	1.11065



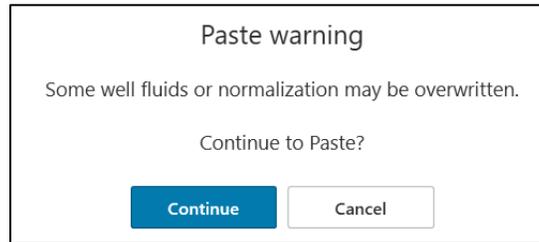


Figure 9 Paste a partial plate

Clicking continue yields the plate shown below:

	1	2	3	4	5	6	7	8	9	10	11	12
A												
B												
C												
D												
E												
F												
G												
H												

- Only the fluid name is required in the row above the template header, all other fluid information is optional.
- Changing the fluid information in the row above the template header will modify the values in the Fluid list. Empty cells in the fluid-information row will be ignored.



NOTE: Empty fluid information cells are treated differently than they are when pasting into the Fluid list ([Paste fluid information to modify the existing fluid list](#)).

Modify well information example

In this example, the original data in [Figure 8](#) has been modified as shown in [Figure 10](#). Rows of Fluid 1 and Fluid 2 have been swapped in columns 1-6. In addition, another copy of the Fluid 1 columns 1-6 has been added to row H. Only Fluids 1 and 2 (not Fluid 3) are copied to the clipboard and pasted into the dispensing software.

Fluid 1	10 mM	DMSO	First Sourc(E7)										
μ M	1	2	3	4	5	6	7	8	9	10	11	12	
A							30	0.197484	0.0013		30	0.197484	0.0013
B	100	10.53874	1.11065	0.117049	0.012335	0.0013	30	0.197484	0.0013	30	0.197484	0.0013	
C							30	0.197484	0.0013	30	0.197484	0.0013	
D													
E							30	0.197484	0.0013	30	0.197484	0.0013	
F	100	10.53874	1.11065	0.117049	0.012335	0.0013	30	0.197484	0.0013	30	0.197484	0.0013	
G							30	0.197484	0.0013	30	0.197484	0.0013	
H	100	10.53874	1.11065	0.117049	0.012335	0.0013							

Fluid 2	10 mM	DMSO	Second Sc G11			By Volume						
nL	1	2	3	4	5	6	7	8	9	10	11	12
A	1000	105.3874	11.1065	1.170485	0.123354	0.013	300	300	300	300	300	300
B							1.974842	1.974842	1.974842	1.974842	1.974842	1.974842
C							0.013	0.013	0.013	0.013	0.013	0.013
D												
E	1000	105.3874	11.1065	1.170485	0.123354	0.013	300	300	300	300	300	300
F							1.974842	1.974842	1.974842	1.974842	1.974842	1.974842
G							0.013	0.013	0.013	0.013	0.013	0.013
H												

Figure 10 Modified Data in Microsoft Excel

Figure 11 shows the result in the dispensing software, with the original plate in the left portion of the figure, and the modified plate in the right portion of the figure. In addition to seeing that the red and green fluids within columns 1-6 have changed places, it can be seen that the original randomization definitions are unchanged, and the normalization and blue fluids have been eliminated.



Figure 11 Original and then modified plate in dispensing software

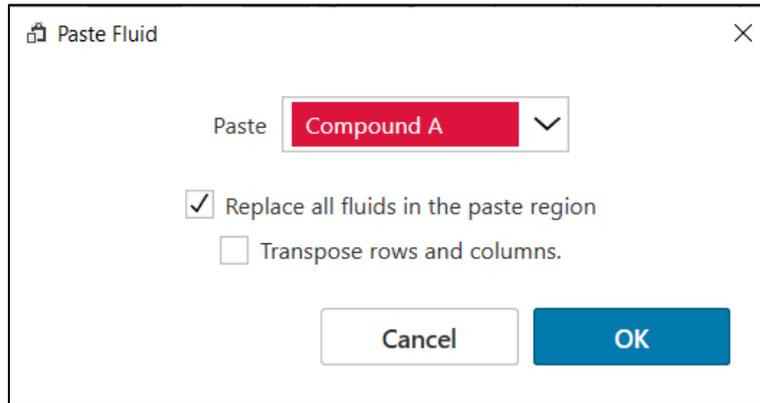
Paste unstructured well information

Well information can be created directly in Microsoft Excel without template information and then pasted into the dispensing software using the **Paste** function. When unstructured well information is pasted, the user will be prompted by the software to identify what fluid to associate with the pasted wells. Pasting will be done in current mode and units for the selected fluid.

In this example, the following array of values is created in Microsoft Excel and copied to the clipboard:

100	50	20	10
5	2	1	0.5
0.2	0.1	0.05	0

When a destination location is selected within a well plate in the Main Operating Window and either **Paste** or **Paste Special** is selected, the software will respond with the following prompt and the user must select a fluid from the drop-down list to paste.



- If **Replace all fluids in the paste region** is checked, then all existing well contents are replaced with the new well values in the paste region.
- If **Transpose rows and columns** is checked, the rows and columns are transposed during paste.

If no fluids exist within the Fluid list, pasting of unstructured well information is not allowed. In this example, well B2 is identified as the destination location, and Fluid 1 is identified as the Paste Fluid. [Figure 12](#) shows the result in the protocol.

	1	2	3	4	5	6	7	8	9	10	11	12
A												
B												
C												
D												
E												
F												
G												
H												

Figure 12 Result of pasting in the example unstructured data into well B2 with Fluid 1 selected.

Appendix C—Matching randomized layout to plate reader data

Once a randomized protocol has been created and run on the dispenser, the dispensing profile needs to be joined with corresponding plate-reader data for analysis. The following shows an example of this procedure.



TIP: The **Data Merge utility**, included in the dispensing software install package, can more easily be used to combine plate reader CSV data with dispense data from an HP report file.

Step 1: Create and run a randomized protocol

[Figure 13](#) shows a typical titration protocol before and after it has been randomized. The randomized protocol is run on the dispenser and the resultant plates are read on a plate reader.

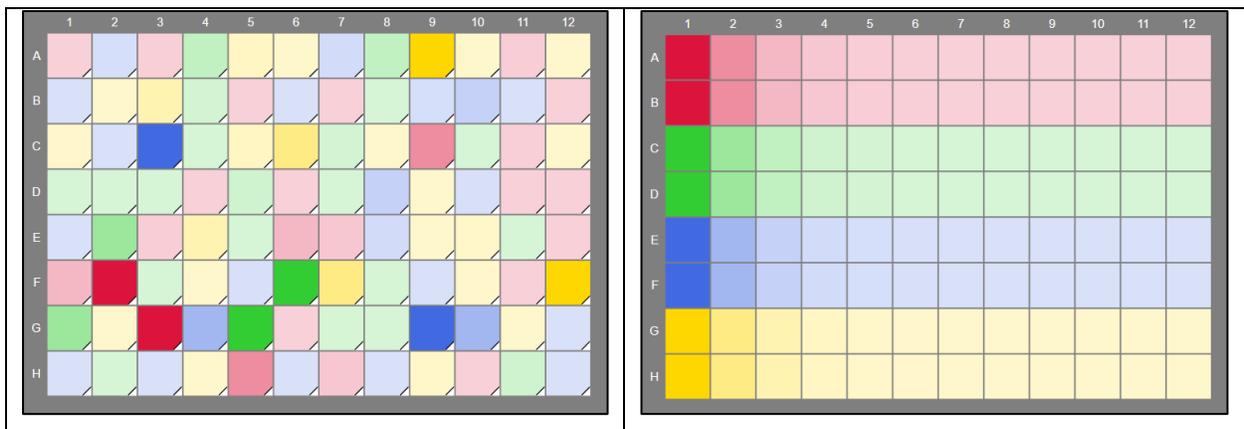


Figure 13 Original Non-Randomized Protocol [left]; Randomized Protocol [right]

NOTE: This example assumes that empty wells are included in the “Tabular Detail” tab of the report (see [Reports](#)). Alternately, the “Tabular” report tab could be used.



Report options

- Include "Don't Dispense" fluid tables
- Include empty wells in "Tabular detail"

Step 2: Read results on plate reader

Figure 14 shows typical results from a plate reader. When a plate reader has the option, it is preferable to use columnar output format, in other words with data sorted A1-H12 (96 plate) or A1-P24 (384 plate) in a single column. Save the plate reader data to an Excel or CSV file for use in Step 4.

```

File Edit Format View Help
Basic assay information
Assay ID: 3229
Assay Started: 12/15/2009 13:29
Assay Finished: 12/15/2009 13:33
Assay Exported: 12/15/2009 13:33
Protocol ID: 100246
Protocol Name: Crystal violet
Serial#: 1020011

Plate well P1tRpt=1 Group=1 Crystal violet
CalcResult
1 A01 0.387
1 A02 0.39
1 A03 0.573
1 A04 0.498
1 A05 0.328
1 A06 0.322
1 A07 0.373
1 A08 0.34
1 A09 0.341
1 A10 0.392
1 A11 0.38
1 A12 1.015
1 B01 0.382
1 B02 0.36
1 B03 0.357
1 B04 0.325
1 B05 0.335
1 B06 0.374
1 B07 0.366
1 B08 0.43
1 B09 0.334
1 B10 0.356
    
```

Figure 14 Typical plate reader output in columnar format

Step 3: Open the dispenser *.xml report file in Excel

Once the *.xml report file has been opened, go to the “Tabular Detail” tab to find the dispense data arranged with each dispense assigned to a row. Note that rows are ordered A1 -H12 (96 plate) or A1-P24 (384 plate) and that in the example below, the dispensed volumes are found in Column ‘P’, while the corresponding concentrations are found in Column ‘N’.

Plate ID	Plate	Well	Dispensed Row	Dispensed Col	Non-random Well	Non-random Row	Non-random Col	Fluid	Fluid Name	Cassette	Dispense Head	Start Time	Dispensed Concentration	Units	Dispensed Volume	Units	Total Well Volume
1	A01	1	1	A05	1	5	3	Fluid 1	1	1	08:59:02.8	0.02799922	µM	0.28	nL	10000.28	
1	A02	1	2	F05	6	5	3	Fluid 3	3	1	08:59:08.4	1.679717807	µM	16.8	nL	100016.8	
1	A03	1	3	A06	1	6	1	Fluid 1	1	1	08:59:02.8	0.599964002	µM	6	nL	100006	
1	A04	1	4	D03	4	3	2	Fluid 2	2	1	08:59:06.0	12.91928761	µM	129.36	nL	100129.36	
1	A05	1	5	G04	7	4	4	Fluid 4	4	1	08:59:11.1	4.637848039	µM	46.4	nL	100046.4	
1	A06	1	6	G08	7	8	4	Fluid 4	4	1	08:59:11.1	0.07799392	µM	0.78	nL	10000.78	
1	A07	1	7	F04	6	4	3	Fluid 3	3	1	08:59:08.4	4.637848039	µM	46.4	nL	100046.4	
1	A08	1	8	C03	3	3	2	Fluid 2	2	1	08:59:06.0	12.91928761	µM	129.36	nL	100129.36	
1	A09	1	9	G01	7	1	4	Fluid 4	4	1	08:59:11.0	99.99099999	µM	1010	nL	101010	
1	A10	1	10	G11	7	11	4	Fluid 4	4	1	08:59:11.1	0.003999998	µM	0.04	nL	10000.04	
1	A11	1	11	A05	1	5	1	Fluid 1	1	1	08:59:02.8	1.679717807	µM	16.8	nL	100016.8	
1	A12	1	12	G07	7	7	4	Fluid 4	4	1	08:59:11.1	0.21599335	µM	2.16	nL	10002.16	
1	B01	2	1	E09	5	9	3	Fluid 3	3	1	08:59:08.5	0.02799922	µM	0.28	nL	10000.28	
1	B02	2	2	H10	8	10	4	Fluid 4	4	1	08:59:11.1	0.00999999	µM	0.1	nL	10000.1	
1	B03	2	3	G03	7	3	4	Fluid 4	4	1	08:59:11.1	12.94322558	µM	129.6	nL	100129.6	
1	B04	2	4	D05	4	5	2	Fluid 2	2	1	08:59:06.0	1.679717807	µM	16.8	nL	100016.8	
1	B05	2	5	B09	2	9	1	Fluid 1	1	1	08:59:02.8	0.02799922	µM	0.28	nL	10000.28	
1	B06	2	6	I07	5	7	3	Fluid 3	3	1	08:59:08.5	0.21599335	µM	2.16	nL	10002.16	
1	B07	2	7	B11	2	11	1	Fluid 1	1	1	08:59:02.8	0.003999998	µM	0.04	nL	10000.04	
1	B08	2	8	C10	3	10	2	Fluid 2	2	1	08:59:06.0	0.00899999	µM	0.099	nL	10000.099	
1	B09	2	9	E05	5	5	3	Fluid 3	3	1	08:59:08.4	1.679717807	µM	16.8	nL	100016.8	
1	B10	2	10	F03	5	3	3	Fluid 3	3	1	08:59:08.4	12.94322558	µM	129.6	nL	100129.6	
1	B11	2	11	F09	6	9	3	Fluid 3	3	1	08:59:08.4	0.02799922	µM	0.28	nL	10000.28	
1	B12	2	12	B12	2	12	1	Fluid 1	1	1	08:59:02.8	0.0013	µM	0.013	nL	10000.013	
1	C01	3	1	H06	8	6	4	Fluid 4	4	1	08:59:11.1	0.599964002	µM	6	nL	100006	
1	C02	3	2	E12	5	12	3	Fluid 3	3	1	08:59:08.5	0.0013	µM	0.013	nL	10000.013	
1	C03	3	3	F01	6	1	3	Fluid 3	3	1	08:59:08.4	99.99099999	µM	1010	nL	101010	
1	C04	3	4	D11	4	11	2	Fluid 2	2	1	08:59:06.0	0.003299999	µM	0.033	nL	10000.033	
1	C05	3	5	H04	8	4	4	Fluid 4	4	1	08:59:11.2	4.637848039	µM	46.4	nL	100046.4	
1	C06	3	6	H02	8	2	4	Fluid 4	4	1	08:59:11.2	35.5029135	µM	360.8	nL	100360.8	
1	C07	3	7	C05	3	5	2	Fluid 2	2	1	08:59:06.0	1.679717807	µM	16.8	nL	100016.8	
1	C08	3	8	G10	7	10	4	Fluid 4	4	1	08:59:11.2	0.00999999	µM	0.1	nL	10000.1	
1	C09	3	9	A02	1	2	1	Fluid 1	1	1	08:59:02.9	35.5029135	µM	360.8	nL	100360.8	
1	C10	3	10	C08	3	8	2	Fluid 2	2	1	08:59:06.0	0.07809939	µM	0.781	nL	10000.781	
1	C11	3	11	B06	2	6	1	Fluid 1	1	1	08:59:02.9	0.599964002	µM	6	nL	100006	

Figure 15 Dispense data Tabular-Detail Tab

Step 4: Paste reader data into the *.xml file.

Paste the plate reader data into the “Tabular Detail” tab of the *.xml dispenser report file. Because both data sets are ordered A1-H12, the plate reader wells are now matched with the report file wells. The pasted wells are highlighted in yellow for visibility.

Figure 16 Dispense data with plate reader data pasted in

Step 5: Re-sort data by fluid name and then “Dispensed concentration”

To get the data sorted by fluid name and then in order of concentration, the Excel ‘data sort’ function can be employed, as shown in [Figure 17](#). The resultant data after sorting can be seen in [Figure 18](#).

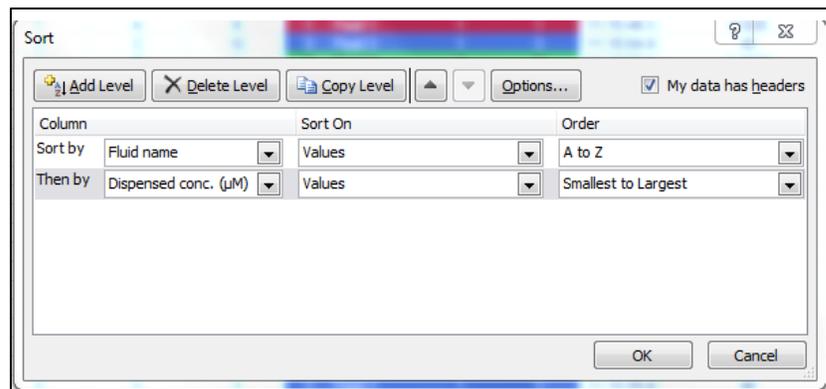


Figure 17 Data sort function in Excel to get randomized data into a useful sequence

Plate ID	Plate	Dispensed Well	Dispensed Row	Dispensed Col	Non-random Well	Non-random Row	Non-random Col	Fluid	Fluid Name	Cassette	Dispense Head	Start Time	Dispensed Concentration	Units	Dispensed Volume	Units	Total Well Volume	Plate	Well	Cal/Result
1	E12	5	12	A12	2	12	1	Fluid 1	1	1	08:59:02.6	0.0013	µM	0.013	nL	100000.013	1	E12	44800.33	
2	B07	2	7	B11	2	11	1	Fluid 1	1	1	08:59:02.8	0.003999998	µM	0.04	nL	100000.04	1	B07	45043.33	
3	G06	7	6	A11	1	11	1	Fluid 1	1	1	08:59:03.0	0.003999998	µM	0.04	nL	100000.04	1	G06	45393.33	
4	D06	4	6	A10	1	10	1	Fluid 1	1	1	08:59:02.9	0.009999999	µM	0.1	nL	100000.1	1	D06	45601.33	
5	D11	4	11	B10	2	10	1	Fluid 1	1	1	08:59:02.9	0.009999999	µM	0.1	nL	100000.1	1	D11	45883.33	
6	A01	1	1	A09	1	9	1	Fluid 1	1	1	08:59:02.8	0.027999922	µM	0.28	nL	100000.28	1	A01	431461.33	
7	B05	2	5	B09	2	9	1	Fluid 1	1	1	08:59:02.8	0.027999922	µM	0.28	nL	100000.28	1	B05	450799.33	
8	D04	4	4	B08	2	8	1	Fluid 1	1	1	08:59:02.9	0.077999592	µM	0.78	nL	100000.78	1	D04	448603.33	
9	F11	6	11	A08	1	8	1	Fluid 1	1	1	08:59:03.0	0.077999592	µM	0.78	nL	100000.78	1	F11	464139.33	
10	D12	4	12	A07	1	7	1	Fluid 1	1	1	08:59:02.9	0.215995335	µM	2.16	nL	100002.16	1	D12	454708.33	
11	H10	8	10	B07	2	7	1	Fluid 1	1	1	08:59:03.0	0.215995335	µM	2.16	nL	100002.16	1	H10	465777.53	
12	A03	1	3	A06	1	6	1	Fluid 1	1	1	08:59:02.8	0.599964002	µM	6	nL	100000.6	1	A03	431641.33	
13	C11	3	11	B06	2	6	1	Fluid 1	1	1	08:59:02.9	0.599964002	µM	6	nL	100000.6	1	C11	459121.33	
14	A11	1	11	A05	1	5	1	Fluid 1	1	1	08:59:02.8	1.679717807	µM	16.8	nL	100016.8	1	A11	459621.33	
15	E03	5	3	B05	2	5	1	Fluid 1	1	1	08:59:02.9	1.679717807	µM	16.8	nL	100016.8	1	E03	467833.33	
16	E07	5	7	B04	2	4	1	Fluid 1	1	1	08:59:02.9	4.637848039	µM	46.4	nL	100046.4	1	E07	46372.33	
17	H07	8	7	A04	1	4	1	Fluid 1	1	1	08:59:03.0	4.637848039	µM	46.4	nL	100046.4	1	H07	465051.22	
18	E06	5	6	A03	1	3	1	Fluid 1	1	1	08:59:02.9	12.94322558	µM	129.6	nL	100129.6	1	E06	460259.33	
19	F01	6	1	B03	2	3	1	Fluid 1	1	1	08:59:03.0	12.94322558	µM	129.6	nL	100129.6	1	F01	456648.33	
20	C09	3	9	A02	1	2	1	Fluid 1	1	1	08:59:02.9	35.95029135	µM	360.8	nL	100360.8	1	C09	460540.33	
21	H05	8	5	B02	2	2	1	Fluid 1	1	1	08:59:03.0	35.95029135	µM	360.8	nL	100360.8	1	H05	464567.02	
22	F02	6	2	B01	2	1	1	Fluid 1	1	1	08:59:02.8	99.99009999	µM	1010	nL	101010	1	F02	444070.33	
23	G03	7	3	A01	1	1	1	Fluid 1	1	1	08:59:02.8	99.99009999	µM	1010	nL	101010	1	G03	463280.33	
24	D07	4	7	C12	3	12	2	Fluid 2	2	1	08:59:06.0	0.0011	µM	0.011	nL	100000.011	1	D07	458679.33	
25	E11	5	11	D12	4	12	2	Fluid 2	2	1	08:59:06.1	0.0011	µM	0.011	nL	100000.011	1	E11	467184.33	
26	C04	3	4	D11	4	11	2	Fluid 2	2	1	08:59:06.0	0.003999999	µM	0.039	nL	100000.039	1	C04	458665.33	
27	D08	4	3	C11	3	11	2	Fluid 2	2	1	08:59:06.1	0.003999999	µM	0.039	nL	100000.039	1	D08	466393.33	
28	B08	2	8	C10	3	10	2	Fluid 2	2	1	08:59:06.0	0.009899999	µM	0.099	nL	100000.099	1	B08	441153.33	
29	H02	8	2	D10	4	10	2	Fluid 2	2	1	08:59:06.2	0.009899999	µM	0.099	nL	100000.099	1	H02	463840.71	
30	D01	4	1	C09	3	9	2	Fluid 2	2	1	08:59:06.1	0.027499924	µM	0.275	nL	100000.275	1	D01	460926.33	
31	E05	5	5	D09	4	9	2	Fluid 2	2	1	08:59:06.1	0.027499924	µM	0.275	nL	100000.275	1	E05	466666.33	
32	C10	3	10	C08	3	8	2	Fluid 2	2	1	08:59:06.0	0.078099999	µM	0.781	nL	100000.781	1	C10	450731.33	
33	G08	7	8	D08	4	8	2	Fluid 2	2	1	08:59:06.2	0.078099999	µM	0.781	nL	100000.781	1	G08	456442.33	
34	D02	4	2	C07	3	7	2	Fluid 2	2	1	08:59:06.1	0.216695304	µM	2.167	nL	100002.167	1	D02	454941.33	
35	F03	6	3	D07	4	7	2	Fluid 2	2	1	08:59:06.1	0.216695304	µM	2.167	nL	100002.167	1	F03	460778.33	
36	H08	8	8	C06	3	6	2	Fluid 2	2	1	08:59:06.1	0.615962057	µM	6.16	nL	100006.16	1	H08	459622.33	
37	G07	7	7	D06	4	6	2	Fluid 2	2	1	08:59:06.2	0.615962057	µM	6.16	nL	100006.16	1	G07	458168.33	
38	B04	2	4	D05	4	5	2	Fluid 2	2	1	08:59:06.0	1.679717807	µM	16.8	nL	100016.8	1	B04	444445.33	

Figure 18 Data set now sorted by Fluid name and then by dispensed concentration

Getting plate reader data into columnar format

If plate reader data is output in spatial plate-layout format, rather than columnar format, several templates are available to enable the conversion to columnar format:

- Spatial Plate to Columnar Template 96: For analyzing 96-well plates.
- Spatial Plate to Columnar Template 384: For analyzing 384-well plates.
- Spatial Plate to Columnar Template 1536: For analyzing 1536-well plates.

These templates are found in the HP Dispensing folder within “My Documents.” Each template has an “Input SpatialFormat **” tab, and an “Output ColumnarFormat **” tab. Reader data in spatial plate-layout format can be pasted into the Input tab, and then the same plate data in columnar format can be copied out of the output tab. [Figure 19](#) shows an example of a 384-plate template for inputting spatial plate-layout data. [Figure 20](#) shows an example of the same data in columnar output format. Once data is arranged in columnar format, it can then be matched to the dispenser report file as shown below.

1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	
1	1.151	1.163	1.194	1.2	1.203	1.232	1.246	1.289	1.283	1.254	1.26	1.259	1.253	1.25	1.248	1.26	1.26	1.252	1.226	1.243	1.238	1.242	1.247	
2	1.281	1.259	1.251	1.259	1.254	1.248	1.245	1.253	1.258	1.256	1.259	1.256	1.267	1.263	1.256	1.254	1.266	1.262	1.241	1.224	1.21	1.213	1.208	1.238
3	1.261	1.261	1.248	1.263	1.261	1.256	1.275	1.277	1.274	1.283	1.281	1.289	1.282	1.281	1.253	1.263	1.255	1.261	1.241	1.22	1.228	1.229	1.239	1.255
4	1.261	1.264	1.272	1.27	1.255	1.264	1.253	1.278	1.265	1.27	1.254	1.269	1.27	1.277	1.263	1.263	1.269	1.282	1.237	1.24	1.26	1.26	1.254	1.254
5	1.273	1.25	1.259	1.251	1.25	1.263	1.264	1.264	1.288	1.254	1.26	1.27	1.271	1.259	1.262	1.261	1.264	1.222	1.214	1.201	1.219	1.22	1.228	
6	1.236	1.228	1.234	1.229	1.235	1.225	1.249	1.245	1.229	1.248	1.229	1.229	1.238	1.247	1.248	1.249	1.25	1.252	1.21	1.203	1.218	1.21	1.204	1.224
7	1.23	1.23	1.212	1.219	1.2	1.22	1.211	1.207	1.227	1.215	1.225	1.232	1.221	1.231	1.223	1.218	1.224	1.206	1.183	1.173	1.166	1.171	1.167	1.177
8	1.205	1.184	1.19	1.192	1.19	1.191	1.197	1.206	1.193	1.189	1.192	1.202	1.195	1.201	1.197	1.2	1.203	1.209	1.168	1.152	1.178	1.162	1.161	1.177
9	1.195	1.197	1.181	1.191	1.186	1.193	1.187	1.186	1.205	1.194	1.201	1.199	1.201	1.194	1.188	1.195	1.202	1.203	1.176	1.164	1.16	1.152	1.161	1.184
10	1.184	1.184	1.179	1.189	1.188	1.199	1.197	1.201	1.208	1.215	1.205	1.212	1.207	1.203	1.192	1.197	1.214	1.177	1.145	1.146	1.137	1.128	1.139	1.148
11	1.181	1.176	1.18	1.176	1.184	1.185	1.183	1.196	1.2	1.197	1.195	1.193	1.218	1.198	1.202	1.194	1.198	1.202	1.177	1.16	1.163	1.152	1.157	1.172
12	1.189	1.194	1.197	1.189	1.199	1.206	1.206	1.197	1.211	1.201	1.202	1.215	1.211	1.208	1.2	1.208	1.205	1.207	1.182	1.172	1.169	1.169	1.164	1.161
13	1.192	1.191	1.184	1.182	1.187	1.195	1.203	1.214	1.2	1.203	1.202	1.203	1.21	1.221	1.194	1.188	1.204	1.194	1.185	1.187	1.195	1.197	1.163	1.182
14	1.192	1.182	1.192	1.191	1.183	1.191	1.212	1.2	1.204	1.206	1.207	1.213	1.214	1.2	1.194	1.205	1.201	1.214	1.182	1.17	1.173	1.167	1.167	1.177
15	1.195	1.2	1.211	1.217	1.212	1.205	1.204	1.215	1.215	1.21	1.218	1.222	1.219	1.224	1.204	1.197	1.208	1.217	1.194	1.186	1.18	1.177	1.173	1.191
16	1.248	1.236	1.236	1.236	1.234	1.232	1.233	1.239	1.226	1.234	1.232	1.24	1.241	1.227	1.233	1.242	1.238	1.234	1.208	1.208	1.2	1.194	1.209	1.188

Paste Plate Reader Data

Figure 19 Sample data input in spatial-plate format

	A	B
1	Well	Response Output
2	A1	1.151
3	A2	1.163
4	A3	1.194
5	A4	1.2
6	A5	1.203
7	A6	1.232
8	A7	1.246
9	A8	1.269
10	A9	1.263
11	A10	1.254
12	A11	1.26
13	A12	1.259
14	A13	1.253
15	A14	1.25
16	A15	1.248
17	A16	1.26
18	A17	1.26
19	A18	1.252
20	A19	1.226
21	A20	1.243
22	A21	1.243
23	A22	1.238
24	A23	1.242
25	A24	1.247
26	B1	1.281
27	B2	1.259
28	B3	1.251

Figure 20 Sample data output in columnar format

Alternative methods for sorting

Note in [Figure 18](#) that several other columns may sometimes be useful as an alternative method for sorting dispensing data after it has been paired to plate-reader data. Specifically, Columns F (Non-Random well), Column G (Non-Random Row), and/or Column H (Non-Random Column) can be used to rearrange randomized plate dispense and plate reader data back into the original non-randomized layout.

Appendix D—Concentration calculations

With DMSO and dilute aqueous solutions, volume to achieve a desired concentration may be calculated without including dispensed volumes; this is a negligible error since the added volumes from the dispenser are small relative to the total assay volume. However, there are cases where the entire assay may be dispensed by the dispenser. For example, in miniaturized well plate based PCR, the volumes added by the dispenser are usually significant and should not be ignored in concentration calculations.

Each protocol plate has a specified **additional volume**. This additional volume is the amount of fluid that the user places in the well in addition to dispensed fluids. The total assay volume will be this **additional volume** plus volume dispensed by the dispenser. In some cases – such as miniaturized PCR where master mix, primer and sample are all from the dispenser – the additional volume for the plate should be set to zero.

Calculation of true concentration must be accomplished iteratively since adding fluid with the dispenser changes the overall assay volume which in turn changes concentrations. Calculation of concentration utilizes well volumes that include both dispensed fluids and user-added Additional volume (see [Plate-associated information](#)).



NOTE: Prior to software release 3.2, concentrations were calculated without including the volume dispensed using the dispenser. This change in concentration calculation means that reported concentrations from earlier versions of the software may be slightly different than those currently reported. This difference is generally negligible relative to other errors in the assay workflow.

Since concentrations are calculated iteratively, there can be conditions created where concentrations cannot be calculated. For example, it is possible to [Normalize](#) wells using **Normalize to % of total volume** with a percentage high enough where a solution does not exist as concentrations are iteratively recalculated. If this occurs, a warning is displayed for affected wells.

Appendix E—Dispense specifications

DMSO dispensing

Selected dispense head cassette specifications for DMSO dispensing:

- 70-100% DMSO.
- Nanoparticles < 1 micron in diameter in stable suspension at concentrations ≤ 0.5%.
- There are additional considerations for small molecule and biomolecule dispensing. See [Dispensing considerations](#) for more information.
- Only use non-Surfactant-Free cassettes for DMSO dispensing.

	Standard-volume cassette	High-volume cassette
Maximum dispense volume per dispense head	10 µL	200 µL
Maximum dispense volume per dispense head – normalization and bulk dispense ¹	20 µL	200 µL
Minimum dispense volume	13 pL	1 nL
Minimum dispense volume – normalization and bulk dispense ¹	1 nL	1 nL
Minimum fill volume	2 µL	50 µL
Maximum dispense volume – each fluid per well	10 µL	10 µL
Dispense increment (per well)	400 pL for > 3.6 nL 20 pL for 17 pL - 3.6 nL 13 pL for < 17 pL	400 pL for > 3.6 nL 20 pL for 1 - 3.6 nL n/a for < 1 nL
Protocol limits (per fluid)	32 plates	32 plates

Notes:

1 – Bulk Dispense is defined as when all dispensed volumes for a fluid are 1nL or greater.

Acetonitrile (ACN) dispensing

Selected dispense head cassette specifications for ACN dispensing:

- Dispensing of ACN is limited to one 384-well plate per dispense head due to its rapid evaporation rate.
- Recommended use of standard volume cassettes due to the rapid evaporation rate of ACN.
- Dispense volume is limited to >10 nL per well and <40 µL per fluid per protocol.
- For combined use with TMTs a maximum concentration of 85 mM TMT is recommended.
- There are additional considerations for small molecule and biomolecule dispensing. See [Dispensing considerations](#) for more information.
- Only use non-Surfactant-Free cassettes for ACN dispensing.

Aqueous dispensing

Selected dispense head cassette specifications for aqueous dispensing, including surfactant and glycerol:

- Aqueous solutions require the addition of surfactant to enable priming by reducing surface tension unless using Surfactant-Free cassettes. Refer to [Table 1](#) for recommended surfactants and concentration ranges. In addition, the presence or absence of glycerol impacts dispensing – there are separate fluid classes dependent on glycerol content.
- Master mix fluids ([Table 2](#)) do not require addition of surfactants and should not be dispensed with Surfactant-Free (SF) cassettes.
- Aqueous compounds can be dispensed without the addition of surfactant using Surfactant-Free (SF) cassettes.
- Total buffer components < 150 mM.
- Nanoparticles < 1 micron in diameter in stable suspension at concentrations ≤ 0.5% by weight.
- Types of biomolecules which either can or cannot be dispensed are listed in [Table 3](#).
- There are additional considerations for small molecule and biomolecule dispensing. See [Dispensing considerations](#) for more information.

	Standard-volume cassette	High-volume cassette
Maximum dispense volume per dispense head	10 µL	200 µL
Maximum dispense volume per dispense head – normalization and bulk dispense ¹	20 µL	200 µL
Minimum dispense volume	11 pL	1 nL
Minimum dispense volume – normalization and bulk dispense ¹	1 nL	1 nL
Minimum fill volume	4 µL	50 µL
Maximum dispense volume – each fluid per well	10 µL	10 µL
Dispense increment (per well) without glycerol	560 pL for > 3.6 nL 11 pL for ≤ 3.6 nL	560 pL for > 3.6 nL 11 pL for 1 - 3.6 nL n/a for < 1 nL
Dispense increment (per well) with glycerol	600 pL for > 3.6 nL 12 pL for ≤ 3.6 nL	600 pL for > 3.6 nL 12 pL for 1 - 3.6 nL n/a for < 1 nL
Protocol limits (per fluid)	1536 wells or 8 plates ^{2,3}	1536 wells or 8 plates ^{2,3}

Notes:

1 – Bulk Dispense is defined as when all dispensed volumes for a fluid are 1 nL or greater.

2 – To minimize evaporation.

3 – Except Aqueous fluids listed in Table 4 that have a protocol limit of 1 plate (384-well)

Fluid Class	Surfactant	Surfactant Concentration	Glycerol
Aqueous	None	N/A	0 - <5%
a + B	Brij 35	0.1 %	0 - <5%
a + B + g	Brij 35	0.1 %	5 - 20%
a + TX	Triton X100	0.1 %	0 - <5%
a + TX + g	Triton X100	0.1 %	5 - 20%
a + Tw	Tween 20	0.3%	0 - <5%
a + Tw + g	Tween 20	0.3%	5 - 20%

Table 1. Aqueous fluid classes

Master Mixes ¹
Brilliant II SYBR Green qRT-PCR 1-Step Master Mix Brilliant III Ultra-Fast SYBR Green qPCR Master Mix Brilliant Multiplex qPCR Master Mix Fast SYBR Green Master Mix iQ SYBR Green Supermix Power SYBR Green PCR Master Mix PowerUp SYBR Green Master Mix QuantiTect SYBR Green PCR Master Mix SsoAdvanced Universal SYBR Green Supermix SYBR Select Master Mix TaqMan Gene Expression Master Mix TaqMan RNA-to-CT 1-Step Kit TaqMan Universal PCR Master Mix

Table 2. Master Mixes

Notes:

1 – The system requires some preparation time to dispense a master mix to ensure optimal performance in every well. The length of this delay may vary from one master mix to another.

Compatible	Not Compatible
Enzymes < 300kDa Antibodies Peptides Oligos and labeled oligos Lipids Small molecules such as ATP, ADP PCR Master mix DNA < 10,000 base pairs	Enzymes > 300kDa Saliva Whole blood Plasma Serum

Table 3. Example compatible biomolecules

Chemical name	Maximum concentration	Recommended dispense volume (per well)	Recommended cassette	Protocol limits (per fluid)
Iodoacetamide (IAA)	53 mM	11 pl - 2.5 µL	High and standard volume SF	1 384-well plate
Tris(2-carboxyethyl)phosphine (TCEP)	100 mM	11 pl - 2.5 µL	High and standard volume SF	1 384-well plate
Dithiothreitol (DTT)	50 mM	11 pl - 2.5 µL	High and standard volume SF	1 384-well plate
Trypsin	0.1 mg/mL	11 pl - 2.5 µL	High and standard volume SF	1 384-well plate
Triethylammonium bicarbonate (TEAB)	100 mM	11 pl - 2.5 µL	High and standard volume SF	1 384-well plate
Trifluoroacetic acid (TFA) ¹	10% aq	10 nL - 2.5 µL ¹	High and standard volume SF	1 384-well plate
Hydroxylamine	5% aq	11 pl - 2.5 µL	High and standard volume SF	1 384-well plate
Formic Acid	4% aq	11 pl - 2.5 µL	High and standard volume SF	1 384-well plate

Table 4. Proteomics fluids qualified for aqueous dispensing using Surfactant-Free (SF) cassettes

Notes:

1 –Recommended TFA dispense volume range between 10 nL and 2.5 µL based on internal testing

Single cell dispensing

Selected dispense head cassette specifications for single cell dispensing:

- Cells suspended in 1X PBS buffer at 200 cells/ μL .
- Cell size range 10 – 25 μm .
- There are additional considerations for cell dispensing. See [Appendix F—Best practices for dispensing - Single cell dispensing](#) for more detail.

	Single cell cassette
Maximum dispense volume per dispense head	20 μL
Minimum single cell isolation volume	480 pL
Minimum fill volume	4 μL
Drop volume	40 pL
Protocol cassette limit	384 wells or 1 plate

Fluid group	Fluid class	Recommended cell size range
Cells	Small	10 – 14 μm
Cells	Medium	15 – 17 μm
Cells	Large	18 – 20 μm
Cells	Extra Large	21 – 25 μm
Sticky Cells	Small	10 – 14 μm
Sticky Cells	Medium	15 – 17 μm
Sticky Cells	Large	18 – 20 μm
Sticky Cells	Extra Large	21– 25 μm

Table 5. Cell fluid groups and classes

Fluid group selection

- **Cells:** this fluid group refers to most standard cell lines.
- **Sticky cells:** this fluid group refers to cell lines that tend to aggregate after trypsinization in suspension. Sticky cells require a larger clearing volume to travel through the microfluidics and to be dispensed into the target well.

Fluid class selection

- The cell size is the diameter of the rounded cells in suspension.

Compatible cell types:

Compatible (10–25 µm cell sizes)	Not compatible
Adherent cell lines Suspension cell lines Disaggregated cell constructs Disaggregated organoids Primary cells	Bacteria Cell organelles (e.g. nuclei) Yeast Virus

Dispensing considerations

The dispensing software will use one of several modes to dispense your fluids. If you have high-volume cassettes available, there are three potential modes.

- Use high-volume cassettes with a maximum dispense volume of 200 µL.
- Use standard-volume cassettes with a maximum dispense volume of 20 µL.
- Use standard-volume cassettes with a maximum dispense volume of 10 µL.

In deciding which mode will be used, the first consideration is the lowest volume to be dispensed into any well using that fluid. If any well has less than one nL, then standard-volume cassettes dispensing 10 µL will be used to ensure good reproducibility with low volume dispensing. If all wells for a fluid are greater than one nL, then [Table 6](#) should be used. The more concentrated the fluid, the less total volume can be reliably dispensed. If the DMSO fluid class is used, or if the biomolecule concentration is not entered in mg/mL, then the most economical option is used by the dispenser.

	Small Molecules	Biomolecules
Standard-volume 10 µL	1000 Da 10 mM	300 kDa or 10,000 base pairs 3 mg/mL
Standard-volume 20 µL	1000 Da 10 mM	300 kDa or 10,000 base pairs 1 mg/mL
High-volume 200 µL	1000 Da 10 mM	300 kDa or 10,000 base pairs 0.3 mg/mL

Table 6. Dispense limits for small molecules and biomolecules

One final consideration is that high-volume cassettes have a greater dead volume than standard-volume cassettes due to the amount of stranded fluid in the large fill cup. If there is an extremely precious fluid, deselect “Use high-volume cassettes” at the start of the protocol run.

Appendix F—Best practices for dispensing

All fluids

To ensure optimum performance, it is recommended that the first plate using a fluid includes a well with volume of 30 nL or greater; this ensures the dispense head is fully ready to dispense other much smaller volumes. Since plate dispense order can vary, it is recommended that all plates follow this recommendation. If your assay design cannot tolerate a dispense volume greater than 30nL, consider dedicating one “scrap” well on each plate to a larger dispense volume. This same well can be used for all fluids on the plate.

DMSO based fluids

- Although very large numbers of fluids and plates are allowed, it is recommended that designs of fewer than 100 fluids and 32 plates are used to reduce user error and to speed up the display processes of the software.
- DMSO fluids with only small molecules less than 1000 Daltons at concentrations no more than 10mM are recommended to be dispensed with the dispenser.
- If prefilling the dispense heads with fluids outside of the dispenser, for example using a robotic system, minimize the time between filling and dispensing. It is recommended that prefilled dispense heads with DMSO based fluids be used within 30 minutes of filling.
- During the Run Protocol operation, it is recommended that the dispenser not be paused for longer than 15 minutes prior to resuming while using DMSO based fluids.
- Only use non-Surfactant-Free cassettes for DMSO dispensing.

Acetonitrile (ACN) based fluids

- ACN fluids with only small molecules less than 1000 Daltons at concentrations no more than 10mM are recommended to be dispensed (except for TMTs, tested to 85 mM)
- Do not prefill the dispense heads with ACN fluids due to evaporation concerns
- During the Run Protocol operation, it is recommended that the dispenser not be paused while using ACN based fluids.
- Shaking after each dispense head is not recommended for ACN fluids.
- Only use non-Surfactant-Free cassettes for ACN dispensing.
- Due to the high evaporation rate of ACN, a limitation of one 384-well plate per dispense head is recommended.

Aqueous based fluids

- The recommended limits for aqueous based fluids are shown when the cursor is moved over the Fluid Class drop-down menu in the Fluid Edit dialog box. For all aqueous fluids it is recommended that:
 - Small molecule compounds be less than 1000 Daltons and be present at no more than 10mM concentration, except for selected Proteomics reagents (see [Appendix E—Dispense specifications, Table 4](#)).
 - Biomolecules be less than 300kD and be present in total at no more than 3mg/mL concentration.
 - Glycerol levels be within the fluid class guidelines of either 0 - <5% or 5 - 20%.
- Surfactant-Free (SF) cassettes should be stored at 4°C to maintain 6-month shelf life.

- The best practices noted below are due to the high evaporation rate of aqueous fluids relative to DMSO:
 - The maximum number of plates in an aqueous fluid protocol is 1536 wells or 8 plates (or 1 plate for proteomics fluids, see [Appendix E—Dispense specifications, Table 4](#)).
 - The amount of aqueous fluid that the user is requested to add to each dispense head is greater than that requested for DMSO fluids, even when the amount to be dispensed is the same. In addition, the minimum volume requested for the fill is 4 μL for aqueous fluids as opposed to a 2 μL minimum requested volume for DMSO fluids. The dispensed volume for each fluid in the protocol is listed in the Fluids List, and the amount to fill into the dispense head is given during Run Protocol.
 - Fill dispense heads with only the amount of aqueous fluid requested at the time of use during Run Protocol. It is not recommended to overfill aqueous fluids in any dispense head.
 - Fill dispense heads with aqueous fluids at the time of use during Run Protocol. It is recommended that dispense heads not be pre-filled offline with aqueous based fluids.
 - During the Run Protocol operation, it is recommended that the dispenser not be paused while using aqueous based fluids.
 - Shaking after each dispense head is not recommended for aqueous fluids.

Single cell dispensing

- For all cell dispensing users must:
 - Use a C1 single cell cassette (limited to 384 wells each to ensure performance).

Recommended single cell sample preparation protocol

- Use cell culture at about 80% confluence pre-dispensing.
- Wash cells with 1X PBS three times before trypsinization.
- Trypsinize and neutralize with complete medium.
- Pipette cell suspension 10 times up and down.
- Take a 1 mL aliquot of the cell suspension and test cell concentration and viability.
- Centrifuge cell suspension at 200 rcf for five minutes.
- Discard supernatant and resuspend cell pellet in 1 mL 1X PBS.
- Filter the cell suspension with a 35 μm filter to form the **stock cell suspension**.
- Test cell concentration and viability.
- Begin a single cell protocol run on the D100 up to the point of inserting the C1 cassette.
- Dilute **stock cell suspension** sample with 1X PBS to 200 cells/ μL to form the **dispensing cell suspension**.
- Immediately load the fill cup of the single cell C1 cassette with the **dispensing cell suspension** without introducing bubbles and initiate dispensing run.

Other recommendations

- Keep **stock cell suspension** at room temperature.
- Prepare a fresh **dispensing cell suspension** if not used within 10 minutes.
- Measure cell concentration and viability of the **stock cell suspension** again to guarantee accurate dilution.
- Vortex (~ three seconds) before loading **cell suspension** into the single cell dispensing cassette.

- Vortex the **dispensing cell suspension** frequently (every three minutes) while not being used.
- For microscopy imaging purposes, pre-fill plates with low volume of fluid (15 µl recommended for 384-well plates) for optimal post-dispense analysis of accuracy.

Cell fluid group and class

Choose the cell fluid group and class that generate the most single cell wells and least non-single cell wells as confirmed by microscopy or other technique such as proteomics or qPCR.

- Choosing the wrong cell fluid group can lead to:
 - More non-single cell wells by selecting Sticky Cells when working with non-sticky cells
 - False positive single cell wells (non-single cell well reported as single cell well) when working with a sticky cell line and selecting the Cells fluid group.
- Choosing the wrong cell fluid class can lead to the following problems:
 - If fluid class is too small, cells will clog the cassette and cause a time out error. There may also be an increase in false positive single cell wells (more wells containing 0 cells will be reported as containing a single cell).
 - If fluid class is too large there may be an increase in false positive single cell wells (more wells containing multiple cells will be reported as containing a single cell).

Time-out

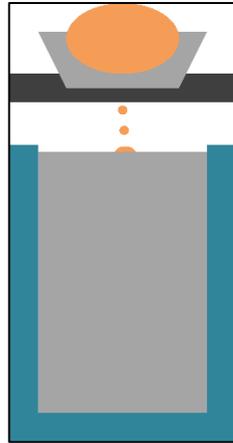
- **Poor single cell sample preparation:** prepare a fresh dispensing cell suspension (ensure thorough pipetting to disaggregate cell clumps), replace cassette and continue single cell dispensing. Stock cell suspension might need to be prepared de novo if error persists.
- **Wrong cell concentration:** prepare a fresh dispensing cell suspension by re-diluting the stock cell suspension to 200 cells/µL in 1X PBS.
- **Wrong fluid class selection:**
 - Test the fluid class that best matches the dispensed cell size according to the cell size range recommendations listed in [Appendix E—Dispense specifications, Table 4](#).
 - Choose the right fluid group depending on the dispensed cell properties. For each test, it is recommended to confirm single cell dispensing accuracy using microscopy or other technique to evaluate positive/false negatives.

C1 single cell cassettes

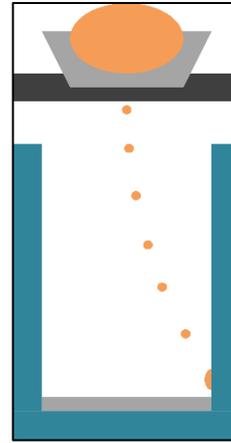
- Do not exceed cassette shelf-life specification: it is not recommended to use C1 cassettes beyond the expiration date.
- It is recommended to refrigerate the C1 cassettes at 4°C to ensure adequate shelf life.

Deep well plates

As distance between the dispense cassette and the target surface increases, dispense placement variation also increases. For this reason, it is best to minimize the dispense distance. When dispensing into deep well plates, it is recommended to dispense into pre-filled plates.



Best practice



Not optimal

Dispensing Software User Guide
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