



Offline Flow Cytometry Analysis Software: Importance of Scaling and Data Visualization

Objectives

- Learn about the flow cytometry open platform data structure and how it facilitates for robust offline analysis
- Learn the importance of data scaling and display features in ensuring your data looks consistent with the native environment
- See specific examples comparing the native display to an offline data analysis software across a variety of platforms and manufacturers

Reasons for Considering Offline Analysis

The analysis software that is provided with the flow cytometer provides the basic features for the most common situations. Offline analysis packages offer specialized features and capabilities that significantly improve the analysis experience or the workflow in the clinical laboratory.

For example, using software listed with the FDA could be beneficial for your documentation requirements. Or consider the effort required to set up LIS connection to a single software package versus adapting and validating connections to multiple software packages. Software features that allow you to establish master report layouts that can be applied your analysis files can help to establish standardized reporting and save technician's time in creating reports. Robust QC reporting capabilities may be important to maintain the needed instrument and assay verification records and streamline audit preparation activities.

Standardized File Format

The ability of the field of flow cytometry to benefit from third party providers of specialized analysis software comes from the decision to standardize the data acquisition file format. In 1984 a standard format for flow cytometry data was proposed. A Data File Standards Committee of the International Society for Analytical Cytology (ISAC) was formed to maintain the standard, called FCS for flow cytometry standard. FCS3.1 is the current standard and has been adopted by most instrument manufacturers.

The raw data file from a flow cytometer is termed list mode. This refers to the fact that the parameter data collected on each event is stored in the order that the event passed through the interrogation point. The FCS standard describes the format that this data is stored into the computer file.

FCS files must have four major segments including HEADER, TEXT, DATA and ANALYSIS. The header section identifies the FCS standard used and outlines the how to find the other sections in the file. The TEXT section contains keywords and keyword values, describing the data set. These include user defined values. The DATA section contains the raw data. An optional ANALYSIS section includes the results of any data processing completed at the time of acquisition.

Offline analysis software takes advantage of this structured file to define the method for loading data regardless of the instrument used for acquisition.

Visualizing Data

Each instrument manufacturer will have settings to scale the data for display during acquisition. Prior to FCS 3.1 data scaling was controlled by the file. With FCS 3.1 data scaling is now controlled by the software doing the analysis. To ensure that the data looks the same in the analysis software as it did during the acquisition, the instrument default preferences need to be applied.

FCS contains a keyword to identify the cytometer used for acquisition. Analysis software can read this keyword, \$CYT, and apply preferences that control the display of data and information. When loading the first file from a given instrument, the preferences for that instrument are established. These can then be refined by the user as needed.

The following example shows the instrument default settings that are available for adjustment in Kalzua C Analysis Software.

Kaluxa C Software Options and Defaults

The left list shows the all loaded names of the acquisition instrument. And each one contains its own Cytometer setting information.

The settings are described as below:

Options and Default	5						x	
Options Defau	lts SCYT Defau	lts						
Defaults affect only newly created plots or data files.								
<pre> control y</pre>								
Gallios	Cytometer Identification SCYT value: Cytometer Identification							
CR 600	SCYT value: <	Default>						
Navios	Parameter Sca	le						
		Default	Min	Max	Decades	Width		
	Linear	V	0	1048576	N/A	N/A	=	
	Log	V	105	1048576	4	N/A		
	Logicle	V	-346	1048576	4.5	0.1		
G	Parameter Priority of signa		s: Area 🔹 He	ight ▼ Width	•			
							- 11	
	Custom data set	t identifier:	=DATASETN	AME()				
Θ	Scale Mode						- 11	
	Scale mode: F	ull Range 🔻					-	
-					R	estore initial se	ettings	
		ОК	Cancel	Apply				

A **\$CYT value,** which is the instrument identifier.

B Parameter Scale grid, it is used to set the default axis display setting.

There are 3 axis types in our Kaluza, "Linear" "Log" and "Logicle". The grid has 3 rows to let the user set display setting for each kind of axis. The "Min" column is the displayed minimum vaue for an axis, and "Max" column is the displayed maximum value for an axis. The "Decades" is only used in Log or Logicle axis. The "Width" is the **Negative Percentage** value for Logicle axis. Use the axis setting from FCS file. If the "Default" column check-box is checked, the Kaluza would work the same as Kaluza 1.5a. User could change the settings by unchecking the "Default" check-box, and input the valid expected value in the corresponding cell of grid.

🕒 Parameter

The priority of signal measurement (Area, Height, Width). The first signal measurement listed must be the one with the highest resolution on the acquisition instrument. When there are multiple measurements captured for a given PMT, Kaluza C uses this order to determine which parameters to use for setting up Compensations and which plots are displayed when Add All Plots is selected.

Data Set Identifier

The default formula used to identify Data Sets when loading new data files(*.lmd, *.fcs). The Data Set Identifier is used in the Data Set column in the Analysis List, as well as anywhere the Data Set name is shown throughout the software.

NOTE The formula may include keywords from the FCS header information, or other custom content.

🔁 Scale Mode

Configures the default plot axis scale:

- Full Range scale would make the axis scale to real event data.
- Legacy scale would make the axis scale to maximum 1024, same as Kaluza 1.3 or before.
- Astrios' scale changes several Kaluza defaults to provide the means to recreate plots that were created in the Astrios acquisition software.
- **NOTE** Astrios is a Scale Mode option only when Astrios is listed as an available instrument on the left side of this window.

*Astrios is for Research Use Only. Not for use in diagnostic procedures.

Options and Defaults X						
Options Defaults SCYT Defaults						
Defaults affect only newly created plots or data files.						
<pre>Custom data set identifier: =DATASETNAME()</pre>						
Gallios CR 600 Scale Mode						
Navios Scale mode: Full Range -						
Quadrant Names						
✓ Use default						
Base name: Default Increment way: Alphabet -						
Postfix name: -+ v						
-+ ++						
+-						
Statistics						
Absolute count calibration keyword: @ABSCALFACTOR						
Sample ID keyword: @SAMPLEID1						
Restore initial settings						
OK Cancel Apply						

🕞 Quadrant Names

The Quadrant Names section allows you to update the naming settings for Quadrant gates. Here, you can choose to use the default gate names or customize the components of Quadrant gate names, including the following:

Base Name: The name of the quadrant begins with the Base Name. The Base Name is the same in every quadrant.

Increment Way: How names are incremented. Options include: Alphabet/Number

G Statistics

Allows you to set/change the Absolute Count calibration keyword.

NOTE The Absolute Count keyword defines the calibration factor of the beads used with this sample if Flow-Count or similar absolute calibration particles are used. If the calibration factor is not provided, this keyword contains "NOT SET".

🕒 LIS

Allows you to set/change the Laboratory Information System (LIS) Sample ID Keyword.

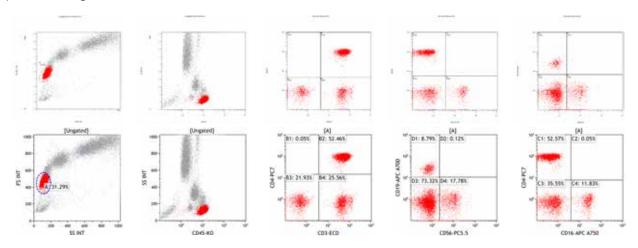
Data Examples

In order to compare data visualization of Kaluza C, software files from various instruments were collected. We obtained the resulting FCS files from the acquisition, reports documenting the user defined plots, indicating the native scaling, and statistics. The FCS files were then loaded into Kaluza C and the same plots and statistics were generated.

Presented plots and values are for illustrative purposes only and are based on customer reported data. Some information has been modified to simplify the presentation. Individual results may vary.

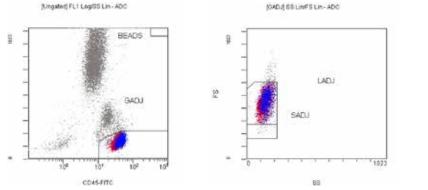
Navios EX Flow Cytometer (Beckman Coulter)

Top row shows Navios EX data analyzed with Navios System software. Bottom row shows the same plots generated within Kaluza C Analysis software after importing the FCS file. Small variations in population statistics is due to the placement of gate A.

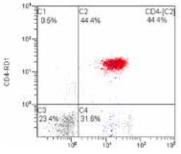


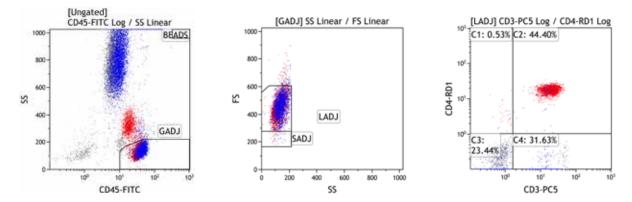
FC 500 Flow Cytometer (Beckman Coulter)

Top row shows FC 500 data analyzed with MXP Acquisition and Analysis software. Bottom row shows the same plots generated within Kaluza C Analysis software after importing the FCS file.



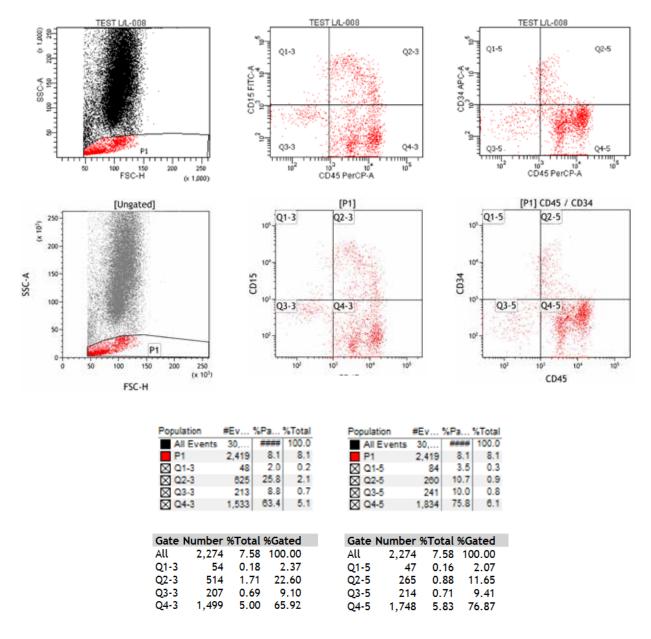
(GADJ AND LADJ) FLA Log/FL2 Log-ADC





FACSCanto (BD Biosciences)

The first row shows FACSCanto data analyzed with DIVA software. The second row shows the same plots generated within Kaluza C after importing the FCS file. Row 3 shows the statistics generated from the DIVA software and row 4 shows the statistics generated from Kaluza C Analysis software. Variations are due to the placement of gate P1.



Conclusion

Flow cytometry acquisition software, typically provided with the instrument by the manufacturer, provides the basic features for the most common situations, however offline analysis packages offer specialized features and capabilities that significantly improve the analysis experience depending on the needs of the laboratory. Industry standards for flow cytometry data files have facilitated the development of offline packages that are optimized for different types of user needs. In choosing a package users will want to ensure that the data is scaled using the same criteria as specified in the acquisition software. Generated plots will look the same in both the acquisition and analysis software, streamlining the workflow. In this paper, we demonstrated how the Kaluza C Analysis offline software package was able to read files from a variety of instruments and suppliers. The data was compared to plots and statistics generated in the native package and provided side-by-side comparisons to compare fidelity. The software will read the parameters from the FCS files and visualize data per the instructions from the manufacturer. An interface with options to adjust these parameters is provided, giving the user exquisite control over their data visualization.



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