

Analyzing Experiments Repeatedly

This article includes the following techniques:

- Linking analyses
- Column math analysis
- User-defined transforms
- Using "Info Constants"
- Creating a using templates
- Saving and applying a method
- Applying a method by example

If you must perform the same analytical and graphing tasks many times, varying only the input data, Prism helps with a number of tools. Scripting, a means of automating data input and output via a simple programming language, is discussed in the Step-by-Step Example "Creating and Running a Prism Script". Three manual techniques are covered in this article:

- <u>Templates</u> allow you to duplicate an entire project in one step. You create the template the first time you set up the project, but store it without the data. To replicate your work, open the template, add the new data, and store the file under a new name. Or insert the template into another file.
- <u>Duplicating a family of sheets</u> replicates your work within the same project file. Simply duplicate a sheet and all sheets linked to it—formatted graphs, analyses, info, and layouts—using a single Prism command. Then replace the input data for the duplicate family.
- <u>Applying a method</u> effectively duplicates a family of sheets, but in this case, the replicated work can come from the same project or a different project. Add a new input data table, then either (a) apply a *saved* method or (b) point to another table in the project to apply *by example* all sheets linked to the latter.

We'll demonstrate these three techniques as applied to a family of linked sheets that produce a simple biochemical activity computation. To make it easier to see the linkage between sheets in the Navigator, open the **Edit** menu and choose **Preferences...** In the Preferences dialog, select the **View** tab, and then activate the **Data+Results view**.

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¹ Adapted from: Miller, J.R., *GraphPad Prism Version 4.0 Step-by-Step Examples*, GraphPad Software Inc., San Diego CA, 2003. *Step-by-Step Examples* is one of four manuals included with Prism 4. All are available for download as PDF files at <u>www.graphpad.com</u>. While the directions and figures match the Windows version of Prism 4, all examples can be applied to Apple Macintosh systems with little adaptation. We encourage you to print this article and read it at your computer, trying each step as you go. Before you start, use Prism's **View** menu to make sure that the Navigator and all optional toolbars are displayed on your computer.

Data+Results view	
 Always show, as well as individual Data and Results folder 	s
O Always show, instead of individual Data and Results folder	s
O Never show	
Show only for large projects	

Quantifying Uptake into Cells

Suppose you want to measure the effects of various inhibitors on uptake of a radiolabeled substrate into suspended cells. You have developed a method for removing aliquots at two times within the "linear" phase of uptake, quenching uptake in the aliquots, and measuring radioactivity in the cell fraction. You are testing a number of inhibitors on the same day using the same cell preparation.

Creating the Data Table

The variables recorded for each uptake measurement are

- Radioactivity at the beginning of the uptake interval (CPM_{Time 1})
- Radioactivity at the end of the uptake interval (CPM_{Time 2})

Their values will be recorded on the data table.

In the Prism Welcome dialog (if Prism is already running, choose **File... New project...**. Make the settings shown below.



Enter the data on the resulting table. The X column holds test article names. One test article is evaluated on each line, with column A and column B holding duplicate radioactivity measurements made 3 minutes apart, at 10 sec and 190 sec after initiation of uptake, respectively.

	X Labels	А		В	
	Test article	CPM Time 1		CPM 1	Time 2
	Х	A:Y1	A:Y2	B:Y1	B:Y2
1	Vehicle control	4292	4334	109152	108930
2	GP-10838	4228	4285	107590	109244
3	GP-10839	4122	4027	65542	66425
4	GP-10842	3588	3511	24518	26533
5	GP-10843	4085	1463	68872	69244

Type a name for this table in the box on the toolbar.

Input data	~

Creating an Info sheet

Variables whose values change from experiment to experiment, but are constant within each experiment, are

- Specific radioactivity of the uptake substrate (cpm/nmol)
- Time interval over which uptake is measured (min)
- Dry weight of cells in each aliquot (mg)

These variables will be recorded in Prism as Info Constants.

From the data table, click the **New** button and choose **New Info...**. In the **Create New Info** dialog, choose to link the Info page to the data table (Input data) we just created.

In the **Constant** column of the Info page, make new entries for specific activity, uptake measurement interval, and dry weight of the cells in one aliquot. Wording isn't critical—just use terms you'll recognize later.

Constant	Value
Experiment Date	Jan-27-2003
Experiment ID	
Notebook ID	
Project	
Experimenter	
Protocol	
SA (cpm/nmol)	511
Time (min)	3
Dry weight (mg)	6.3

In the **Value** column, enter the numbers shown. These are values are common to all measurements represented on the data table (Raw CPM). They will be used to perform the conversion and normalization of raw CPM values to uptake in nmol substrate/min/mg dry weight.

Computing Uptake

We will compute uptake as

$$\frac{CPM_{_{Time\,2}} - CPM_{_{Time\,1}}}{Specific \ Activity \cdot Uptake \ Interval \cdot Dry \ Weight}$$

We'll compute this quantity for each pair of CPM measurements (CPM at 10 sec and CPM at 190 sec) on the original data table using two linked analyses. We'll compute $CPM_{Time 2} - CPM_{Time 1}$ using a Column Math analysis, then divide the results by the values for specific activity (SA in cpm/nmol), uptake interval (Time in min), and dry weight (in mg) found on the Info page using a User-defined Y transform.

Column Math Analysis

From the data table, click the **Analyze** button. In the **Analyze Data** dialog, choose **Remove baseline and** column math from the **Data manipulations** list.

Prism performs a **Column Math** analysis by computing the differences between values in pairs of adjacent columns (A and B, C and D, etc.) all the way across a data table or Results sheet, ignoring empty cells, and placing the differences on a new Results sheet. In the Parameters dialog, indicate that the subtrahends (numbers to *be*

subtracted, also termed "baseline values") are in column A. This means that Prism is to subtract values in columns A, C, E... from values in B, C, F... respectively.

Complete the selections in the Parameters dialog as follows:

Parameters: Remove Baseline and Column Math		
Where are baseline values?	Calculate	
🔿 Column 💽	 Difference: Value - Baseline 	
O Data sets (columns) B, D, F	◯ Sum: Value + Baseline	
 Data sets (columns) A, C, E First row Last row Mean of first 3 rows. Mean of last 3 rows. Erase those rows from the analysis Chinear baseline Assume the baseline is linear with X. Use the values predicted from the regression line. 		
 Don't assume linear baseline. Use the the second sec	cond replicate of the baseline only with the second replicate the baseline replicates, and use the average with each	
New graph		
Create a new graph of the results		
	Help Me Decide Cancel OK	

When you leave the dialog, Prism creates a Results table to hold the differences. Note that the distinction between duplicate determinations has been maintained, and Prism has treated the replicates as "matched" per our setting in the Parameters dialog.

Test article	B-A		
Х	A:Y1	A:Y2	
Vehicle control	104860.000	104596.000	
GP-10838	103362.000	104959.000	
GP-10839	61420.000	62398.000	
GP-10842	20930.000	23022.000	
GP-10843	64787.000	67781.000	

Name this Results sheet:

1 🗸

User-Defined Transform

From this Results sheet, click **Analyze**. In the **Analyze Data** dialog, choose **Transforms** from the **Data manipulations** list.

In the **Parameters: Transforms** dialog, indicate that you will be using a user-defined Y transform.

Function List	User-defined Y functions	*

Click the **Add...** button to open the **Equation** dialog. Give your transform a name, and enter the transform equation in the **Equation** box, as shown below.

Name:	Adjustment for spec act, time, weight
- Four	tion
E qua	don
Y=Y,	/SpecAct/Time/Weight

The parameters "SpecAct", "Time", and "Weight", on the right-hand side of the equation are arbitrary—use terms that make sense to you. We'll specify their values in the next dialog. Click **OK** to return to the **Parameters: Transforms** dialog.

Prism analyzes the user-defined transform and displays the three parameters that it finds. Click in the value box next to SPECACT, and choose **SA (cpm/nmol)** from the **Link and Info Constant** dialog. Fill in the values for TIME and WEIGHT similarly.

Parameters	Value	Value
SPECACT	(511.0)	Link an Info Constant 🛛 🛛 🛛
TIME	(3.0)	[No linked constant]
WEIGHT	(6.3)	Input data info
Replicates O Transform	individual Y values	Time (min) = 3.0 Dry weight (mg) = 6.3

Choose to **Transform the average of replicates** and check the box to **Create a new graph of the results**. A new Results sheet appears with the final transformed values. You can change the heading for column A to increase readability.

Test article	nmol/min/mg
Х	Y
Vehicle control	10.844
GP-10838	10.785
GP-10839	6.410
GP-10842	2.275
GP-10843	6.863

Rename the Results sheet:

nmol/min/mg 🛛 🔽

Instead of applying one user-defined transform, you could do three successive divisions using the **Y=Y/K** transform from the **Standard functions** list, filling in the value for K from the **Link an Info Constant** dialog each time. That would avoid having to enter the user-defined transform and would allow you to see all intermediate values, but would also clutter the project with additional Results sheets.

Click the **Graphs** tab to view the graph of the transformed data. In the figure below, some additional editing has been done. You may be particularly interested in showing the baseline labels at an angle—double-click on the baseline to open the **Format Axes** dialog, make sure the **X axis** tab is selected, and choose angled labeling.



Templates

Using a Template Alone

The project file we just created will the basis for our template. Before saving a template, it's usually a good idea although not strictly necessary—to erase any data on the table *that will change during subsequent use of the template*.

Delete the values for specific activity, time, and dry weight on the Info page if you wish. This is not mandatory.

Switch to the data table ("Input data"), select all data below the column headings.

	X Labels	А		В	
	Test article	CPM Time 1		CPM Time 2	
	Х	A:Y1 A:Y2		B:Y1	B:Y2
1	Vehicle control	4292	4334	109152	108930
2	GP-10838	4228	4285	107590	109244
3	GP-10839	4122	4027	65542	66425
4	GP-10842	3588	3511	24518	26533
5	GP-10843	4085	1463	68872	69244
6					

Press the **Delete** *key*—don't use the **Delete** command on the **Edit** menu; you want to maintain the data cells, but without the data. If you see a message that tells you about the data being linked to the graph, disregard it and click **OK**.

Don't switch to the graph at this point. That could break the link to the graph and cause the template not to work correctly.

Place the cursor in the top cell (row 1) of the X column, then select File... Save Special... Save Template.

Fill out the dialog box as shown below. Note that you may indicate here whether you wish the template to be stored on your computer or on another computer networked to it. You may also generate different categories under which templates are stored. The description at the bottom of the dialog is optional; this message will be displayed as an aid to any user opening the template.

Save Template	
Save the template on: The network server, for all Prism On my hard drive	n users to share
Choose a category:	
Sample templates	Add New Category
Choose a name:	
 Create new template 	Cell uptake
Overwrite an existing template	Enzyme kinetics template
Create the template from: This data table and its info sheel The entire project	ets, analyses and graphs
Description and/or instructions (optio	nal):
Enter test article names (CoLX) and I Time 2 (CoLB). Enter identifying info, weight (mg) on info sheet. View upta "nmol/min/mg".	CPMs in duplicate at Time 1 (Col A) and specific activity (cpm/nmol), time (min), cell ske normalized to cell weight on sheet
	Help Cancel OK

Now close the file, and we'll see how to use the template.

Choose **File...New Project** to open the Welcome dialog. Set the radio button for a **Template** and select the template that you just created.

Welcome to Graph	ıPad Prism	
GraphPad	M	Version 4.00.228 January, 14, 2003
To start: 💿 Creat	e a new project 🛛 🔿 Open an exi	sting file
Choose: 🔿 Type	of graph 🛛 🔘 Format of data table	e 💽 Template
Template category:	Sample templates	✓
Cell uptake Enzyme kinetics temp Exam score histogram Linear time course Protein Assay std. cu RIA t test	olate n rve	

Click **OK**, and you'll get a new project formatted as your template. If you created a description, you will see it now. When you close the description box, the cursor should be at the top of the X column (note that the column headings are still there, if you didn't delete them before you saved the template).

Template description and/or instructions
Template:C:\PROGRAM FILES\GRAPHPAD\PRISM 4\TEMPLATES\SAMPLE
Enter test article names (CoIX) and CPMs in duplicate at Time 1 (CoIA) and Time 2 (CoIB). Enter identifying info, specific activity (cpm/nmol), time (min), cell weight (mg) on info sheet. View uptake normalized to cell weight on sheet "nmol/min/mg".
ОК

You can also insert a template into a project that you have open. With any data table in view, click the **New** button and then choose **New Data Table (+ Graph)...** Choose **Insert template** and proceed as described above. Now enter some new data:

	X Labels	А		В	
	Test article	CPM Time 1		CPM Time 2	
	Х	A:Y1 A:Y2		B:Y1	B:Y2
1	Vehicle control	6378	6539	162140	163934
2	GP-10844	5936	5877	134951	133710
3	GP-10845	5216	5455	130946	131687
4	GP-10846	5819	5710	91859	93122
5	GP-10848	4938	4897	36834	39886

Switch to the Info page and fill in or replace the values in the last three rows.

SA (cpm/nmol)	642
Time (min)	3
Dry weight (mg)	7.4

Now switch to the Results page named nmol/min/mg to view the automatically updated uptake values,

	X Labels	Α
	Test article	nmol/min/mg
	Х	Y
1	Vehicle control	10.986
2	GP-10844	9.011
3	GP-10845	8.839
4	GP-10846	6.085
5	GP-10848	2.346

then click on the Graphs tab to view the graph:



When you save your work, remember to use the File... Save As... command and give the file a new name.

Inserting a Template into Another Project

We have just covered one way to use a template—alone, opening the template with its empty table, filling the table with data to produce updated results, and storing the file under a new name.

You may also insert a template into an existing file. Click the **New** button, then choose **New Data Table** (+Graph).... In the Create New Table dialog, choose Insert template, change the template category if necessary, and select the template. The contents of the template are merged into the open file.

Applying a Method

If you use the same treatment (analyses or graphs, or both) with various data tables, try applying a method.

Saved Method

When you *save* a method, you can apply it later within the same project file or to another project file. From a data table that is linked to analyses and/or graphs that you wish to reproduce (you may use the template that we've created in this example), click **File... Save Special... Save Method**. Make entries in the **Save Method** dialog as shown below, and then click **OK** to save the method.

Sa	we Method		×
	Choose a category:		
	Sample methods		
		Add New Category	
	Choose a name:		~
	 Create new method 	Cell uptake	
	Overwrite an existing method	Enzyme kinetics	
	Description and/or instructions (optio	nal):	
	Method for converting CPM at Time	1 (Col A) and Time 2 (Col B) to uptake in	
	activity, time, and cell weight on info	page as necessary.	
		2	

To use the method, start with the project file created earlier in this article (see "Quantifying Uptake into Cells" earlier in this article) or a similar file. Add a new table, with new input data, and rename each data table so that they can be distinguished easily. The organization of the project file might look like this:



With the new table selected, click **Analyze**. In the **Analyze Data** dialog, choose to use a saved method, then select the category and specific method. If desired, enter a description of the method and a text string to be prefixed to the names of each newly generated sheet.

Analyze Data	×
Analysis	
🔘 Built-in analysis.	
 Use saved method. 	
O Method by example. Analyze and graph the same as another table in this project.	
Select a method:	
Choose a category Choose a method	
Sample methods Cell uptake	
Enzyme kinetics	
Exam score histogram	
Description	
Method for converting CPM at Time 1 (CoI A) and Time 2 (CoI B) to uptake in nmol substrate/min/mg and graphing results. After applying method, edit specific activity, time, and cell weight on info page as necessary.	
Prefix sheet names with *Sept 9 Help Cancel OK	j

Click **OK** to apply the method to the data table you selected earlier. Prism applies the analyses and generates a new graph based on the new data.



If any of the information is to be amended on the info page (including the date), do that now, then display the appropriate Results sheets and graph to verify that the results have been updated.

Method by Example

If you wish to apply to a data table work that is associated with another data table in the *same project*, you can apply that work by example. Let's return to the project file in the previous section as it appeared before applying the saved method (at that point we had just created the new data table "Sept 9 input data" and entered new data on that table.



Click **Analyze**, then choose **Method by example**. Select the data table to which the method (analyses and graph) is linked, i.e., indicate the desired method by pointing to an example of its application. At the bottom of the dialog, provide a distinguishing prefix for the new sheet names if desired.

Analyze Data		×
Analysis		
O Built-in an	alysis. d method.	
 Method by 	y example. Analyze and graph the same as another table in this project.	
	Select the table to use	
	CPM2-CPM1:Subtract baseline data	
	Input data	
Prefix sheet nan	nes with *Sept 9 Help Cancel OK	

When you exit this dialog, Prism applies the analyses and creates a new graph, just as it did when the saved method was applied.

🗁 Project
🚊 🧰 Data with Results
🖨 🔳 Input data
🖌 🖌nmol/min/mg
🖻 🏢 Sept 9 input data
🛛 🛃*Sept 9 CPM2-CPM1
f]*Sept 9 nmol/min/mg
🗉 🧰 Data Tables
🖨 🧰 Info
🕤 🚹 Project info 1
🕤 🚹 Input data info
🕤 🚹 *Sept 9 Input data info
🖻 🧰 Results
🖨 🧰 Graphs
🔤 🔟 nmol/min/mg graph
🛄 Sept 9 input data graph
🛄 *Sept 9 nmol/min/mg graph
- Cayouts