



Medical-Biological
Research & Technologies



QUANT ASSAY SOFTWARE MANUAL FOR MPP-96 PHOTOMETER

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Purpose

This program is designed to operate the photometer MPP-96 and analyse the results obtained from it.

Using QuantAssay it is possible to program the analysis of the following assays:

- Quantitative assays: the ability to install up to 20 standards and choose fit model from 5/4 parameter logistic, linear and piecewise linear models
- BestFit function for the selection of the best calibration curve.
- Multiplex analysis - up to 7 different tests on the same plate
- Qualitative assays: the ability to install up to 8 types of controls (weak positive, strong positive, negative, etc.)
- Avidity / affinity assay
- Save, load and export results
- Create visual reports

This manual describes how to install the program, control the device, create and edit assays, analyse the results and troubleshoot the program.

Installation

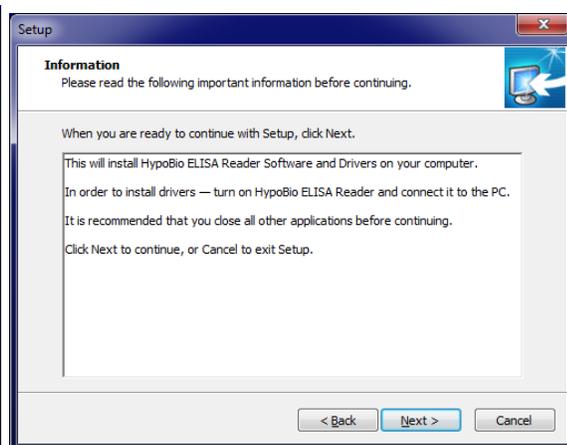
Welcome window



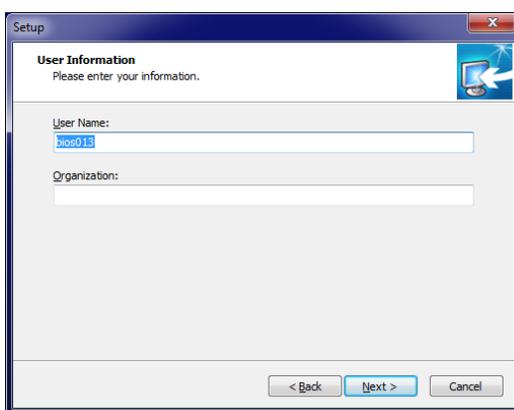
License Agreement



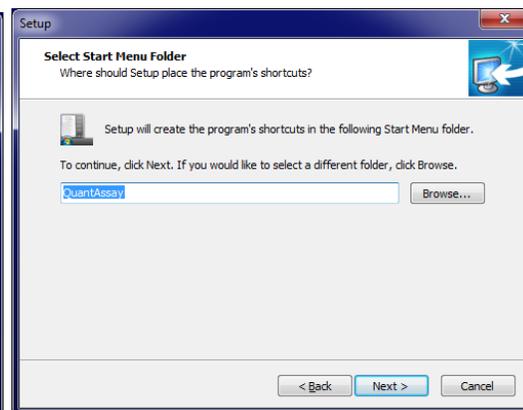
Information Window



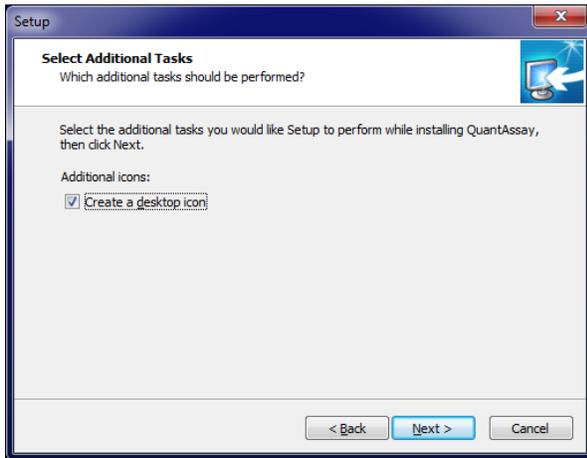
information of the user



the path of the installation



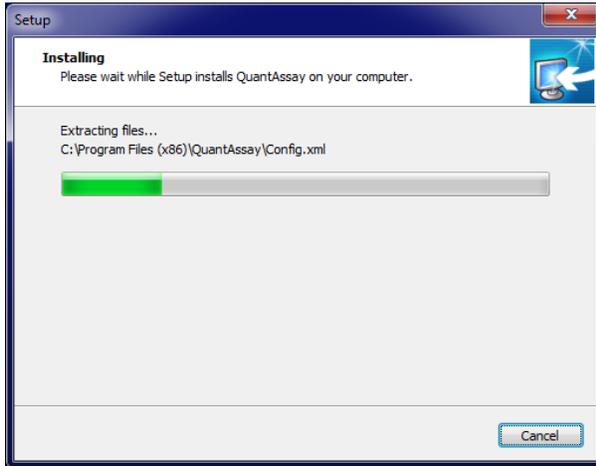
Advanced Settings



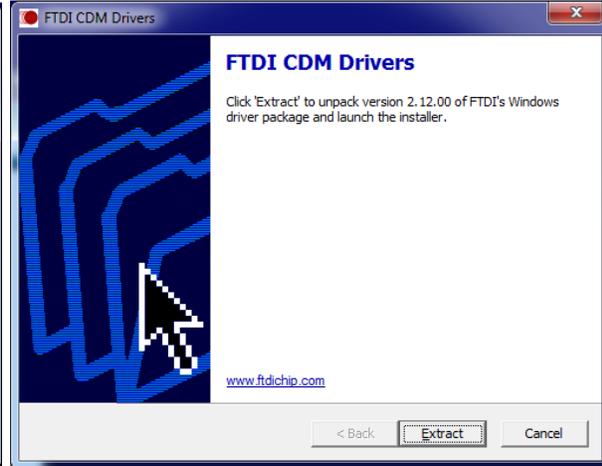
Ready to Install



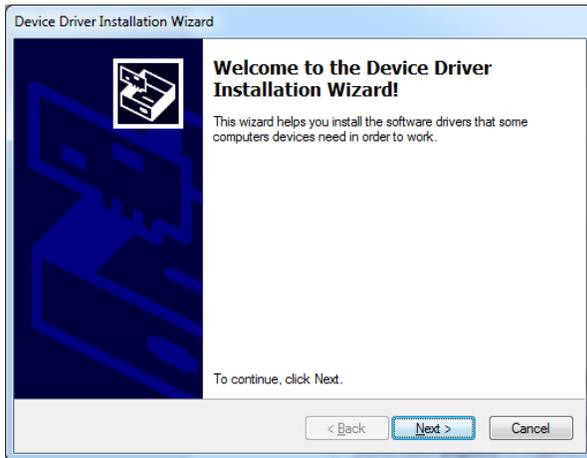
process software installation



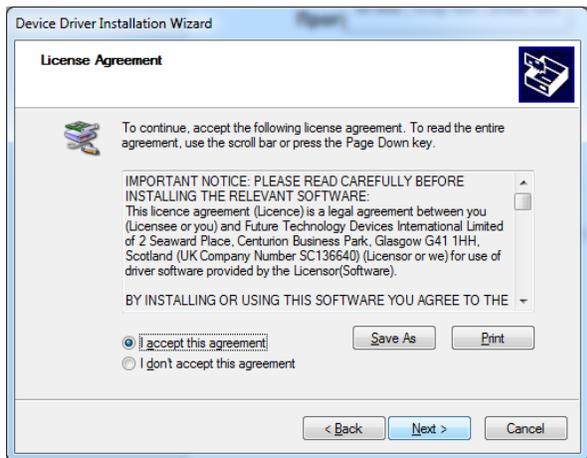
The installation of the drivers



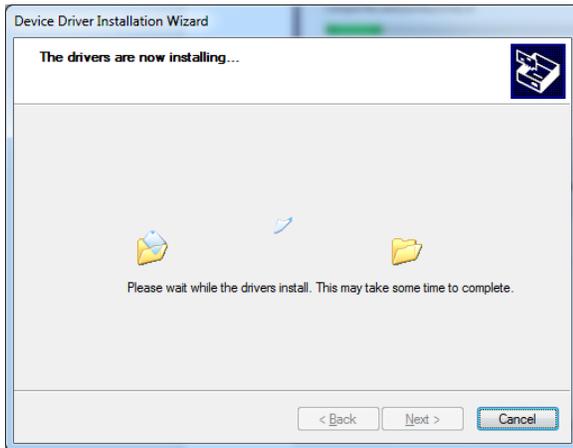
Drivers installation process



License Agreement



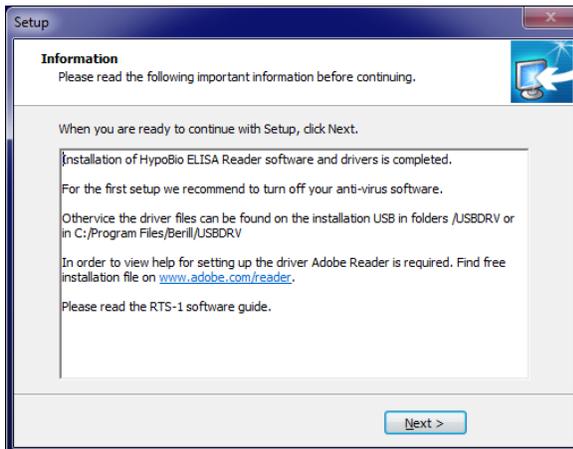
The installation of drivers



the installation of drivers is finished



finished installing of the software



The program is ready to start.



Account control: Administrators and common users rights

1. Administrator rights: now you are able to set to access levels for the ordinary users and master users.

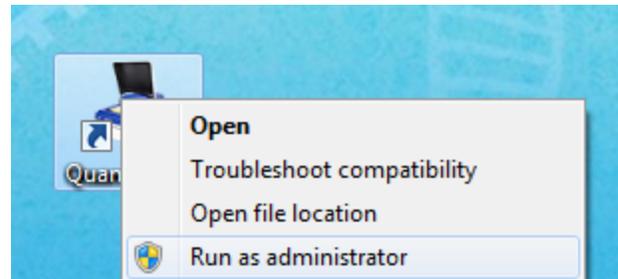
Ordinary users can:

- Use software
- Browse Assays
- Save templates

Master users can:

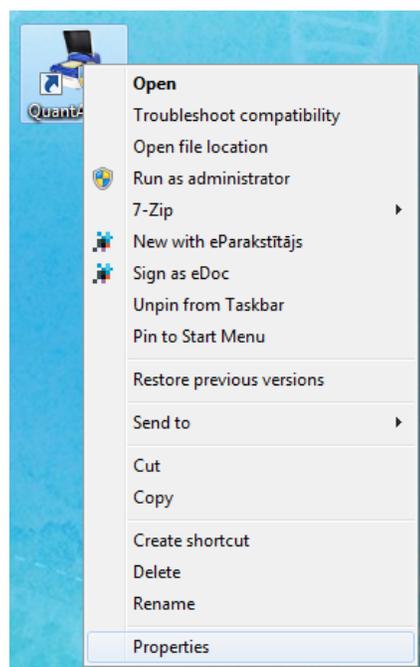
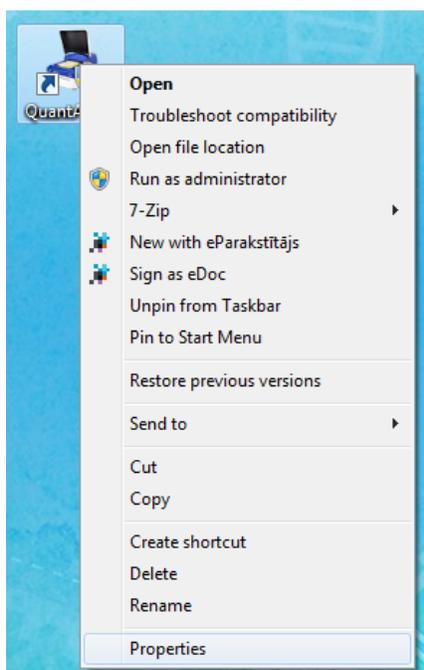
- Use software
- Create/Edit Assays
- Save templates

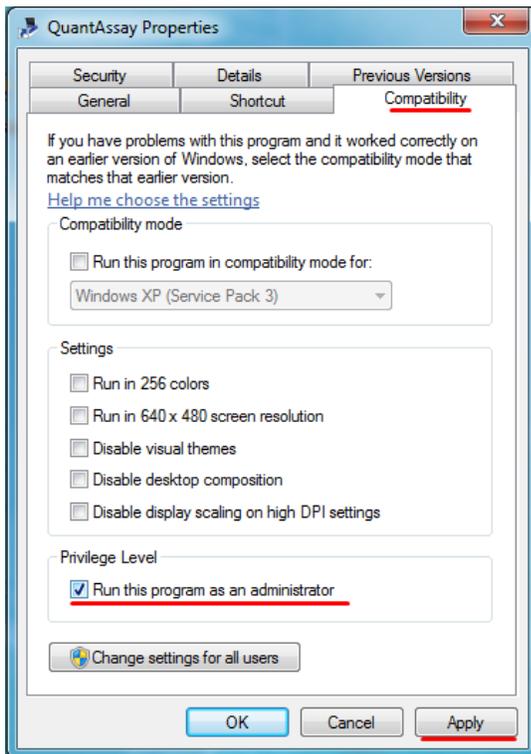
So, the ordinary users cannot change or create assays. If you have a single user who is also a master user than this feature can get annoying when creating or editing assays, so to use the software without being asked each time do the following: Run the software as administrator.



But user would need to do that every time he uses software.

Or if user wants to set this forever: Go to Properties/Compatibility tick the "Run this program as an administrator" checkbox and apply the changes.



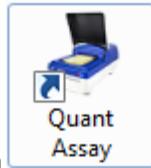


Note! Because of the that modification, we had to move all assays to common documents folder, and your user would need to do that manually. (If you were using versions below 0.7.x.x)

Here is an instruction how to move files:

- 2.1. Open new the software and close it. (This will create needed folders)
- 2.2. Copy all assays: In Program Files(x86)/QuantAssay find folder Methodics. Copy this folder to: C:/Users/Public/Public Documents/QuantAssay/ then Replace conflict files if being asked.

Performing a Measurement



1. Open the program
2. Go to the tab "Available Devices"



3. Select the wavelengths at which you want to measure

Wavelength

| | |
|--|------------------------------------|
| <input checked="" type="checkbox"/> 405 nm | <input type="checkbox"/> Channel 1 |
| <input type="checkbox"/> 450 nm | <input type="checkbox"/> Channel 2 |
| <input type="checkbox"/> 490 nm | <input type="checkbox"/> Channel 3 |
| <input type="checkbox"/> 620 nm | <input type="checkbox"/> Channel 4 |

4. Optional: enter the reference channel and if you would like to mix the plate before the measurement:

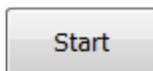
Enable reference Ref. filter, nm

Mix before measure

Mixing

| | | |
|----------------------|----|----|
| Mixing amplitude, mm | 8 | ▲▼ |
| Frequency, 1/s | 12 | ▲▼ |
| Time | 4 | ▲▼ |

5. Click on the "Start" button



6. Then, in approx. 5 to 15 sec., the program will automatically open the tab "Input Data", which will display the measurement results:

| | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 |
|---|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|
| A | 0.001 | 0.001 | 0.001 | 0.002 | 0.002 | 0.002 | 0.002 | 0.002 | 0.003 | 0.003 | 0.003 | 0.003 |
| B | 0.000 | 0.001 | 0.001 | 0.001 | 0.001 | 0.001 | 0.002 | 0.002 | 0.003 | 0.003 | 0.003 | 0.003 |
| C | 0.001 | 0.001 | 0.002 | 0.001 | 0.001 | 0.002 | 0.002 | 0.001 | 0.002 | 0.002 | 0.002 | 0.002 |
| D | 0.001 | 0.001 | 0.002 | 0.003 | 0.003 | 0.003 | 0.003 | 0.003 | 0.003 | 0.003 | 0.003 | 0.003 |
| E | 0.001 | 0.001 | 0.001 | 0.001 | 0.001 | 0.001 | 0.002 | 0.003 | 0.003 | 0.003 | 0.003 | 0.003 |
| F | 0.000 | 0.001 | 0.001 | 0.001 | 0.001 | 0.002 | 0.002 | 0.002 | 0.002 | 0.002 | 0.002 | 0.002 |
| G | 0.001 | 0.001 | 0.002 | 0.002 | 0.002 | 0.002 | 0.002 | 0.002 | 0.002 | 0.002 | 0.002 | 0.002 |
| H | 0.001 | 0.001 | 0.002 | 0.002 | 0.003 | 0.003 | 0.003 | 0.003 | 0.003 | 0.004 | 0.004 | 0.004 |

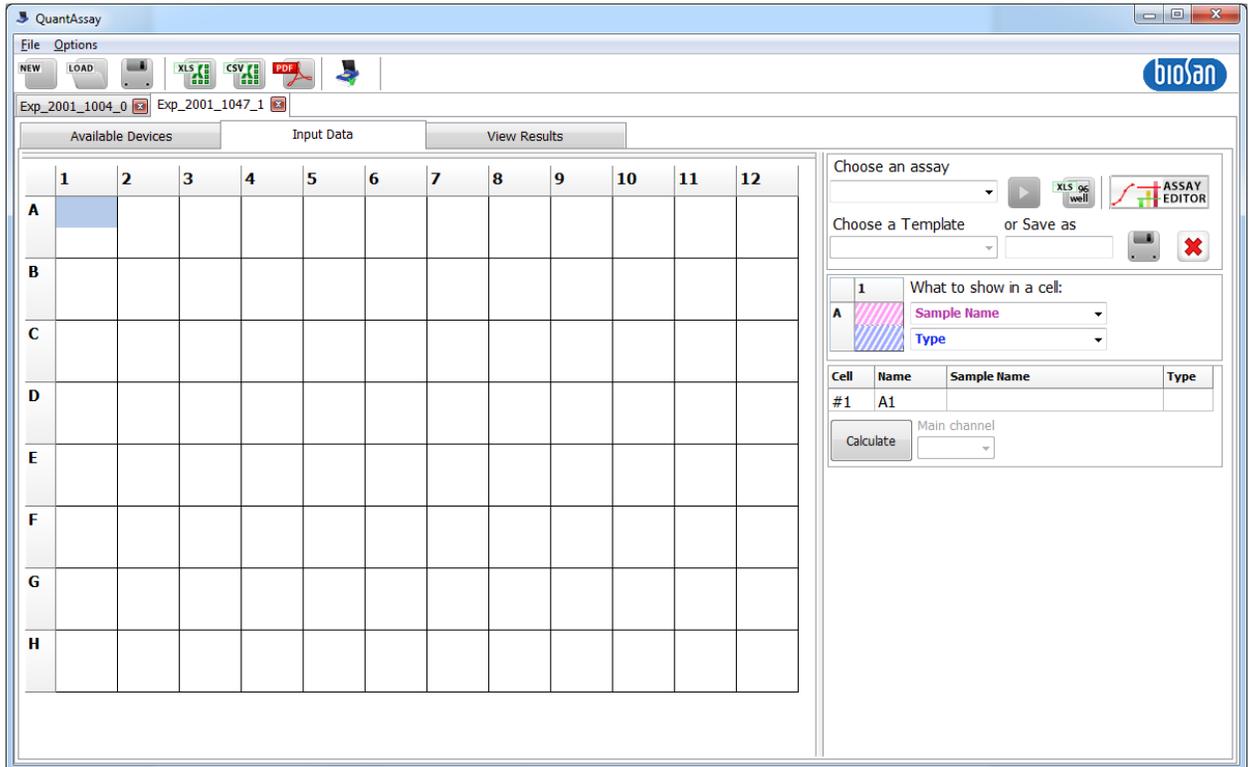
7. To save the experiment in the format of Quant Assay file, click on the “Save” button
8. To save the data in the format of the plate, click on XLS button, which is located next to "Assay editor" button
9. To save the data in .csv .xls .pdf formats, click on the corresponding icons



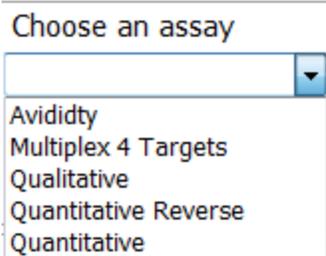
Running a test

1. Open the software

window after opening the program



2. Select an assay (here are listed predefined assays):



- a. Avidity
 - b. Multiplex 4 Targets
 - c. Qualitative
 - d. Quantitative reverse
 - e. Quantitative
3. Each assay is described in more details in the section **Assay Editor**. Here we describe

the use of the simplest assay - Qualitative

4. Qualitative assay:

This assay adopted to put a specified threshold value of Optical Density (OD): A sample is regarded as positive if the corresponding OD value is equal to or greater than the threshold (OD critical), which in this example is calculated by the formula:

$$= \text{Negative Control (N1)} + 0.2$$

where N1 - Is the mean OD value for the negative control samples.

Quality control is also taken into account by following conditions:

- the OD value of the Positive Control (P1) are at least 0.8 OD, where P1 - Is the mean OD value the positive control
- the OD value of the Negative Control (N1) are less than 0.2 OD, where N1 - is the average OD value for the negative control

5. Fill virtual plate with:
Types of samples:



- Test sample



- Background (the average value of those samples will be deducted from the whole plate, the deducted values can only be observed in the Results tab, data input tab will remain the same)



- Positive Control 1



- Negative Control 1 (Threshold/OD critical is calculated based on OD value of those samples)



- Remove the sample



- In this field are specified a name (constant), suffix (counted), and a group (counted). For example, if you add a test sample, it will

be referred to as Smp 1 and will apply to group 1, and the counter of suffix and groups will jump to 2, as shown in the following picture:



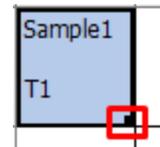
| | 1 | 2 | 3 | 4 |
|---|------------|---|---|---|
| A | Smp1 T1 | | | |

6. The methods of filling the plate:

- a. For a quick filling of the plate with test samples: fill in one of the wells (eg A1), with a test sample.

| | 1 | 2 | 3 | 4 | 5 |
|---|---------------|---|---|---|---|
| A | Sample1 T1 | | | | |
| B | | | | | |
| C | | | | | |
| D | | | | | |

- b. In order to fill all the remaining wells with remaining samples, place the mouse cursor on the small square in the lower right corner of the cell, hold the left mouse button, and lead to the desired cell (as in Excel).



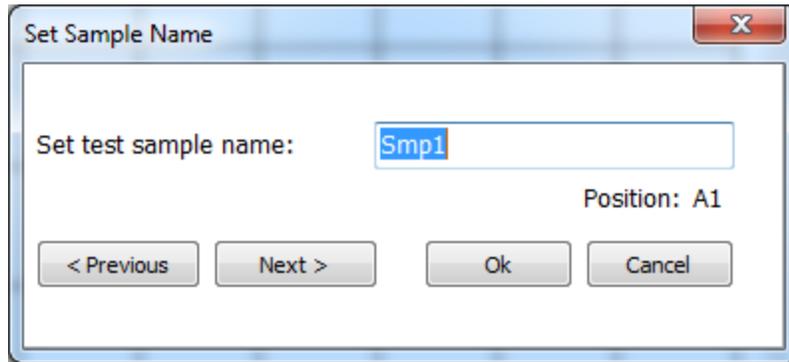
After

| | 1 | 2 | 3 |
|---|---------------|---|---|
| A | Sample1 T1 | | |
| B | | | |
| C | | | |

Before

| | 1 | 2 | 3 |
|---|---------------|---------------|---------------|
| A | Sample1 T1 | Sample4 T4 | Sample7 T7 |
| B | Sample2 T2 | Sample5 T5 | Sample8 T8 |
| C | Sample3 T3 | Sample6 T6 | Sample9 T9 |

- c. To enter the name of the sample, click on the desired cell by double-clicking the mouse. The following window appears:



To confirm the name: press the OK button. To cancel, press: Cancel. To move to the next / previous cell, click on the appropriate button.

- d. To make sample repeats, select two adjacent cells in which the sample is and press on the "Sample" button. To fill the remaining part of the plate in this pattern -- hold down left mouse button on the little black square and drag the mouse to the desired cell. If the samples are filled in 3, 4, etc. repetitions, fill the appropriate number of adjacent cells.

Before

| | 1 | 2 | 3 | 4 |
|---|---------------|---------------|---|---|
| A | Sample1 T1 | Sample1 T1 | | |
| B | | | | |
| C | | | | |

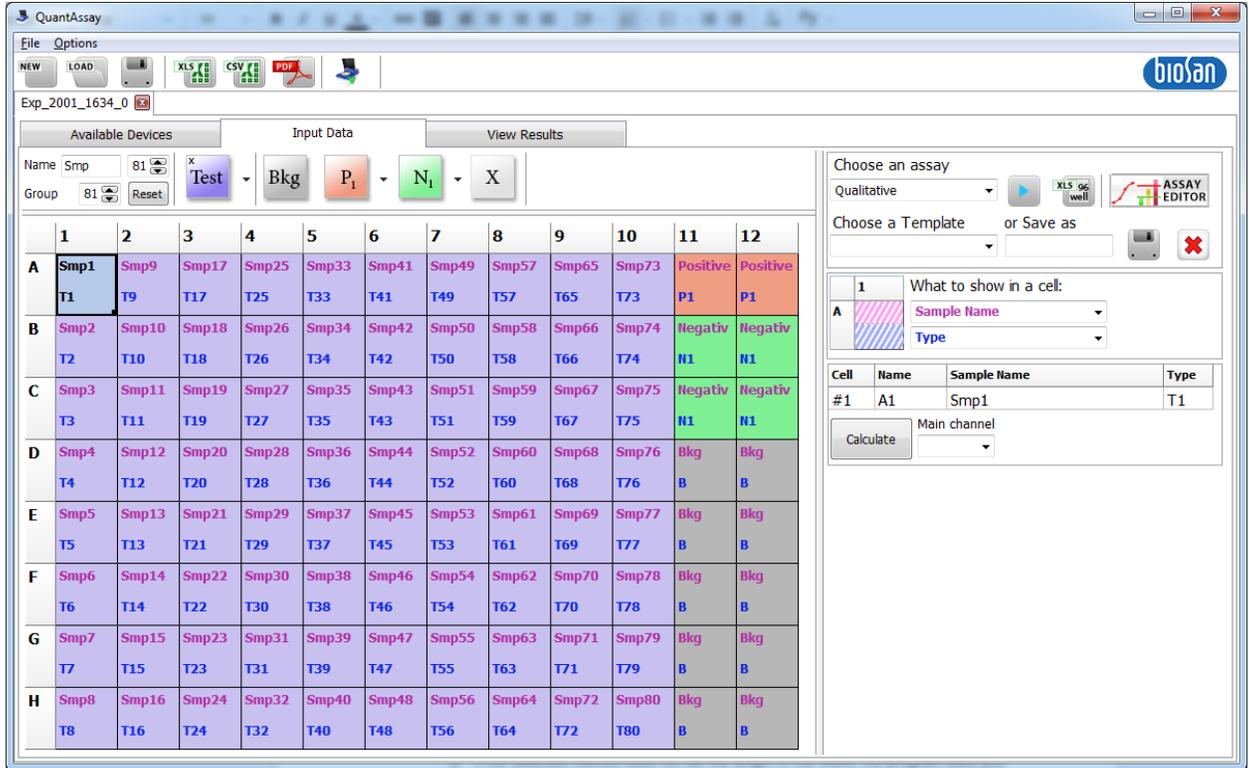
After

| | 1 | 2 | 3 | 4 |
|---|---------------|---------------|---------------|---------------|
| A | Sample1 T1 | Sample1 T1 | Sample4 T4 | Sample4 T4 |
| B | Sample2 T2 | Sample2 T2 | Sample5 T5 | Sample5 T5 |
| C | Sample3 T3 | Sample3 T3 | Sample6 T6 | Sample6 T6 |

- e. Fill in the controls: for positive controls select P1; for negative controls, select the

N1.  

7. Example of a filled plate



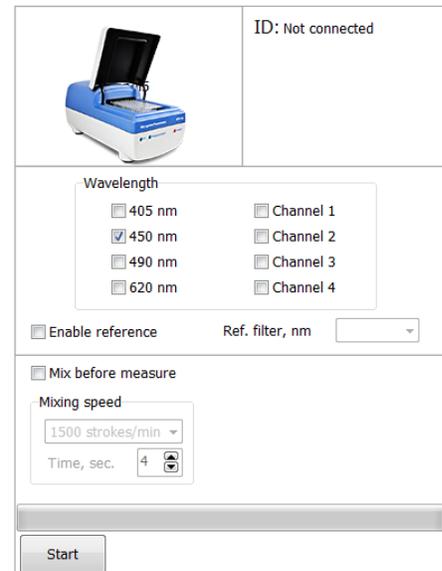
8. to save the template of the plate, enter its name in the "Save As" and press the save

or Save as



icon

9. To start the measurement, click on the "Start" 
 - a. If the selected method the wavelength is not set, then program will jump to the Available Devices tab, where you can set the wavelength and other parameters
 - b. Press "Start" when ready



10. Next, the program will take you back to the “Data Input” tab.

The screenshot shows the QuantAssay software interface. The main window is titled "QuantAssay" and has a menu bar with "File" and "Options". Below the menu bar is a toolbar with icons for "NEW", "LOAD", "XLS", "CSV", "PDF", and a printer icon. The current experiment name is "Exp_2001_1634_0".

The interface is divided into three tabs: "Available Devices", "Input Data", and "View Results". The "Input Data" tab is active. It shows a grid of wells with the following data:

| | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 |
|---|----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-------------------|-------------------|
| A | Smp1 0.000 | Smp9 0.000 | Smp17 0.000 | Smp25 0.000 | Smp33 0.000 | Smp41 0.000 | Smp49 0.000 | Smp57 0.000 | Smp65 0.000 | Smp73 0.000 | Positive 0.000 | Positive 0.000 |
| B | Smp2 -0.001 | Smp10 -0.001 | Smp18 -0.001 | Smp26 -0.001 | Smp34 -0.001 | Smp42 -0.001 | Smp50 -0.001 | Smp58 0.000 | Smp66 0.000 | Smp74 0.000 | Negativ 0.000 | Negativ 0.000 |
| C | Smp3 0.000 | Smp11 0.001 | Smp19 0.001 | Smp27 0.001 | Smp35 0.002 | Smp43 0.002 | Smp51 0.002 | Smp59 0.002 | Smp67 0.002 | Smp75 0.003 | Negativ 0.002 | Negativ 0.003 |
| D | Smp4 0.000 | Smp12 0.001 | Smp20 0.001 | Smp28 0.001 | Smp36 0.001 | Smp44 0.001 | Smp52 0.001 | Smp60 0.001 | Smp68 0.001 | Smp76 0.001 | Bkg 0.001 | Bkg 0.001 |
| E | Smp5 0.000 | Smp13 0.001 | Smp21 0.001 | Smp29 0.001 | Smp37 0.001 | Smp45 0.001 | Smp53 0.001 | Smp61 0.001 | Smp69 0.001 | Smp77 0.001 | Bkg 0.001 | Bkg 0.002 |
| F | Smp6 0.000 | Smp14 0.001 | Smp22 0.002 | Smp30 0.002 | Smp38 0.002 | Smp46 0.002 | Smp54 0.002 | Smp62 0.002 | Smp70 0.002 | Smp78 0.002 | Bkg 0.002 | Bkg 0.002 |
| G | Smp7 -0.001 | Smp15 -0.001 | Smp23 -0.001 | Smp31 -0.001 | Smp39 -0.001 | Smp47 -0.001 | Smp55 -0.001 | Smp63 -0.001 | Smp71 -0.001 | Smp79 -0.001 | Bkg -0.001 | Bkg -0.001 |
| H | Smp8 0.001 | Smp16 0.001 | Smp24 0.001 | Smp32 0.001 | Smp40 0.001 | Smp48 0.001 | Smp56 0.001 | Smp64 0.001 | Smp72 0.001 | Smp80 0.001 | Bkg 0.001 | Bkg 0.001 |

The right-hand panel contains the following controls:

- Choose an assay:** Qualitative (dropdown), with a play button and "XLS og well" icon.
- Choose a Template or Save as:** (dropdown), with a save icon and a red X icon.
- What to show in a cell:** A dropdown menu showing "Sample Name" and "450 nm".
- Cell configuration table:**

| Cell | Name | Sample Name | Type |
|------|------|-------------|------|
| #1 | A1 | Smp1 | T1 |
- Calculate:** A button with a dropdown menu showing "Main channel" and "450 nm".

11. To view the results in the table format, go to “View Results” tab.

| Cell | Type | Sample Name | Group | OD 450 nm | Result 1 | Result 2 | Mean (OD) | Standard Deviation (OD) | Coefficient of Variation (OD) |
|------|------|---------------------|-------|-----------|----------|----------|-----------|-------------------------|-------------------------------|
| A1 | T1 | Smp1 | 1 | 0.000 | - | 0.00 | 0.000 | 0.000 | 0.00% |
| A2 | T9 | Smp9 | 9 | 0.000 | - | 0.00 | 0.000 | 0.000 | 0.00% |
| A3 | T17 | Smp17 | 17 | 0.000 | - | 0.00 | 0.000 | 0.000 | 0.00% |
| A4 | T25 | Smp25 | 25 | 0.000 | - | 0.00 | 0.000 | 0.000 | 0.00% |
| A5 | T33 | Smp33 | 33 | 0.000 | - | 0.00 | 0.000 | 0.000 | 0.00% |
| A6 | T41 | Smp41 | 41 | 0.000 | - | 0.00 | 0.000 | 0.000 | 0.00% |
| A7 | T49 | Smp49 | 49 | 0.000 | - | 0.00 | 0.000 | 0.000 | 0.00% |
| A8 | T57 | Smp57 | 57 | 0.000 | - | 0.00 | 0.000 | 0.000 | 0.00% |
| A9 | T65 | Smp65 | 65 | 0.000 | - | 0.00 | 0.000 | 0.000 | 0.00% |
| A10 | T73 | Smp73 | 73 | 0.000 | - | 0.00 | 0.000 | 0.000 | 0.00% |
| A11 | P1 | Positive control P1 | | 0.000 | Error | | 0.000 | 0.000 | 5.29% |
| A12 | P1 | Positive control P1 | | 0.000 | Error | | 0.000 | 0.000 | 5.29% |
| B1 | T2 | Smp2 | 2 | -0.001 | - | -0.01 | -0.001 | 0.000 | 0.00% |
| B2 | T10 | Smp10 | 10 | -0.001 | - | -0.01 | -0.001 | 0.000 | 0.00% |
| B3 | T18 | Smp18 | 18 | -0.001 | - | -0.01 | -0.001 | 0.000 | 0.00% |
| B4 | T26 | Smp26 | 26 | -0.001 | - | -0.01 | -0.001 | 0.000 | 0.00% |
| B5 | T34 | Smp34 | 34 | -0.001 | - | -0.01 | -0.001 | 0.000 | 0.00% |
| B6 | T42 | Smp42 | 42 | -0.001 | - | -0.01 | -0.001 | 0.000 | 0.00% |
| B7 | T50 | Smp50 | 50 | -0.001 | - | -0.01 | -0.001 | 0.000 | 0.00% |
| B8 | T58 | Smp58 | 58 | 0.000 | - | 0.00 | 0.000 | 0.000 | 0.00% |
| B9 | T66 | Smp66 | 66 | 0.000 | - | 0.00 | 0.000 | 0.000 | 0.00% |
| B10 | T74 | Smp74 | 74 | 0.000 | - | 0.00 | 0.000 | 0.000 | 0.00% |
| B11 | N1 | Negative control | | 0.000 | OK | | 0.001 | 0.001 | 112.87% |
| B12 | N1 | Negative control | | 0.000 | OK | | 0.001 | 0.001 | 112.87% |
| C1 | T3 | Smp3 | 3 | 0.000 | - | 0.00 | 0.000 | 0.000 | 0.00% |

12. To export data in PDF, Excel and CSV click on the corresponding icon



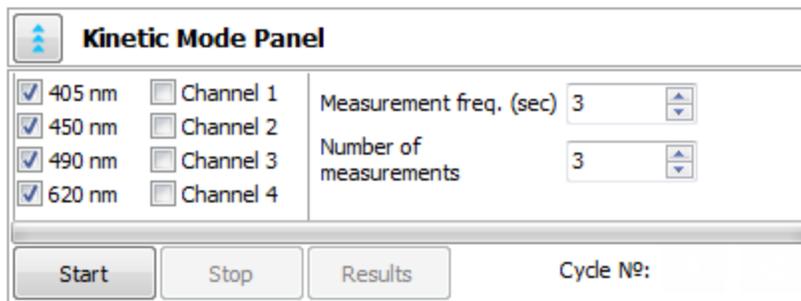
13. To save the experiment in QuantAssay format, click Save icon



Kinetic Mode

To make measurements over time do the following:

Go to Input Data tab and find following panel in the bottom right corner:



The screenshot shows a software panel titled "Kinetic Mode Panel". It contains several controls for setting up measurements:

- Four wavelength options are listed on the left, each with a checked checkbox: 405 nm, 450 nm, 490 nm, and 620 nm.
- Four channel options are listed on the right, each with an unchecked checkbox: Channel 1, Channel 2, Channel 3, and Channel 4.
- Two numerical input fields are on the right: "Measurement freq. (sec)" set to 3 and "Number of measurements" set to 3. Both fields have up and down arrow buttons.
- At the bottom, there are three buttons: "Start", "Stop", and "Results".
- To the right of the buttons is a label "Cycle Nº:" followed by a blank space for a number.

Here simply choose channels, set measurement frequency (in seconds) and number of measurements (in the example above software will do 12 measurements with 3 seconds intervals between).

You can stop the measurements any time by click stop, to get the results click on the Results and in the new tab press XLS button, which will export data to Excel.

If you want to make more measurement, simply put the maximum number of measurements (99999).

Quick conversion table.

1 min = 60 sec, 10 min = 360 sec, 1 hour = 21 600 sec, 2 hour = 23 200 sec.

Assay Editor

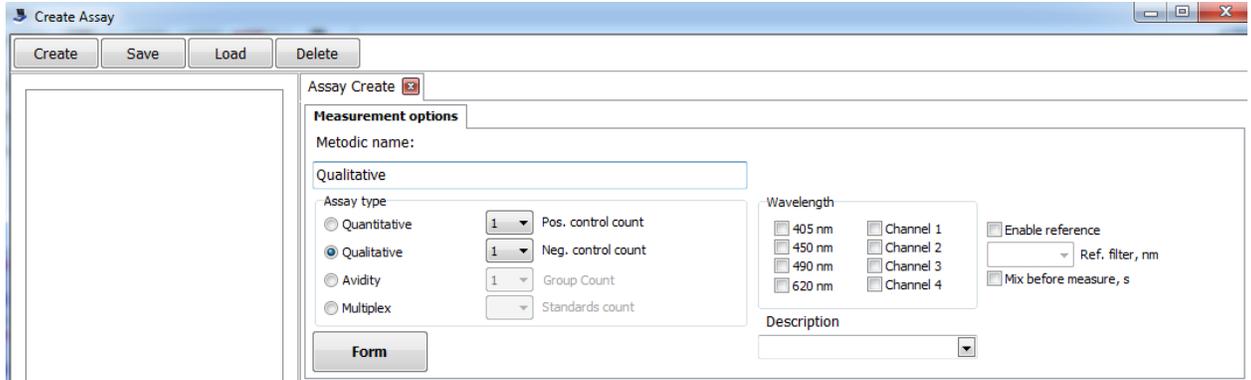
1. Assay Editor will allow you to program the following types of analysis:
 - Qualitative analysis
 - Quantitative analysis: linear and reverse
 - Analysis of avidity
 - Multiplex assay
2. for each type of the assay it is possible to define:
 - the number of types of positive controls (strong, weak, etc.)
 - the number of types of negative controls (no 1' 2' antibody conjugate, water sample)
Note: Each type of the Control can be analyzed separately from the rest of the positive or negative controls
 - For multiplex analysis, you can select the number of targets (antigens)
 - Primary wavelength channel
 - Reference channel (OD values obtained on the reference channel will be subtracted from the OD values obtained on the primary wavelength channel)
 - for quantitative methods: the choice of the calibration curve between the “Best Fit” and piecewise linear models. (Best Fit will automatically select the model with highest coefficient of determination (R^2) among the: 5 parameter logistic, 4 parameter logistic, linear and various regression models.
 - description of the assay

Creating Qualitative assay

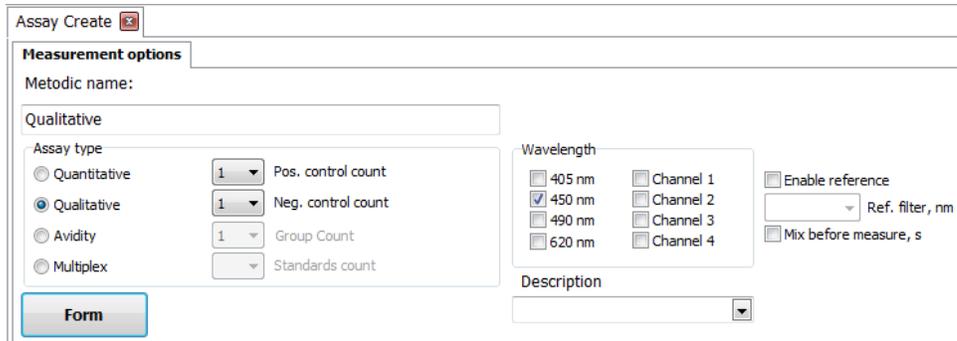
1. For example we need to create a qualitative assay with following criteria:
Measurement channel at 450 nm.
In this assay the sample will analysed as positive, if the corresponding OD value is equal or greater than the Critical (Threshold) OD, which is calculated by the formula:
$$= NC1 + 0.2 OD$$
where NC1 is the Average OD of Negative Control 1.
Quality control of Negative and positive controls should meet following criteria:
 - OD value of the positive control must be greater than 1 OD
 - OD value of the negative control must be less than 0.1 OD

The following steps show how to create this assay:

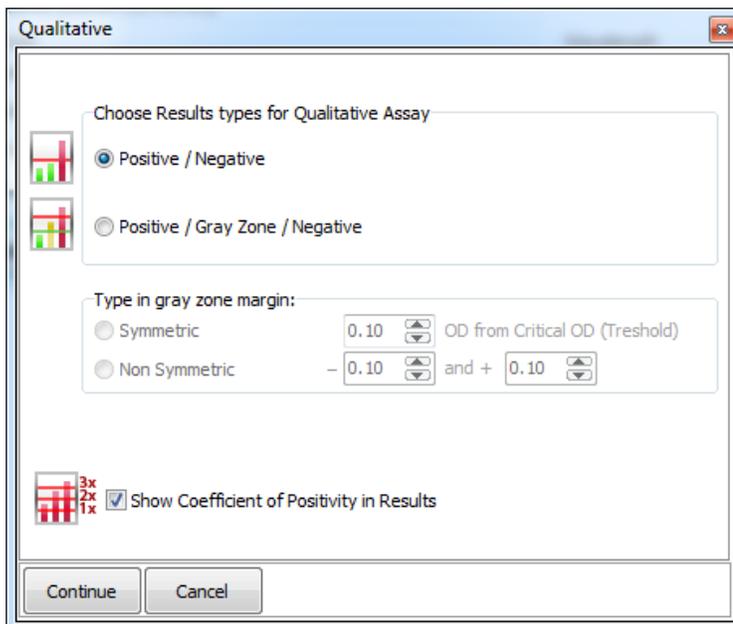
2. Click on “Create” button. Following window will appear:



3. Enter the name of the assay e.g. “Qualitative”, select the type of assay: “Qualitative”, leave the number of the positive/negative Controls: e.g. 1, set the wavelength to: 450 nm. Give a short description of the assay. Click **Form**



4. In the next window you can select the types of results for this assay:



- 1) “Positive / Negative” -- according, if the sample OD is greater or equal to the Critical

(Threshold) OD, the result will be marked as "Positive", else the sample will be marked as "Negative"

2) "Positive / Gray Zone / Negative" -- according, if the sample OD is greater or equal to the threshold OD plus the value indicated in the box "Gray zone = +/-", the result will be marked as "Positive", else if the sample will be between the threshold OD plus/minus OD value indicated in the box below the result will be marked as "Gray Zone", else the sample will be marked "Negative"

Positive / Gray Zone / Negative

Type in gray zone margin:

Symmetric 0.10 OD from Critical OD (Treshold)

Non Symmetric - 0.10 and + 0.10

If you leave the tick in the check box "Show Coefficient of Positivity in Results" that would output the ratio of test sample divided by threshold OD.

Click "Continue" .

5. As we see, the assay editor automatically filled most of the fields in order to analyze the results and to perform quality control. Here is what is being filled automatically and what it means:

6. Tab "Variables and formula"

| Variables and formulas | | |
|------------------------|---------------------------|-----------|
| Variable | Description | Formula |
| [C] | Critical OD | [N1] +0.1 |
| [F] | Coefficient of positivity | [T_0]/[C] |

Two variables were created: [C] and [F], where [C] - is the Critical (Threshold) OD, and [F] - is the ratio of test sample divided by Critical (Threshold) OD or, how we call it — Coefficient of Positivity.

Critical (Threshold) OD is calculated by the formula [N1] +0.1, where [N1] - is the average value of negative control 1. So if N1 is 0.1, than Critical OD will be equal to 0.2 OD

7. Next, we need to perform quality control and analyze our test samples:

Tab "Results Interpretation".

As we see, the assay editor automatically fills most of the fields. Here is what is being filled automatically and what it means:

| Result interpretation | | | | | |
|-----------------------|-------------|----------|-------|----------|-------|
| For variable | Conditional | Result 1 | | Result 2 | |
| | | True | False | True | False |
| [T] | [T]>[C] | + | - | [F] | [F] |
| [P1] | [P1]>1 | OK | Error | | |
| [N1] | [N1]<0.2 | OK | Error | | |

Columns:

- In the column "For variable" you can set for which variable following conditional will be used, e.g. variable [T] means that the conditional and results filled in the next fields will be used for the test samples, to chose another variable, right-click on the field under the column and select an appropriate variable.

- In the column "Conditional" are specified conditional formula by which the "Results 1 and 2" are interpreted, the condition is being interpreted by logical operation "IF, THAN" , and outputs the result in "Result 1 and 2" sub-columns "True" or "False".

In our example:

Condition [T]> [C] means that if the test sample OD ([T]) is greater than the critical OD ([C]), then the "Result" 1 will be "+".

In the column "Result 2", regardless of the condition, positivity coefficient [F] will always be outputted.

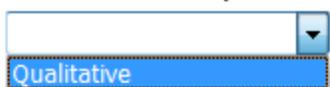
Further, quality control:

For the negative control ([N1_0] is the same as [N1]) is written condition [N1] <0.2, which means that if the OD of neg. control is less than 0.20, then "Result 1" outputs "Ok", if not, then "Error".

For the positive control ([P1_0] is the same as [P1]) the condition is [P1]> 1.0 that is, if the OD of Pos. control is more than 1.0, then "Result 1" outputs "Ok", else "Error".

8. Save the assay and close the window of "Assay Editor"
9. Choose your newly created assay and run it:

Choose an assay



Create a Qualitative Reverse assay with Negative/Suspect/Positive results

This example will feature IDEXX® Pseudorabies Virus gpI Antibody Test Kit®. Go to the calculations chapter of their manual.

First of all, create the assay

The screenshot shows the 'Create Assay' window. The 'Create' button is highlighted. The 'Assay name' field is filled with 'Pseudorabies Virus gpI Antibody Test Kit'. Under 'Assay type', 'Qualitative' is selected. Under 'Wavelength', '620 nm' is selected. The 'Form' button is at the bottom.

Type in the name: Pseudorabies Virus gpI Antibody Test Kit

Assay type select Qualitative

Wavelength set 620nm

Controls

$$NC\bar{x} = \frac{A1 A(650) + A2 A(650) + A3 A(650)*}{3}$$

$$PC\bar{x} = \frac{A4 A(650) + A5 A(650)}{2}$$

*Example shows Negative Control run in Triplicate.

We see that there is used 1 PC and 1 NC, since our software is always calculating the mean of each of the controls (no matter the number of replicates), we don't need to do anything additional here.

Leave the Pos./Neg. control group count as 1 each. Click Form.

Assay type

Quantitative

Qualitative

Avidity

Multiplex

1 Pos. control count

1 Neg. control count

1 Group Count

Standards count

Form

In the next window, select Results type as Positive/Gray Zone/ Negative
 Leave Symmetric gray zone
 Deselect Show coefficient of Positivity in Results, or you can leave it, if you wish.
 Press Continue

Qualitative

Choose Results types for Qualitative Assay

Positive / Negative

Positive / Gray Zone / Negative

Type in gray zone margin:

Symmetric 0.10 OD from Critical OD (Treshold)

Non Symmetric - 0.10 and + 0.10

Show Coefficient of Positivity in Results

Continue Cancel

Now we need to enter the S/N ratio:

Samples

$$S/N = \frac{\text{Sample A(650)}}{NC\bar{x}}$$

In the Variables and formulas change Description for the [F] variable to S/N, and change it's corresponding formula to

[T_0]/[N1]

You can use right click for selecting OD Sample > 0 ,
type in “/” ,
use right click for selecting K (Negative control) 1 > 0

| Variable | Description | Formula |
|----------|---------------------------|------------|
| [C] | Critical OD | [N1]+0.1 |
| [F] | Coefficient of positivity | [T_0]/[N1] |

Go to Result interpretation table, clear all Conditional fields and type in the new conditions from the Calculations chapter of the kit's manual.

First of all we need to enter the Validation criteria, which is:

Validity criteria

$$NC\bar{x} - PC\bar{x} \geq 0.300$$

In the Results interpretation table, for variables P1 and N1, write following conditionals:

[N1]-[P1]>=0.3 , if True = Ok if False = Error,

you can use Right click for selecting the controls.
See below for finished example screenshot.

| Result interpretation | | | | |
|-----------------------|------------------|----------|-------|---|
| For variable | Conditional | Result 1 | | F |
| | | True | False | |
| [P1] | [N1]-[P1] >= 0.3 | OK | Error | |
| [N1] | [N1]-[P1] >= 0.3 | OK | Error | |

Now go to interpretation chapter of the kit's manual, which is:

15 Interpretation:

| Negative | Suspect | Positive* |
|------------|-------------------|------------|
| S/N > 0.70 | 0.60 < S/N ≤ 0.70 | S/N ≤ 0.60 |

*Confirm all positives in duplicate.

Note: IDEXX has instrument and software systems available which calculate results and provide data summaries.

For Variables T (test samples) , write following conditionals:

[F]>0.7 ; True = Negative

([F]>0.6) && ([F]<=0.7) ; True = Suspect

[F]<=0.6 ; True = Positive

you can use Right click for selecting the samples and logical operators.

See below for finished example screenshot.

| Result interpretation | | | | |
|-----------------------|---------------------------------------|----------|-------|-----|
| For variable | Conditional | Result 1 | | Re: |
| | | True | False | |
| [T] | $[F] > 0.7$ | Negative | | |
| [T] | $([F] > 0.6) \ \&\& \ ([F] \leq 0.7)$ | Suspect | | |
| [T] | $[F] \leq 0.6$ | Positive | | |
| [P1] | $[N1] - [P1] \geq 0.3$ | OK | Error | |
| [N1] | $[N1] - [P1] \geq 0.3$ | OK | Error | |

Your assay is good to go!

Create a Quantitative Assay

We want to create a quantitative assay with following criteria:

Measurement channel is 450 nm., with reference channel at 620 nm and mix before measuring. 6 standards with concentrations of: 0, 5, 10, 25, 100, 500 International Units (IU) are being used. Calibration curve should be fitted automatically by choosing the best fitting curve (based on R^2 value) and test samples concentrations will be calculated by using that curve.

We want the test samples OD value of which are greater than OD value of Standard 1 to be marked as positive samples.

We want to exclude extrapolation.

Quality control of Standards and of Negative and Positive controls should meet following criteria:

- Each standard of a higher concentration should have OD greater than the lower standard ($OD_{\text{standard}_0} < OD_{\text{standard}_1}$, $OD_{\text{standard}_1} < OD_{\text{standard}_2}$, etc)
- OD value of the positive control should be greater than 1 OD
- OD value of the negative control should be less than 0.1 OD

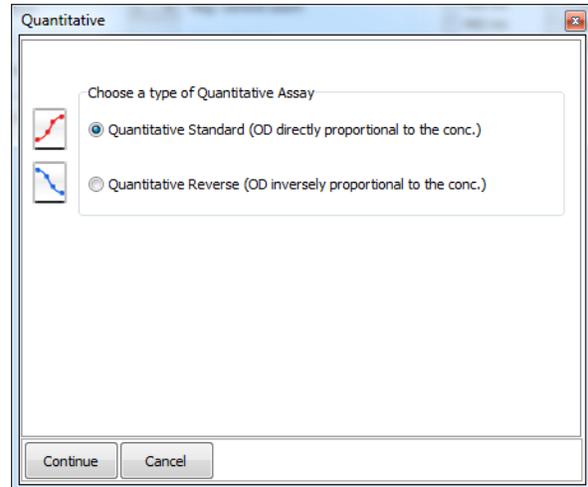
The following steps show procedure of creation of this assay

1. Click on “Create” button. Following window will appear:

The screenshot shows a software window titled "Measurement options". It contains the following fields and controls:

- Assay name:** A text input field containing "Assay Name (05.04 12:39:35)".
- Assay type:** A group of radio buttons and dropdown menus. "Quantitative" is selected. The dropdowns are set to "1" for "Pos. control count", "1" for "Neg. control count", "1" for "Group Count", and "2" for "Standards count".
- Wavelength:** A group of checkboxes for "405 nm", "450 nm", "490 nm", and "620 nm". "450 nm" is checked. There are also checkboxes for "Channel 1", "Channel 2", "Channel 3", and "Channel 4".
- Enable reference:** A checkbox that is checked.
- Ref. filter, nm:** A dropdown menu.
- Mix before measure, s:** A checkbox that is checked.
- Description:** A text input field.
- Curve fit method:** A dropdown menu set to "Best fit (Recommended)".
- Form:** A button at the bottom left.

2. Enter the name of the assay eg “Qualitative”, select the type of assay: “Qualitative”, set the number of standards to 6, leave the number of the “Positive”/“Negative Controls”: eg 1, set the “Wavelength”: 450 nm, set enable reference, select 620 nm and set “Mix before measure, s”. Give a short “Description” of the assay. Leave selection on the “Best fit (Recommended)” in the “Curve fit method” or select the default curve that will be used for calculation. Click “Form”



Measurement options

Assay name:

Assay type

| | | |
|---|--------------------------------|--------------------|
| <input checked="" type="radio"/> Quantitative | <input type="text" value="1"/> | Pos. control count |
| <input type="radio"/> Qualitative | <input type="text" value="1"/> | Neg. control count |
| <input type="radio"/> Avidity | <input type="text" value="1"/> | Group count |
| <input type="radio"/> Multiplex | <input type="text" value="6"/> | Standards count |

Wavelength

| | |
|---|------------------------------------|
| <input type="checkbox"/> 405 nm | <input type="checkbox"/> Channel 1 |
| <input checked="" type="checkbox"/> 450 nm | <input type="checkbox"/> Channel 2 |
| <input type="checkbox"/> 490 nm | <input type="checkbox"/> Channel 3 |
| <input checked="" type="checkbox"/> 620 nm(Ref) | <input type="checkbox"/> Channel 4 |

Enable reference
 Ref. filter, nm
 Mix before measure, s

Constants list

Description

Curve fit method

3. In the next window you can select the type of the quantitative assay: “Quantitative Standard” or “Quantitative Reverse” (“Reverse” means that with the increase of concentration the OD is decreasing, “Standard” means that with the increase of concentration the OD is also increasing)
4. As we see, the assay editor automatically fills most of the fields to analyze the results and to perform quality control. Here is what is being filled automatically and what it means:
5. Tab “Variables and formulas”

| Variables and formulas | | Standards |
|------------------------|---------------------------|--------------|
| Variable | Description | Formula |
| [C] | Critical OD | $[N1] + 0.1$ |
| [F] | Coefficient of positivity | $[T_0]/[C]$ |
| | | |

Two variables were created: [C] and [F], where [C] - is the “Critical (Threshold) OD”, and [F] - is the ratio of test sample divided by threshold OD or “coefficient of positivity”. In our example both formulas are irrelevant as we will quantify the results by fitting plots via standards. What we need to set is the concentration of standards, to do that click on the Standards tab

| Variables and formulas | | Standards |
|------------------------|---------------|-----------|
| Variable | Concentration | Units |
| [S0] | | |
| [S1] | | |
| [S2] | | |
| [S3] | | |
| [S4] | | |
| [S5] | | |

In column “Variable” [S0], [S1], etc. stands for Standard 0, Standard 1, etc. In column “Concentration” fill in the concentration values. In field “Units” choose “IU” (international units).

| Variables and formulas | | Standards |
|------------------------|---------------|-----------|
| Variable | Concentration | Units |
| [S0] | 0 | IU |
| [S1] | 5 | IU |
| [S2] | 10 | |
| [S3] | 25 | |
| [S4] | 100 | |
| [S5] | 500 | |

- Next, we need to perform quality control and analyze our test samples:
Tab “Results Interpretation”.
As we see, the assay editor automatically filled most of the fields. Here is what is being filled automatically and what it means:

| Result interpretation | | | |
|-----------------------|------------------------------------|----------|--------------|
| For variable | Conditional | Result 1 | |
| | | True | False |
| [S0] | [S0] < [S1] | OK | Error |
| [S1] | [S1] < [S2] | OK | Error |
| [S2] | [S2] < [S3] | OK | Error |
| [S3] | [S3] < [S4] | OK | Error |
| [S4] | [S4] < [S5] | OK | Error |
| [S5] | [S4] < [S5] | OK | Error |
| [T] | (([SMin] < [T]) && ([T] < [SMax])) | In Range | Out of Range |
| [P1] | [P1] > 1 | OK | Error |
| [N1] | [N1] < 0.2 | OK | Error |

Columns:

- In the column "For variable" you can set for which variable following conditional will be used, eg variable [S0] means that the conditional and results filled in the next fields will be used for the Standard 0. To chose other variable, right-click on the field under the column.

- In the column "Conditional" are specified conditional formula by which the "Results 1 and 2" are interpreted, the condition is being interpreted by logical operation "IF, THAN" , and outputs the result written in "Result 1" sub-columns "True" or "False".

In our example:

Condition [S0] < [S1] means that if the Standard 0 ([S0]) is less than the Standard 1 ([S1]), then the "Result 1" will be "Ok", else it will be "Error".

Further:

Analysis of test samples:

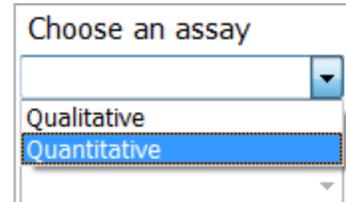
| | | | |
|-----|------------------------------------|----------|--------------|
| [T] | (([SMin] < [T]) && ([T] < [SMax])) | In Range | Out of Range |
|-----|------------------------------------|----------|--------------|

The conditional (([Smin] < [T]) && ([T] <= [Smax])) means that any test sample ([T]) that is

greater than Standard minimum ([S1]) and less or equal to Standard maximum ([S5]) will be outputted as “In range” in “Result 1”, else it will be outputted as “Out of range”
That is how we can exclude extrapolation. **However the calculated concentration value will be outputted in both cases.**

For the negative control ([N1_0] is the same as [N1]) is written a condition [N1] <0.2, which means that if the OD of neg. control is less than 0.2 OD, then “Result 1” outputs "Ok", else outputs "Error".

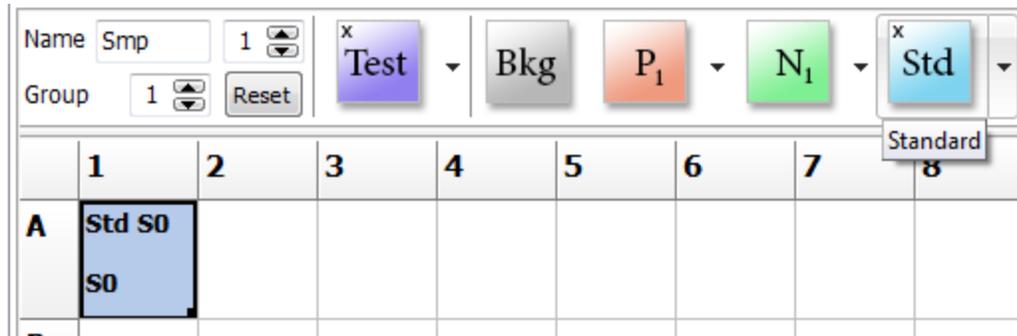
For the positive control ([P1_0] is the same as [P1]) the condition is [P1]>1.0 that is, if the OD of Positive control is more than 1.0, then “Result 1” outputs "Ok", else "Error".



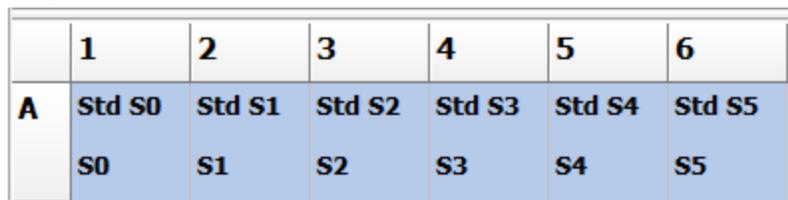
7. Save the assay and close the window of “Assay Editor”
8. Choose your newly created assay and run it:

9. In order to add Standards do the following:

- a. Select the well with Standard 0 (if in duplicate select 2 wells)

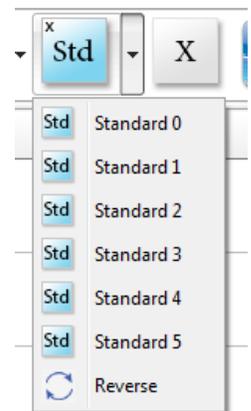


- b. Drag the black square till the well with the last standard



Or manually chose a well and add a needed standard by choosing from drop box

- c. Setting samples and controls, and obtaining results is the same as in Qualitative assay



Create a quantitative assay with concentration based interpretation

There is a possibility to base your interpretation of results not based on OD, but on calculated concentration, to do that, do the following: when setting conditional, use concentration type of value (e.g. [O])

| | |
|------------------------------|------------------|
| [S2] | 3 |
| [S3] | 4 |
| Result interpretation | |
| For variable | Condition |
| [S4] | [S4] < [S5] |
| [S5] | [S4] < [S5] |
| [T] | |

Concentration

Critical OD ▶

Coefficient of positivity ▶

Standards ▶

Wild Card ▶

Logical operators ▶

Formula (Sample OD within standard range)

In the following example, all test samples with calculated concentration greater than 3 units, will result as a positive (+) result, other will output as negative (-). Standards and Controls can be interpreted in the same way.

| | | | |
|-----|---------|---|---|
| [T] | [O] > 3 | + | - |
|-----|---------|---|---|

Create a quantitative assay with qualitative interpretation

In the following example: all test samples with calculated concentration less than 1 unit, will result as a negative (-) result, samples with calc. concentration value from 1 (including) to 3, will output as a gray zone (+/-) result, samples with calc. concentration greater or equal to 3, will output as positive (+) result.

| | | | |
|-----|---------------------|-----|--|
| [T] | [O] < 1 | - | |
| [T] | [O] >= 1 && [O] < 3 | +/- | |
| [T] | [O] >= 3 | + | |

Create an avidity assay

The screenshot shows a software window titled 'Assay Create' with a sub-tab 'Measurement options'. The 'Metodic name:' field contains 'Assay Name (27.01 17:30:40)'. Under 'Assay type', 'Avidity' is selected with a value of '1' in the adjacent dropdown, and 'Pos. control count' is set to '1'. Other options include 'Quantitative', 'Qualitative', and 'Multiplex'. The 'Wavelength' section has '450 nm' selected, with other options at 405 nm, 490 nm, and 620 nm. Channel selection includes 'Channel 1' through 'Channel 4'. Additional checkboxes for 'Enable reference', 'Ref. filter, nm', and 'Mix before measure, s' are present. A 'Description' field is at the bottom. A 'Form' button is located at the bottom left.

1. Avidity assays analyses the samples by calculating the Index of Avidity (IA) of positive test samples. Index of Avidity is the ratio of optical density of a sample in the presence of a dissociating agent (dissociation ELISA) to an optical density of the same sample without dissociating agent (direct ELISA).

We want to create an avidity assay with following criteria:

So we begin by selecting a type of assay: Avidity.

Measurement channel is set to e.g. 450 nm.

Analyzed sample is regarded as positive if sample's $OD_{\text{of direct ELISA}}$ is greater or equal to the Critical (Threshold) OD which is calculated by the formula:

= $NC1 + 0.2 OD$, where NC1 is the Average OD of Negative Controls 1.

Index of avidity shall be calculated for all positive test samples by the formula:

= $OD_{\text{dissociation ELISA}} / OD_{\text{direct ELISA}}$

All positive samples should be divided into 3 groups:

- Samples with avidity index less than 0.30 (or 30%), those samples will be marked as "+" (e.g. low avidity antibodies)
- Samples with avidity index greater than 0.30 (or 30%) and less or equal to 0.50 (or 50%), those samples will be marked as "++" (e.g. normal avidity antibodies)
- Samples with avidity index greater than 0.50 (or 50%), those samples will be marked as "+++" (high avidity antibodies)

Quality control of negative and positive controls should meet following criteria:

- $OD_{\text{direct ELISA}}$ of the positive control must be greater than 1 OD and the index of avidity greater than 0.30 (30%);
- OD value of the negative control must be less than 0.1 OD

The following steps show procedure of creation of this assay

2. Create a new assay, click "New"
3. Enter the name of the assay, eg Avidity, select the type of assay: Avidity, set the wavelength channel. Number of controls: 1 Negative Control, 1 Positive Control. Give a short description of the assay. Click "Form"

Measurement options

Metodic name: Avidity

Assay type

- Quantitative 1 Pos. control count
- Qualitative 1 Neg. control count
- Avidity 2 Avidity count
- Multiplex Standards count

Wavelength

- 405 nm Channel 1
- 450 nm Channel 2
- 490 nm Channel 3
- 620 nm Channel 4

Enable reference

Ref. filter, nm

Mix before measure, s

Description

Form

4. In the following window you can set the analysis of the results:

Avidity

Choose Results types for Avidity Assay

- Positive / Negative
- Positive / Gray Zone / Negative

Gray zone margin:
by 1.00 to 2.00 higher than OD critical (treshold) value

Type in avidity index margins for positive samples and it's result

| | Margin | Result |
|----------|-----------------|--------|
| If AI < | 0.30 | + |
| If AI >= | 0.30 and 0.50 < | ++ |
| If AI >= | 0.50 | +++ |

Show Avidity index in Results

Continue Cancel

- 1) Positive / Negative, e.g. according if the OD of the sample is greater than or less than the threshold OD, program will output results "Positive" or "Negative", Avidity index will be calculated only for positive test samples
- 2) Positive / Gray Zone / Negative -- according, if the sample OD is greater or equal than the threshold OD multiplied by the value indicated in the field "to" (e.g. 2), the result will

be marked as "positive", else if the sample will be between the threshold OD multiplied by the value in the value indicated in the field "by" (e.g. 1) and "to" (e.g. 2) the result will be marked as Gray Zone, else the sample will be marked as "Negative".

Avidity index will be calculated only for positive test samples

- Next, you need to type in avidity index' margins for positive test samples and corresponding result:

Type in avidity index margins for positive samples and it's result

| Margin | Result |
|--------------------------|--------|
| If AI < 0.30 | + |
| If AI >= 0.30 and 0.50 < | ++ |
| If AI >= 0.50 | +++ |

 Show Avidity index in Results

In this example avidity index (AI) below or equal to 0.30, will output "+" in results
 If the avidity index (AI) is between 0.30 and 0.50, will output "++" in results,
 and if the avidity index (AI) is equal to or greater than 0.50, will output "+++" in results
 Leave the checkbox "Show avidity index in results" that would output the AI in the results.

Click "Continue."

- As we see, the assay editor automatically fills most of the fields in order to analyze the results and perform quality control.

Here is what is being filled automatically and what it means:

- Tab "Variables and formulas"

Variables and formulas

| Variable | Description | Formula |
|----------|--------------|-------------|
| [C] | Critical OD | [N1] +0.1 |
| [R] | Sample Ratio | [T_1]/[T_0] |

Two variables were created: [C] and [R], where [C] - is the Critical (Threshold) OD, and [R] - is the Avidity Index.

Critical (Threshold) OD is calculated by the formula [N1] +0.1, where [N1] - is the average value of negative control 1. So if N1=0.1, than Critical OD = 0.2 OD

[R] is calculated by the formula [T_1] / [T_0], where [T_1] is sample with a dissociating agent (dissociation ELISA) - and [T_0] - sample without dissociating agent (direct ELISA)

8. Next, we need to perform quality control and analyze our test samples:

As we see, the assay editor automatically fills most of the fields. Here is what is being filled automatically and what it means:

| Result interpretation | | | | | |
|-----------------------|-----------------------------------|----------|-------|----------|-------|
| For variable | Conditional | Result 1 | | Result 2 | |
| | | True | False | True | False |
| [T] | [T_0]<[C] | - | | | |
| [T] | [R]<0.3 && [T_0]>=[C] | + | | [R] | |
| [T] | [R]>=0.3 && [R]<0.5 && [T_0]>=[C] | ++ | | [R] | |
| [T] | [R]>=0.5 && [T_0]>=[C] | +++ | | [R] | |

Columns:

- In the column "For variable" you can set for which variable following conditional will be used, e.g. variable [T] means that the conditional and results filled in the next fields will be used for the test samples, to chose another variable, right-click on the field under the column and choose a needed variable.

- In the column "Conditional" are specified conditional formula by which the "Results 1 and 2" are interpreted, the condition is being interpreted by logical operation "IF, THAN" , and outputs the result written in "Result 1" sub-columns "True" or "False".

In our example:

Conditional [T_0]< [C] means that if the test sample's OD_{of direct ELISA} ([T_0]) is less than the critical OD ([C]), then the "Result" 1 will be "-".

Conditional [R]<0.3 && [T_0]>=[C] means, that if the Avidity Index is less than 0.3 **AND** the OD_{of direct ELISA} is greater or equal to Threshold OD, then the "Result 1" will be "+" and Avidity Index will be written in "Result 2"

Conditional [R]>=0.3 && [R]<0.5 && [T_0]>=[C] means, that if the Avidity Index is greater or equal to 0.3 **AND** is less than 0.5 **AND** the OD_{of direct ELISA} is greater or equal to Threshold OD, then the "Result 1" will be "++" and Avidity Index will be written in "Result 2"

Conditional [R]>=0.5 && [T_0]>=[C] means, that if the Avidity Index is greater or equal to 0.5 **AND** the OD_{of direct ELISA} is greater or equal to Threshold OD, then the "Result 1" will be "+++" and Avidity Index (Avidity Index) will be written in "Result 2"

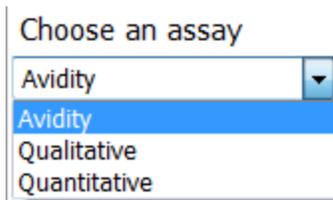
Further, quality control

| Result interpretation | | | |
|-----------------------|-------------------------------------|----------|-------|
| For variable | Conditional | Result 1 | |
| | | True | False |
| [P1_0] | [P1_0]>=[C] && ([P1_1]/[P1_0]>=0.3) | OK | Error |
| [P1_1] | [P1_0]>=[C] && ([P1_1]/[P1_0]>=0.3) | OK | Error |
| [N1_0] | [N1_0]<[C] | OK | Error |
| [N1_1] | [N1_1]<[C] | OK | Error |

For the positive controls ([P1_0] and [P1_1]) we need to check if OD_{of direct ELISA} is greater than the threshold OD ([P1_0]>=[C]) **AND** the avidity index should be greater or equal to 0.3, since we do not have a variable for positive controls' avidity indexes, we need to specify it separately either in Tab "Variables and Formula" or specify it here: ([P1_1]/[P1_0]>=0.3), the "Result 1" outputs "Ok", else "Error".

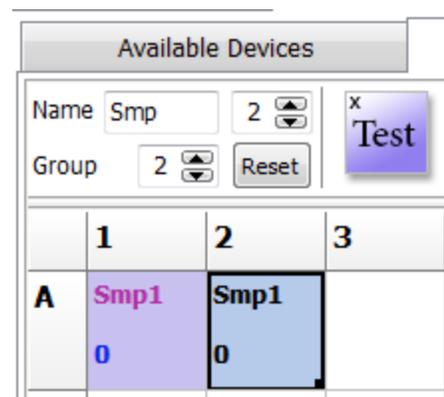
For the negative control ([N1_0] and [N1_1]) we need to check if OD_{of direct ELISA} is lower than the Critical (Threshold) OD, so the expression [N1_0]<[C] means that if the OD of negative control is less than Critical (Threshold) OD, then "Result 1" outputs "Ok", if not, then "Error".

9. Save the assay and close the "Assay Editor"
10. Choose your newly created assay and run it:
11. Choose the assay from the list and run it:

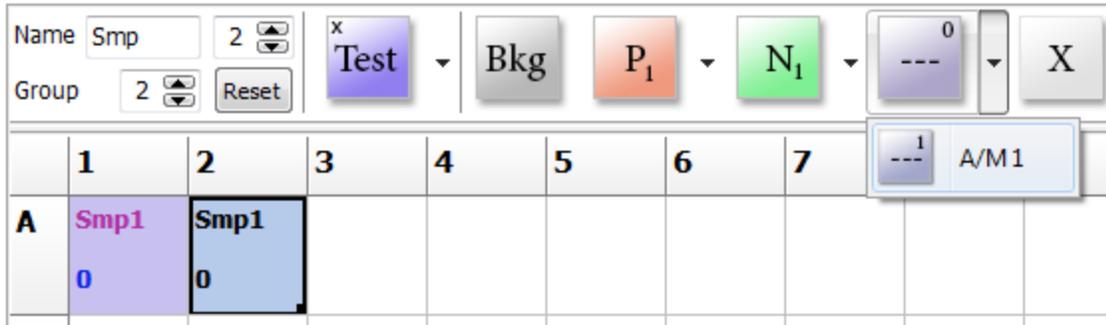


12. In order to fill the samples do the following:
Set 2 samples in two adjacent wells (or controls) and select the sample with protein dissociating agent (e.g. urea).

Click on the droplist near the "----0" button, A/M 1 button



will appear, click on it.



13. Now the sample in A2 is with urea, the bottom part of the well will now show “Control Reagent” type. **Note:** The well with urea will now appear with a slightly different color. Select any other well in order to see it. Controls will not appear with different color.

| | 1 | 2 |
|---|-----------|-----------|
| A | Smp1 0 | Smp1 1 |

14. To fill the plate with the same pattern, select both wells and drag the mouse till the end.

| | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 |
|---|-----------|-----------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| A | Smp1 0 | Smp1 1 | Smp9 0 | Smp9 1 | Smp17 0 | Smp17 1 | Smp25 0 | Smp25 1 | Smp33 0 | Smp33 1 | Smp41 0 | Smp41 1 |
| B | Smp2 0 | Smp2 1 | Smp10 0 | Smp10 1 | Smp18 0 | Smp18 1 | Smp26 0 | Smp26 1 | Smp34 0 | Smp34 1 | Smp42 0 | Smp42 1 |
| C | Smp3 0 | Smp3 1 | Smp11 0 | Smp11 1 | Smp19 0 | Smp19 1 | Smp27 0 | Smp27 1 | Smp35 0 | Smp35 1 | Smp43 0 | Smp43 1 |
| D | Smp4 0 | Smp4 1 | Smp12 0 | Smp12 1 | Smp20 0 | Smp20 1 | Smp28 0 | Smp28 1 | Smp36 0 | Smp36 1 | Smp44 0 | Smp44 1 |
| E | Smp5 0 | Smp5 1 | Smp13 0 | Smp13 1 | Smp21 0 | Smp21 1 | Smp29 0 | Smp29 1 | Smp37 0 | Smp37 1 | Smp45 0 | Smp45 1 |
| F | Smp6 0 | Smp6 1 | Smp14 0 | Smp14 1 | Smp22 0 | Smp22 1 | Smp30 0 | Smp30 1 | Smp38 0 | Smp38 1 | Smp46 0 | Smp46 1 |
| G | Smp7 0 | Smp7 1 | Smp15 0 | Smp15 1 | Smp23 0 | Smp23 1 | Smp31 0 | Smp31 1 | Smp39 0 | Smp39 1 | Smp47 0 | Smp47 1 |
| H | Smp8 0 | Smp8 1 | Smp16 0 | Smp16 1 | Smp24 0 | Smp24 1 | Smp32 0 | Smp32 1 | Smp40 0 | Smp40 1 | Smp48 0 | Smp48 1 |

15. When setting controls, be sure to put "A/M 1" again. **Note** Controls with urea will not appear with different color, to check if you have entered control with urea, choose "Control Reagent" in "What to show in a cell", controls with urea will have "1" in the bottom part of the cell

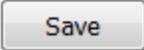
| | | | | |
|---|---|-------------------------|----------|----------|
| | 1 | What to show in a cell: | Positive | Positive |
| A |  | Sample Name | 0 | 1 |
| |  | Control Reagent | Negativ | Negativ |
| | | | 0 | 1 |

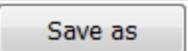
16. Obtaining results is the same as in other assays.

Tools for assay editing

Note, that only master users (administrator account in windows) can create or edit assays.

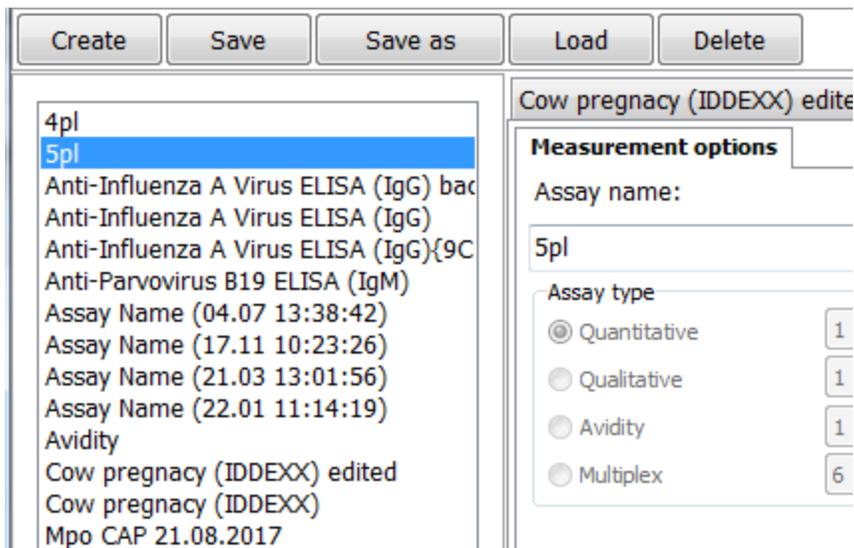
 - Creates a new assay

 - Saves the assay

 - Saves a copy of an assay (including all referenced templates)

 - Loads an assay

 - Deletes a selected assay from the list of assays.



The screenshot shows a software interface for editing assays. At the top, there are five buttons: 'Create', 'Save', 'Save as', 'Load', and 'Delete'. Below these is a list of assays. The '5pl' assay is selected and highlighted in blue. The list includes:

- 4pl
- 5pl
- Anti-Influenza A Virus ELISA (IgG) bac
- Anti-Influenza A Virus ELISA (IgG)
- Anti-Influenza A Virus ELISA (IgG){9C
- Anti-Parvovirus B19 ELISA (IgM)
- Assay Name (04.07 13:38:42)
- Assay Name (17.11 10:23:26)
- Assay Name (21.03 13:01:56)
- Assay Name (22.01 11:14:19)
- Avidity
- Cow pregnancy (IDDEXX) edited
- Cow pregnancy (IDDEXX)
- Mpo CAP 21.08.2017

To the right of the list is a panel titled 'Cow pregnancy (IDDEXX) edite'. It has a tab labeled 'Measurement options'. Below the tab, there is a field for 'Assay name:' containing '5pl'. Underneath, there is a section for 'Assay type' with four radio button options, each with a corresponding count in a small box:

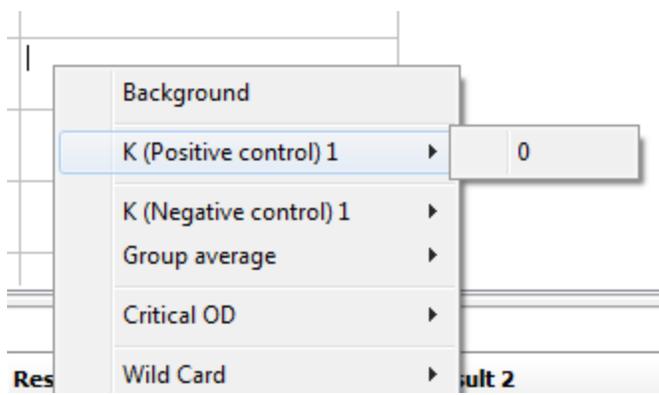
- Quantitative 1
- Qualitative 1
- Avidity 1
- Multiplex 6

Using new variables: Wildcards

While creating new assays, users can use other than preset variables like Critical OD [C] or Coefficient of positivity [F]. Those variables are called Wildcards [W], and user can use 7 new variables per assay.

One of the examples of use:

User needs to see the ratio for positive and negative controls, for that he needs to do the following: add new variables by right clicking on the Variable empty cell and selecting a Wildcard, add a suitable description, under the formula column input following: Pos. control divided by critical OD: [P1]/[C], **input of variables can only be done by right clicking on that cell**, mathematical operators (+, -, *, /) can only be inputted from keyboard.



same for negative and other pos. control.

| Variables and formulas | | |
|------------------------|-----------------------|----------|
| Variable | Description | Formula |
| [W1] | ratio for Pos control | [P1]/[C] |
| [W2] | ratio for neg control | [N1]/[C] |
| | | |

| Result interpretation | | | | | |
|-----------------------|-------------|----------|-------|----------|-------|
| For variable | Conditional | Result 1 | | Result 2 | |
| | | True | False | True | False |
| [N1] | [N1]<0.4 | OK | Error | [W2] | [W2] |
| [P1_0] | [P1]>1 | Ok | Error | [W1] | [W1] |

Logical operations in the interpretation of results

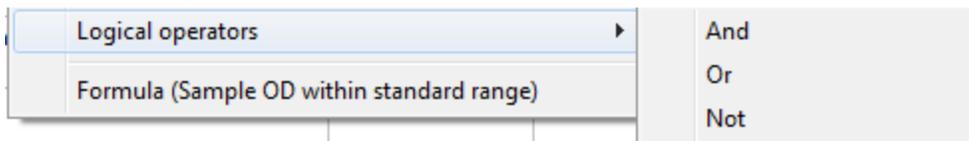
Logical expressions can take one of two values - "true" or "false". Logic operations are used for complex logical expressions. We use logical operations as conditions for determining the results of the program.

For example, Condition:

| For variable | Conditional | Result 1 | |
|--------------|-----------------------|----------|----------|
| | | True | False |
| [T] | [T_0]>=1 && [T_1] >=2 | Positive | Negative |

Here we have two conditions: OD sample type T_0 and OD sample type T_1, if the OD of both samples is greater than or equal to 1, then the conditions of "Yes" and as a result in front of the sample It will be written the result of "laid." If not, the result will be written "Neg."

to set other logical operators, choose from the menu by clicking the right button of the mouse



Using Standard Deviation

If you are using replicates, It is also possible to use Standard Deviation value for the calculations. Below is the example where the Critical OD is being calculated by OD value of Negative Control plus 3 Standard deviations.

| Variables and formulas | | |
|------------------------|-------------|---------------|
| Variable | Description | Formula |
| [C] | Critical OD | [N1]+3*[N1_S] |

Models for quantitative analysis

For building calibration curves we

1. 5-parameter logistic model
2. 4-parameter logistic model
3. linear model
4. Piecewise linear model

5-parameter logistic model (5PL)

5-parameter logistic or 5PL nonlinear regression model that is used to analyze data in biological or immunological samples, such as ELISA or curves dose / response. It differs from the 4PL or 4-parameter logistic model in that it is asymmetric function and is better suited for immunological or biological data.

We use 2 5PL formulas:

$$F(x) = A + \frac{D}{\left(1 + \left(\frac{x}{C}\right)^B\right)^E} \qquad F(x) = \frac{A - D}{\left(1 + \left(\frac{x}{C}\right)^B\right)^E} + D$$

$$F(x) = A + (D/(1+(x/C)^B)^E) \quad \text{or} \quad F(x) = (A-D)/(1+(x/C)^B)^E + D$$

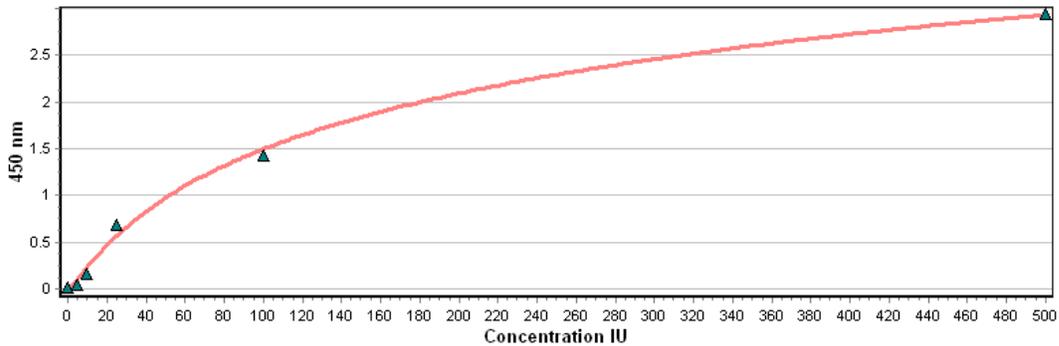
where:

- A — the OD value for the minimum asymptote
- B — the Hill slope
- C — the concentration at the inflection point
- D — the OD value for the maximum asymptote
- E — is the asymmetry factor

5 Parameter Logistics 1

$$y = A + (D / (1 + (X/C)^B))^E$$

A = 388.50, D = -388.51, C = 24.19, B = 1.19, E = 0.00
R-Square = 1.00



4-parameter logistic model (4PL)

4-parameter logistic or 4PL nonlinear regression model is used to analyze data in a biological or immunological samples, such as ELISA or curve dose / response. in 4PL 4

Formula:

$$F(x) = \frac{A - D}{1 + \left(\frac{x}{C}\right)^B} + D$$

$$F(x) = (A - D) / (1 + ((x/C)^B)) + D$$

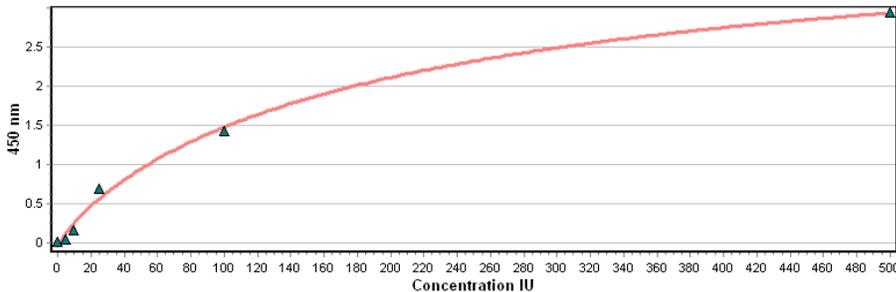
where:

- A — the OD value for the minimum asymptote
- B — the Hill slope
- C — the concentration at the inflection point
- D — the OD value for the maximum asymptote

4 Parameter Logistics

$$y = (A - D) / (1 + (x/C)^B) + D$$

A = -0.05, D = 4.30, C = 203.66, B = 0.87
R-Square = 0.99

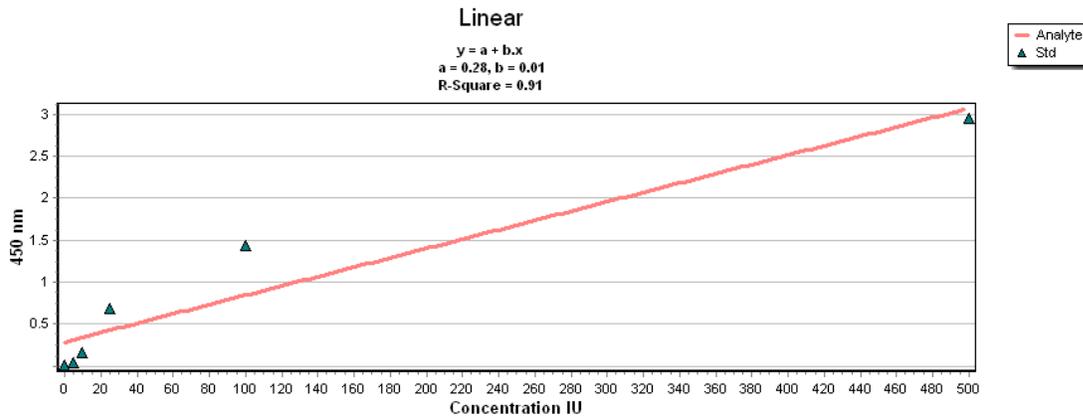


Linear model

linear function - the function of the form

$$y = kx + b$$

basic functions: increment of the function is proportional to the increment of the argument (concentration).

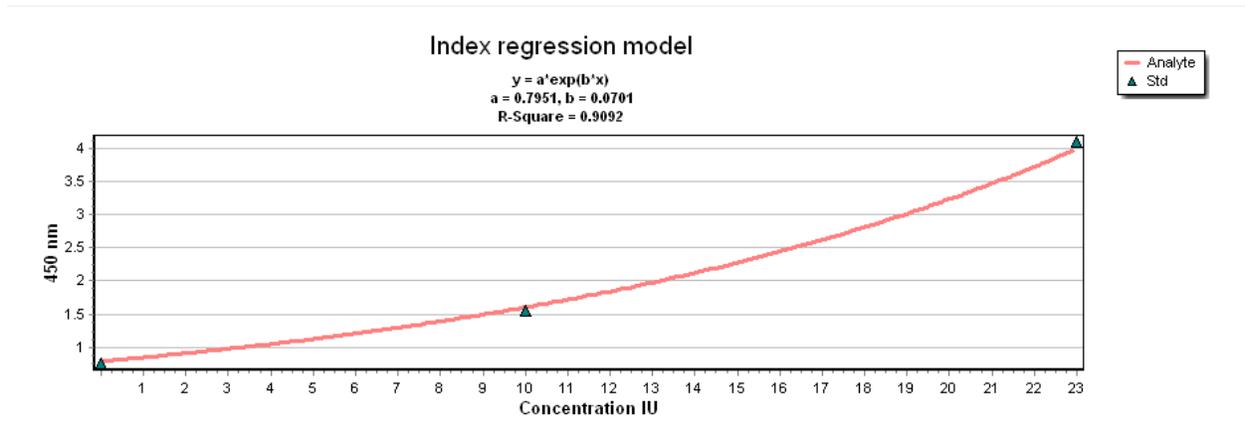


The piecewise linear model

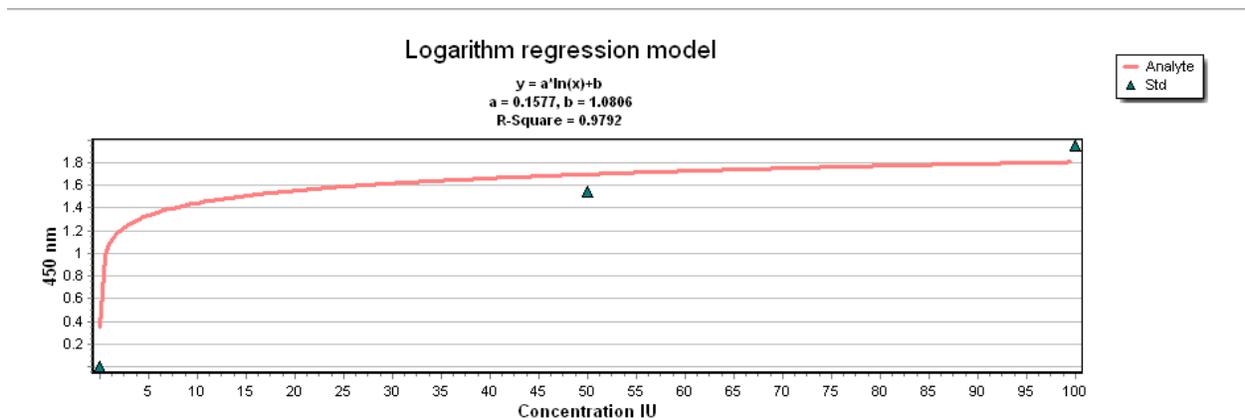
A piecewise linear function is a function defined on the set of points and is linear between each interval.



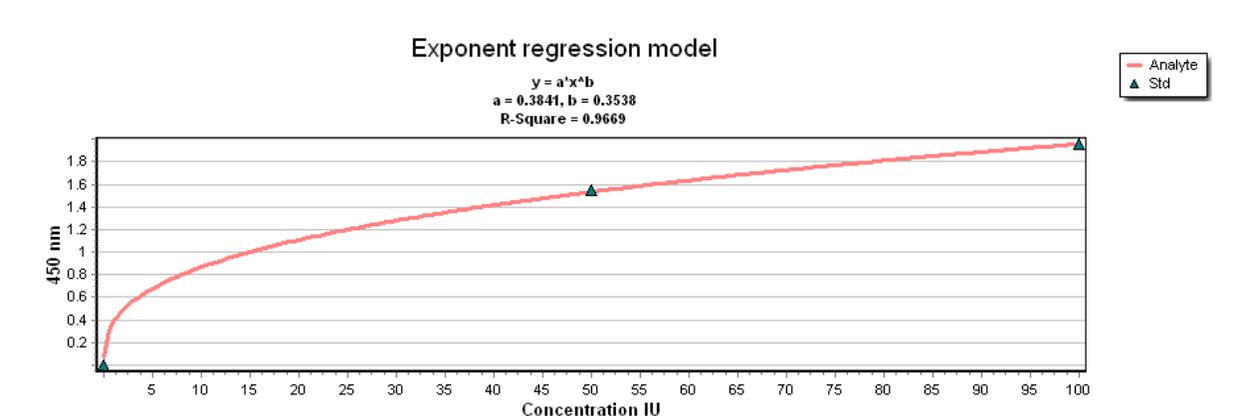
The index regression model



The logarith regression model



The exponent regression model



The cubic spline model

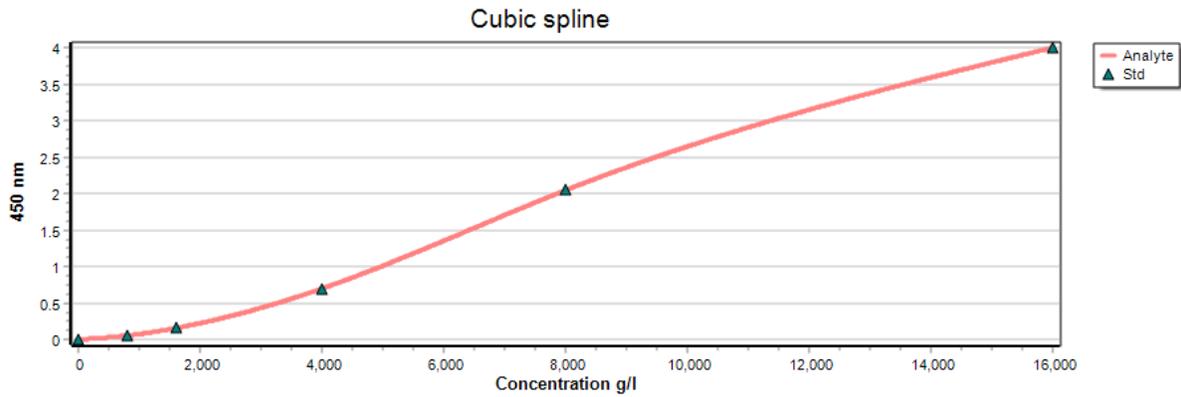


Chart [calibration] tab

| Standards | Given Concentration | Calculated Concentration | OD 450 nm | Residuals | %Recovery | Sample Name | Cell | Show |
|-----------|---------------------|--------------------------|-----------|-----------|-----------|-------------|------|-------------------------------------|
| S0 | 0 | *1.412 IU | 0.0001 | 1.411 | NA | Std S0 | A1 | <input checked="" type="checkbox"/> |
| S0 | 0 | 1.981 IU | 0.0014 | 1.980 | NA | Std S0 | A2 | <input checked="" type="checkbox"/> |
| S1 | 5 | 5.062 IU | 0.4005 | 0.062 | 101.244% | Std S1 | B1 | <input checked="" type="checkbox"/> |
| S1 | 5 | 4.826 IU | 0.3748 | -0.174 | 96.527% | Std S1 | B2 | <input checked="" type="checkbox"/> |
| S2 | 10 | 9.986 IU | 0.7626 | -0.014 | 99.860% | Std S2 | C1 | <input checked="" type="checkbox"/> |
| S2 | 10 | 9.918 IU | 0.7590 | -0.082 | 99.176% | Std S2 | C2 | <input checked="" type="checkbox"/> |
| S3 | 20 | 22.074 IU | 1.1766 | 2.074 | 110.372% | Std S3 | D1 | <input checked="" type="checkbox"/> |
| S3 | 20 | 20.849 IU | 1.1471 | 0.849 | 104.247% | Std S3 | D2 | <input checked="" type="checkbox"/> |
| S4 | 50 | 45.665 IU | 1.5478 | -4.335 | 91.330% | Std S4 | E1 | <input checked="" type="checkbox"/> |
| S4 | 50 | 45.007 IU | 1.5415 | -4.003 | 90.102% | Std S4 | E2 | <input checked="" type="checkbox"/> |

5 Parameter Logistics 1

$y = A + \frac{D(1+(X/C)^B)^E}{1+(X/C)^B}$

A = 19.1644, D = -19.1637, C = 2.4277, B = 18.6169, E = 0.0015

R-Square = 0.9991

Use Best Fit feature

5 Parameter Logistics 1

Set X-axis to log scale

Set Y-axis to log scale

Show samples

Allow Extrapolate

Recalculate

Here you can select a needed model, by removing the tick from the field "Use the best fit feature" Next: select a model from the list below.

Use Best Fit feature

5 Parameter Logistics 1

4 Parameter Logistics

5 Parameter Logistics 1

5 Parameter Logistics 2

Linear

Piecewise linear

Recalculate

Then click on the “Recalculate” button.

User can switch X, Y axis to log scale, as well as to show samples on the curve and enable/disable extrapolation (for last feature Recalculate button should be pressed).

User can export calibration data to .xls file.

Loading a standards curve

First you need to create your curve.

Open the program and load a quantitative experiment, like below:

QuantAssay v0.7.1.2

File Options

NEW LOAD XLS CSV PDF

bioan

Exp_1805_1034_0 test for quality test for quant

Available Devices Input Data View Results Chart

Name Smp Group 1 Reset Test Bkg P₁ N₁ Std X Load

| | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 |
|----------|------------------|------------------|----------------|----------------|-----------------|-----------------|--------|--------|--------|--------|--------------------|--------------------|
| A | Std S0 0.0080 | Std S0 0.0090 | Smp2 0.0080 | Smp2 0.0090 | Smp11 0.3407 | Smp14 0.4770 | 0.4293 | 1.6330 | 2.5920 | 4.3000 | Positive 1.0002 | Positive 1.0000 |
| B | Std S1 0.0600 | Std S1 0.0650 | Smp3 0.0600 | Smp3 0.0650 | Smp12 0.2839 | Smp15 0.3975 | 0.3577 | 1.3608 | 2.1600 | 3.6000 | Positive 1.0003 | Positive 1.0000 |
| C | Std S2 0.1600 | Std S2 0.1690 | Smp4 0.1600 | Smp4 0.1690 | Smp13 0.2366 | Smp16 0.3312 | 0.2981 | 2.0000 | 1.8000 | 3.0000 | Negativ 0.2002 | Negativ 0.2000 |
| D | Std S3 0.6900 | Std S3 0.7200 | Smp5 0.6900 | Smp5 0.7200 | 0.1183 | 0.6300 | 0.1491 | 0.5670 | 0.9000 | 1.5000 | Negativ 0.2001 | Negativ 0.1999 |
| E | Std S4 2.0000 | Std S4 2.1000 | Smp6 2.0000 | Smp6 2.1000 | 0.0592 | 0.3150 | 0.0745 | 0.2835 | 0.4500 | 0.7500 | Bkg 0.0001 | Bkg 0.0001 |
| F | Std S5 4.0001 | Std S5 4.0000 | Smp7 4.0001 | Smp7 4.0000 | 0.0296 | 0.1575 | 0.0373 | 0.1418 | 0.2250 | 0.3750 | Bkg 0.0001 | Bkg 0.0001 |
| G | 2.0001 | 2.0000 | 2.0001 | 2.0000 | 0.0148 | 0.0788 | 0.0186 | 0.0709 | 0.1125 | 0.1875 | Bkg 0.0001 | Bkg 0.0001 |
| H | Smp1 4.2000 | 0.2990 | 1.0000 | 1.0000 | 0.0074 | 0.0394 | 0.0093 | 0.0354 | 0.0563 | 0.0938 | Bkg 0.0001 | Bkg 0.0001 |

Choose an assay
test for quant

Choose a Template or Save as
Plate_23.01.2018 16:31

What to show in a cell:
A Sample Name
450 nm

| Cell | Name | Sample Name | Type |
|------|------|-------------|------|
| | | | |

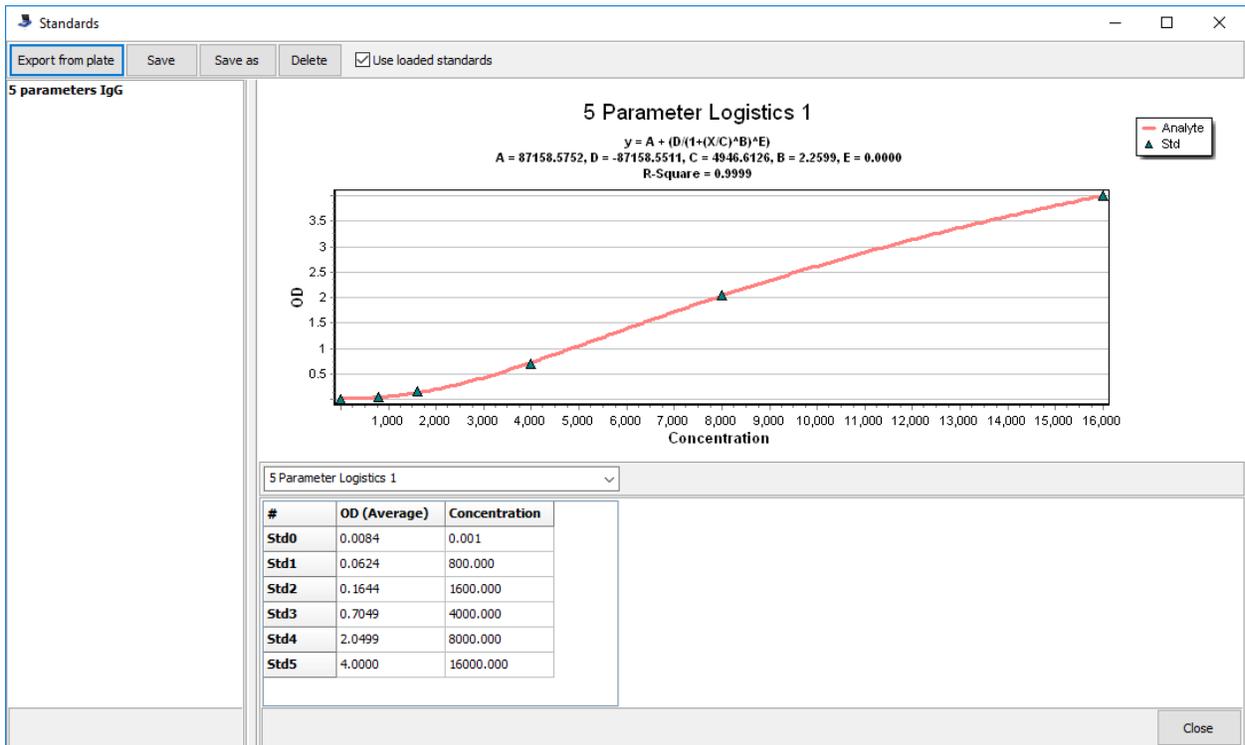
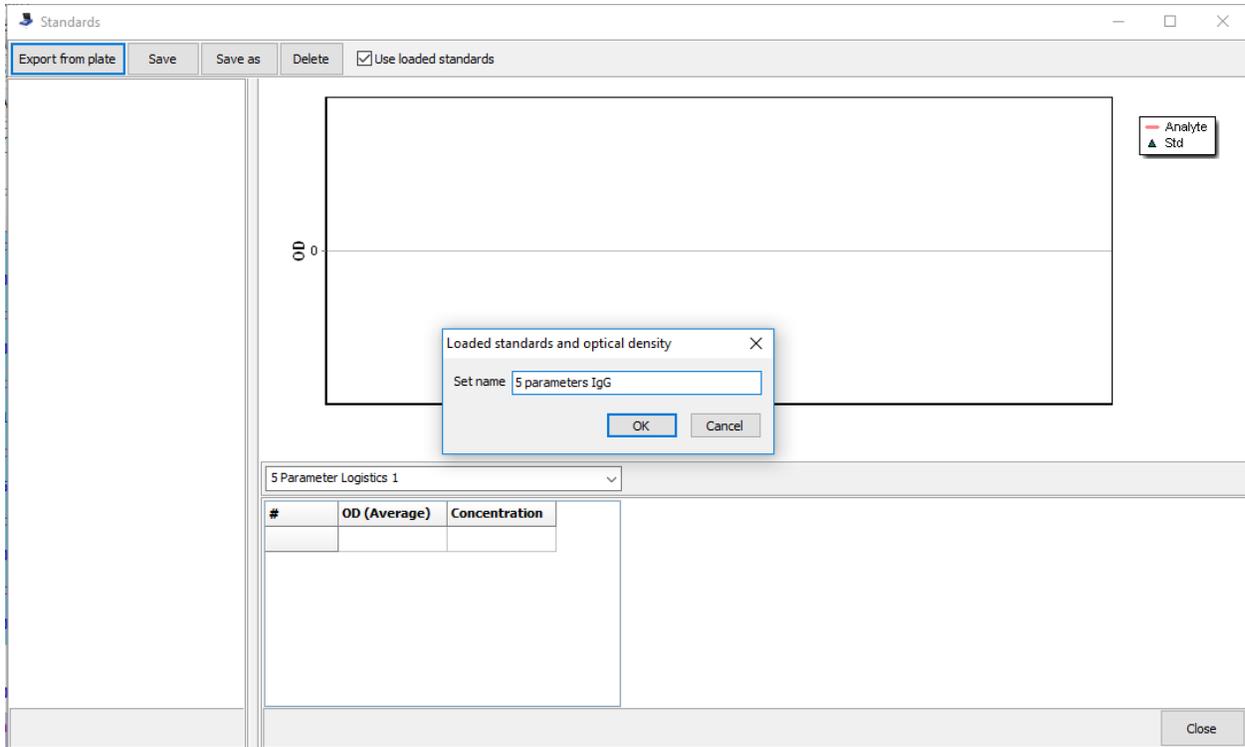
Calculate Main channel
450 nm

Kinetic Mode Panel

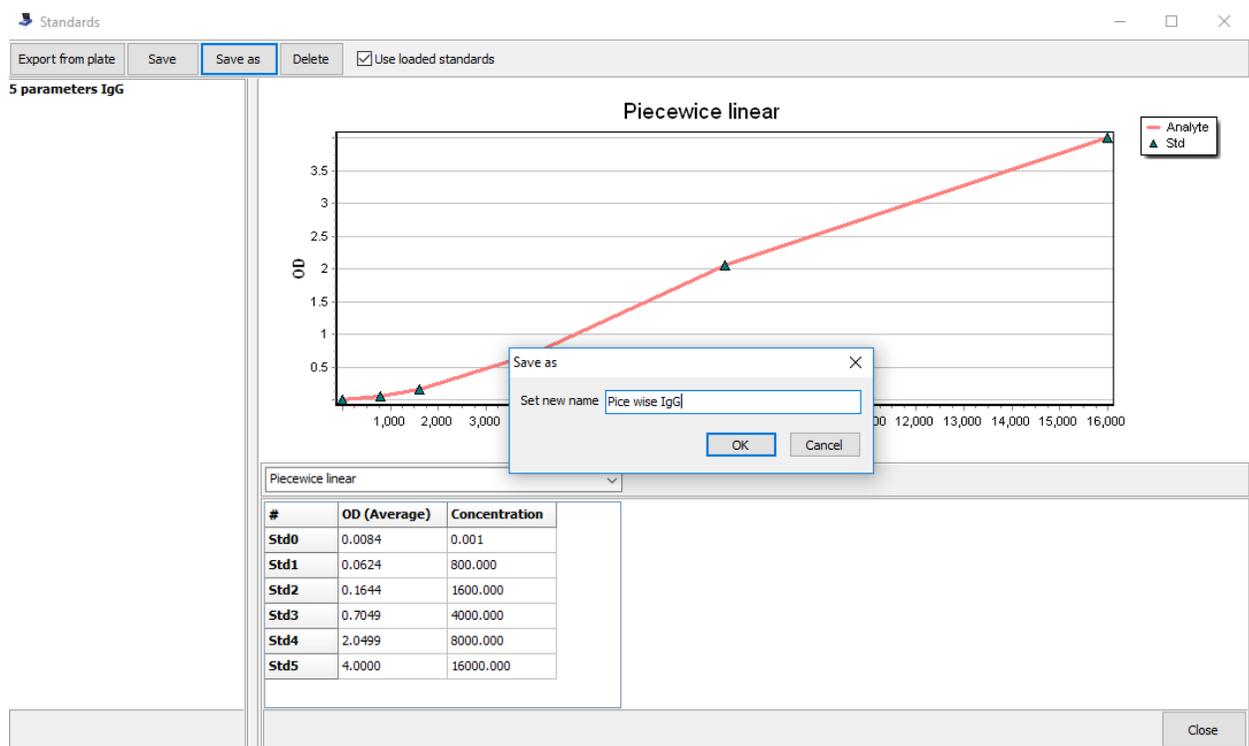
Press on load curve icon:



In this windows click on the Export from plate, Set the name of the curve and press Ok. The curve is now saved for the later use.



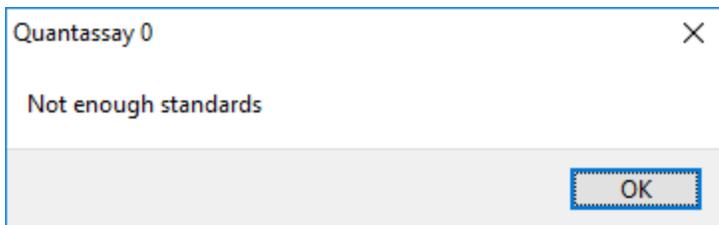
Here you can now set other type of curve and Save it as new curve.



Now you need to load it in the new experiment.

Set you samples and measure the plate.

You will get following message:

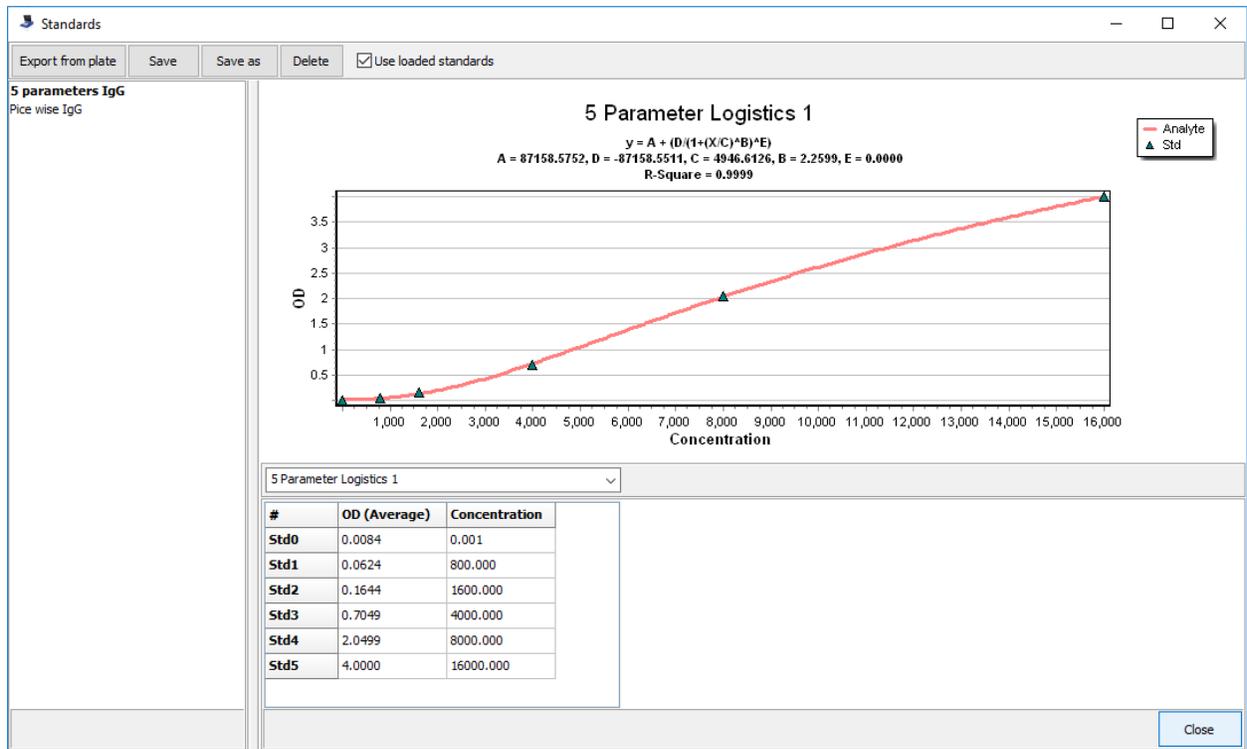


Press ok, go to Input Data tab.

Press on load curve icon:



Select the curve you need and close the windows, your results will be calculated:



If you do not want to use loaded curve, then go to Input Data: Load Curve, and disable "Use loaded standards" box. Use loaded standards

Results tab

| Available Devices | | Input Data | | | View Results | | | Chart | | | |
|-------------------|------|---------------------|-------|-----------|--------------|---------------------|--------------------|--------------------------|-----------|-------------------------|---------------------------|
| Cell | Type | Sample Name | Group | OD 450 nm | Result 1 | Given Concentration | Mean Concentration | Calculated Concentration | Mean (OD) | Standard Deviation (OD) | Coefficient Variation (%) |
| A1 | S0 | Std S0 | | 0.0001 | OK | 0 IU | *1.412 IU | *1.412 IU | 0.0008 | 0.0007 | 86.768% |
| A2 | S0 | Std S0 | | 0.0014 | OK | 0 IU | 1.412 IU | 1.981 IU | 0.0008 | 0.0007 | 86.768% |
| A3 | T1 | Smp1 | 1 | 1.9557 | Out of Range | | *103.001 IU | *103.350 IU | 1.9541 | 0.0017 | 0.086% |
| A4 | T1 | Smp1 | 1 | 1.9524 | Out of Range | | *103.001 IU | *102.653 IU | 1.9541 | 0.0017 | 0.086% |
| A5 | T9 | Smp9 | 9 | 0.0001 | Out of Range | | *1.412 IU | *1.412 IU | 0.0001 | 0.0000 | 0.000% |
| A6 | T9 | Smp9 | 9 | 0.0001 | Out of Range | | *1.412 IU | *1.412 IU | 0.0001 | 0.0000 | 0.000% |
| A7 | T17 | Smp17 | 17 | 0.0011 | In Range | | 2.030 IU | 1.922 IU | 0.0018 | 0.0007 | 36.541% |
| A8 | T17 | Smp17 | 17 | 0.0025 | In Range | | 2.030 IU | 2.086 IU | 0.0018 | 0.0007 | 36.541% |
| A9 | T25 | Smp25 | 25 | 4.1524 | Out of Range | | *14338.244 IU | *12074.361 IU | 4.2262 | 0.0738 | 1.747% |
| A10 | T25 | Smp25 | 25 | 4.3000 | Out of Range | | *14338.244 IU | *17041.094 IU | 4.2262 | 0.0738 | 1.747% |
| A11 | P1 | Positive control P1 | | 1.9738 | OK | | *104.028 IU | *107.207 IU | 1.9590 | 0.0148 | 0.758% |
| A12 | P1 | Positive control P1 | | 1.9441 | OK | | 104.028 IU | 100.946 IU | 1.9590 | 0.0148 | 0.758% |
| B1 | S1 | Std S1 | | 0.4005 | OK | 5 IU | 4.943 IU | 5.062 IU | 0.3876 | 0.0128 | 3.315% |
| B2 | S1 | Std S1 | | 0.3748 | OK | 5 IU | 4.943 IU | 4.826 IU | 0.3876 | 0.0128 | 3.315% |
| B3 | T2 | Smp2 | 2 | 1.9268 | In Range | | 100.098 IU | 97.461 IU | 1.9399 | 0.0132 | 0.680% |
| B4 | T2 | Smp2 | 2 | 1.9531 | In Range | | *100.098 IU | *102.808 IU | 1.9399 | 0.0132 | 0.680% |
| B5 | T10 | Smp10 | 10 | 0.0016 | In Range | | 2.123 IU | 2.002 IU | 0.0031 | 0.0015 | 49.231% |
| B6 | T10 | Smp10 | 10 | 0.0046 | In Range | | 2.123 IU | 2.184 IU | 0.0031 | 0.0015 | 49.231% |
| B7 | T18 | Smp18 | 18 | 0.0029 | In Range | | 2.095 IU | 2.113 IU | 0.0026 | 0.0003 | 11.518% |
| B8 | T18 | Smp18 | 18 | 0.0023 | In Range | | 2.095 IU | 2.074 IU | 0.0026 | 0.0003 | 11.518% |
| B9 | T26 | Smp26 | 26 | 0.0040 | In Range | | 2.115 IU | 2.162 IU | 0.0029 | 0.0011 | 35.684% |
| B10 | T26 | Smp26 | 26 | 0.0019 | In Range | | 2.115 IU | 2.040 IU | 0.0029 | 0.0011 | 35.684% |
| B11 | T34 | Smp34 | 34 | 0.0028 | In Range | | 2.133 IU | 2.109 IU | 0.0033 | 0.0005 | 13.847% |
| B12 | T34 | Smp34 | 34 | 0.0038 | In Range | | 2.133 IU | 2.153 IU | 0.0033 | 0.0005 | 13.847% |

This tab displays results in the following columns:

Cell #

Type

Sample name

Group

OD *** nm

Result 1 and 2

Give concentration (for quantitative assays, the blue font and * marked are extrapolated values)

Mean concentration (for quantitative assays, the blue font and * marked are extrapolated values)

Calculated concentration (for quantitative assays)

Mean OD

Standard deviation of OD (for samples repeats)

Coefficient of Variation of OD (for samples repeats)

For multiplex, avidity and qualitative assays columns relating to the concentration are not displayed.

For avidity methods column A/M indicates what sample was diluted with a dissociating agent (0 -- not diluted, 1 -- diluted)

For multiplex methods column A/M displays the group of the
Also the results table can be sorted by column or rows.

In order to output results in PDF, Excel and CSV click on a corresponding icon



LIS export

When the experiment is finished, click on the LIS export button to start.

Export builder

Save Cancel

Export settings

Filename: Exp_2301_1612_0_0 Extension: csv txt

Separator type: ; , TAB Other

Identifier types

Cell Standard Deviation (OD)
 Type Coefficient of Variation (OD)
 Sample Name Assay name
 A/M Conc. units
 Group
 OD 450 nm
 Result 1
 Result 2
 Given Concentration
 Mean Concentration (g/l)
 Calculated Concentration (g/l)
 Mean (OD) (g/l)

Include headers
 Rewrite file with same name

Check all Uncheck all

Export content

| | |
|---|--------------------------------|
| 1 | Type |
| 2 | Sample Name |
| 3 | OD 450 nm |
| 4 | Calculated Concentration (g/l) |
| 5 | Conc. units |
| 6 | Cell |

Select the file name extension for your data. You can choose either .csv or .txt format.

Choose the needed separator type:

Separator type

; , TAB Other

Select the identifiers (headers) you want to export.

Identifier types

| | |
|--|--|
| <input checked="" type="checkbox"/> Cell | <input type="checkbox"/> Standard Deviation (OD) |
| <input checked="" type="checkbox"/> Type | <input type="checkbox"/> Coefficient of Variation (OD) |
| <input checked="" type="checkbox"/> Sample Name | <input type="checkbox"/> Assay name |
| <input type="checkbox"/> A/M | <input checked="" type="checkbox"/> Conc. units |
| <input type="checkbox"/> Group | |
| <input checked="" type="checkbox"/> OD 450 nm | |
| <input type="checkbox"/> Result 1 | |
| <input type="checkbox"/> Result 2 | |
| <input type="checkbox"/> Given Concentration | |
| <input type="checkbox"/> Mean Concentration (g/l) | |
| <input checked="" type="checkbox"/> Calculated Concentration (g/l) | |
| <input type="checkbox"/> Mean (OD) (g/l) | |

Select if you want to export the header names

Include headers

Export content panel is visualizing the the exportable headers.

Export content

| | |
|----------|--------------------------------|
| | |
| 1 | Cell |
| 2 | Type |
| 3 | Sample Name |
| 4 | OD 450 nm |
| 5 | Calculated Concentration (g/l) |
| 6 | Conc. units |

The Rewrite file with same name checkbox will rewrite the file with same name without prompting confirmation from you.

Rewrite file with same name

When finished press on Save button and select the path for exporting.

Temporary saves

By default software autosaves each measurement.

Measurements can be found in “Documents/QuantAssay/Temporary saves”

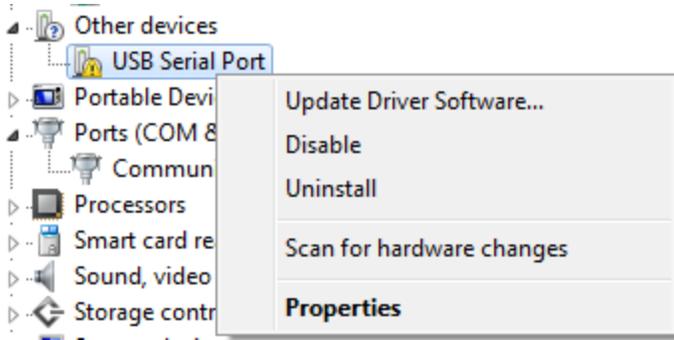
This feature autosaves up to 50 measurements, if you have 50 measurements already, then the earliest measurements from the list will be overwritten.

Troubleshooting

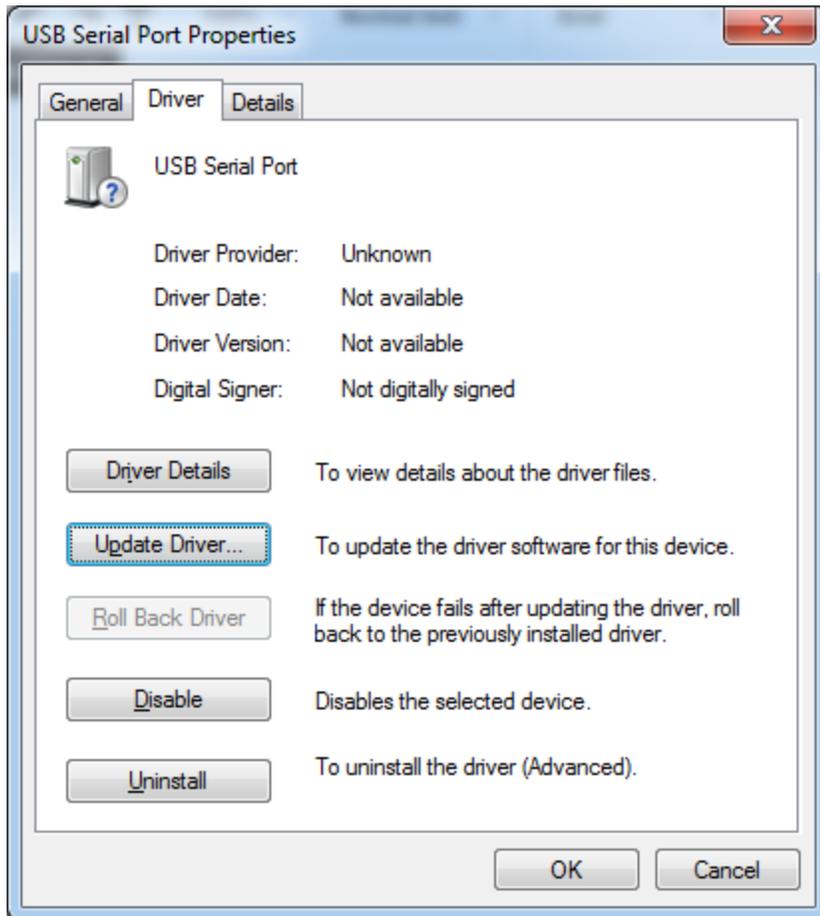
World practice shows that software vendors in the case of software malfunctioning indicate that user have accepted of the license agreement by which the software was provided as is or/and the shortcomings of operating system compatibility with the PC hardware, which leads to errors of or reduced productivity of the program. Unfortunately, we state that this practice model is the best for us and we have to stick to it. But, we would be grateful if you send captured errors to software@biosan.lv, so that we can identify the cause and, possibly, make the program better.

1. - The device can not connect to the computer.
 - 1.1 Check that the USB cable is firmly connected to the PC and to the instrument, try to eject and inster both ends.
 - 1.2 Try restarting your devices/software/computer, if it does not help, then reinstall the software.
 - 1.3 If the problem persists, go to the point 4 of this troubleshooting
2. - The program can not close, says that the experiment is still going, but I stopped it.
 - Try pressing the Play button on the toolbar (to start the experiment), and click on the Stop button, wait 5 seconds and then try to close it. If this does not work, open Task Manager (Ctrl + Shift + Esc) and close all processes “quantassay.exe”
3. - Device does not respond to the program
 - Try to turn on or off the device, if necessary, try to plug it off and then on.
4. - Drivers can not be installed
 - 1. Try to give administrator rights to the user who installs the program
 - 2. If previous step did not help, try the following:
Go to Control Panel/Device Manager
Expand the Other devices line:





Click on Properties, Select Driver tab, click on the Update Driver:



Select Search automatically for updated driver software (you should have internet connection on)

- ➔ **Search automatically for updated driver software**
Windows will search your computer and the Internet for the latest driver software for your device, unless you've disabled this feature in your device installation settings.

After installing the driver, following message should appear:

Windows has successfully updated your driver software

Windows has finished installing the driver software for this device:



If you do not have internet connection, please use computer with internet to download latest drivers from: <http://www.ftdichip.com/Drivers/VCP.htm> under “Available as setup executable”. Then transfer and install the driver on the computer where the units are connected.

If it did not help, please try to find a solution here:
<https://support.microsoft.com/en-us/help/2654149/error-usb-device-not-recognized-when-you-try-to-access-a-usb-external-hard-drive>.

If that did not help, try to connect the instrument to the different computer, to check if the problem is on the computer or instrument end.

Also try to change the USB cable.

5. - My problem is not described here.
-While working with the program a problem that is not described here may arise. There is an universal solution: reboot the computer and / or reinstall the program. Check the website for the latest software updates
6. - I have connected the devices to USB 2.0 SS (super speed) terminals and my computer shuts down constantly, indicating that there is some sort of an error with FTDI driver.
- Please avoid connecting to USB 2.0 SS, connect the devices to the standard USB 2.0.
Data for connecting to USB 3.0 terminals is yet not available.

Disclaimer

1. Program is provided "as is", as it was stated in the license agreement.
2. This documentation may not coincide with the latest version of the program. In this case, we are sorry and hope for your understanding, and we would be grateful if you could point us the inconsistencies, also we hope that the interface is really intuitive and does not require thorough explanation. In any case, write to software@biosan.lv , we will be happy to guide you!