BD LSRFortessa Standard Operation Protocol

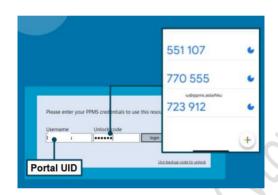
A. Log in the Window and Tracker

1. Press Ctrl+Alt+Delete

2. Account: User

Password: posted under the bottom of the computer screen

3. Log in Tracker





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B. Log in the BD FACSDiva Software

- Key in username and password and click OK to log in.
 *Please contact the Technical Staff in charge if you do not have an account.
 **Please contact the Technical Staff in charge if you forget password.

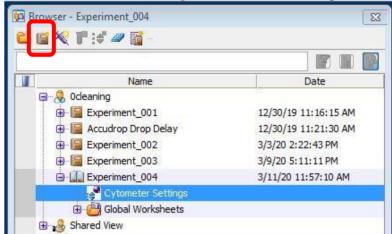
2. Click Use CST Settings when pop-up message as below is shown.



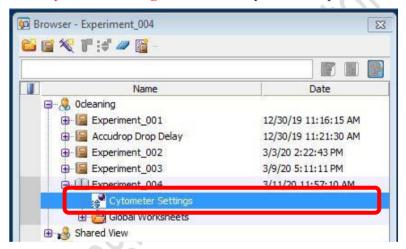
C. Experiment Setup

1. Setup New Experiment

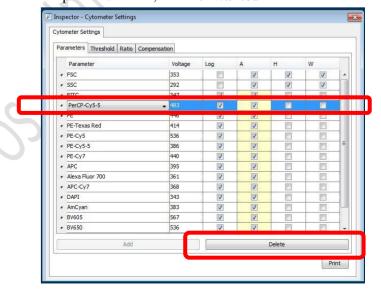
1.1 Go to Browser, Click New Experiment icon. A new experiment will be created



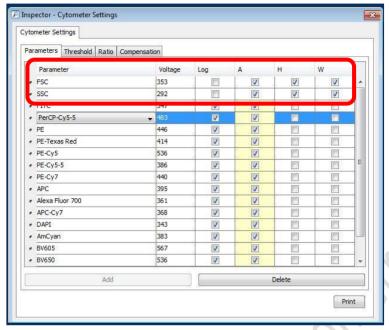
1.2 Click Cytometer Settings under the newly created Experiment



1.3 Go to Inspector Window, select unwanted channels and click Delete



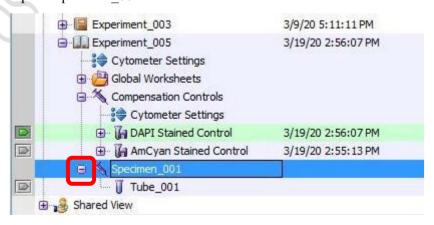
1.4 Check H and W boxes of FSC and SSC



- 1.5 Keep Log boxes of FSC and SSC unchecked
- 1.6 Keep Log boxes of all fluorescence channels checked
 - * If you are doing **cell cycle or DNA content** analysis, please keep **log box** of your DNA specific fluorescence channel **unchecked**.
- 2. Setup Compensation Please refer to FACSDiva compensation automated protocol
- 3. Setup Plots and Tables
 - 3.1 Go to Browser, Click New Specimen icon



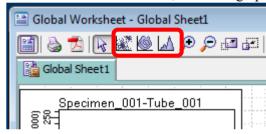
3.2 Expand Specimen 001

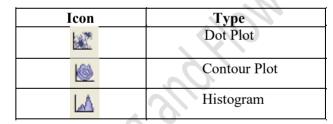


3.3 Click the tube **pointer** of Tube 001

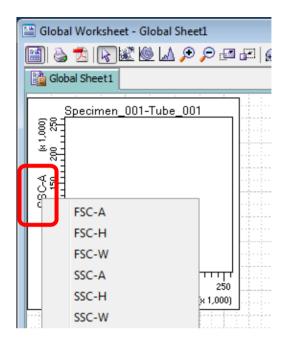


3.4 Go to Global Sheet Window, Click the graph type icon

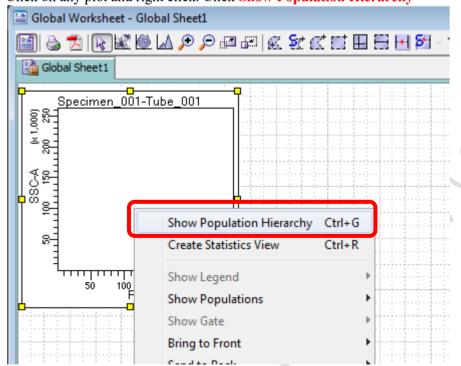




- 3.5 Click on the blank area of Global Worksheet window to create a new plot.
- 3.6 Mouse over the axis label and right click. Select the parameters of interest from the list.



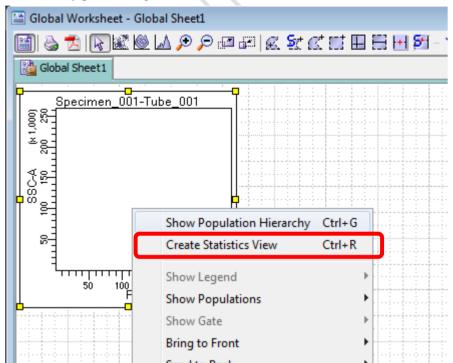
- 3.7 Repeat step 3.4 3.6 until all plots needed is created.
- * Essential Plots: FSC-A VS SSC-A; FSC-H VS FSC-W; SSC-H VS SSC-W
- 3.8 Click on any plot and right click. Click **Show Population Hierarchy**



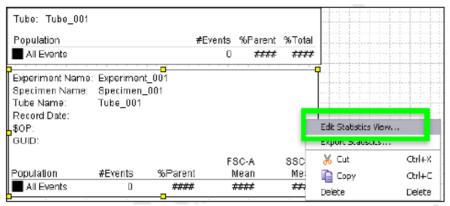
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Imaging and Flow Cytometry Core

3.9 Click on any plot and right click. Click Create Statistics View



3.10 Right Click on Statistics View table, select **Edit Statistics View** to select statistics of interest to be shown in the table.



3.11 Click Statistics Tab, check the boxes of interested statistics and then click OK

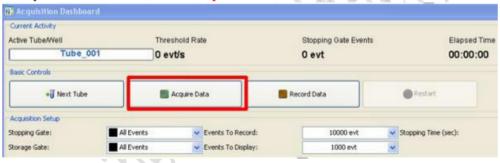


D. Sample Acquisition

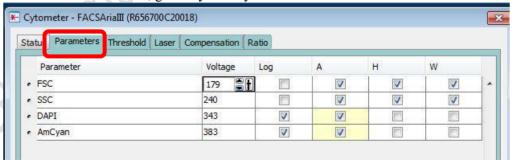
- 1. Gently tab or pulse vortex your sample and put your sample tube on SIP (Sample injection port).
 - *Run the unstained/negative control sample before stained samples.
 - !! DO NOT return the support arm to the center without putting a tube filled with liquid on SIP!!
- 2. Press RUN and LOW on the fluidics control panel.



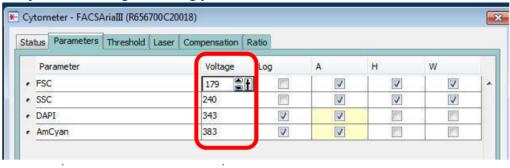
3. Go to Acquisition Dashboard, Click Acquire Data



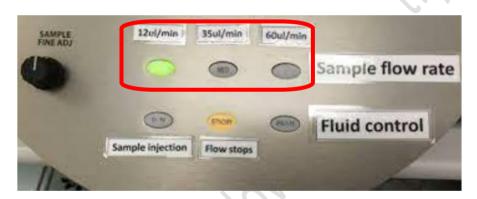
4. When Threshold rate > 0, go to Cytometry window and Click Parameters tab.



5. Adjust PMT Voltage accordingly



6. Adjust Sample flow rate on the fluidics control panel if needed (Maximum Threshold rate 8000 evt/s)



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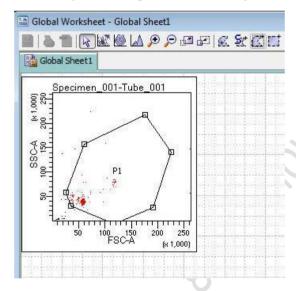
E. Create Gates

1. Go to Global Sheet Window, Click the type of gate needed



Icon	Туре
Œ	Polygon Area Gate
tat	Rectangle Area Gate
	Quantrad Gate
	Interval Gate

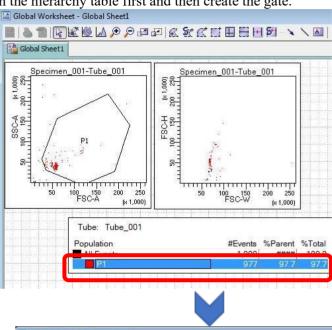
2. Draw the gate on the plot of interest to gate out target cluster /peak

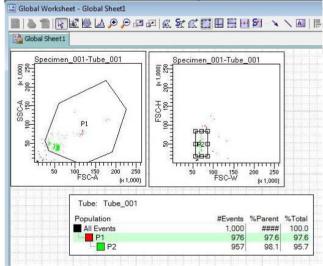




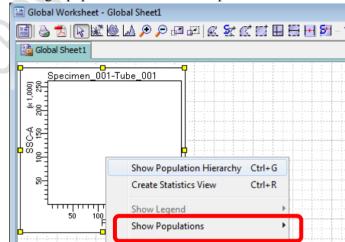
3. If you want to create a new population out of parent population, highlight the parent gate on the hierarchy table first and then create the gate.

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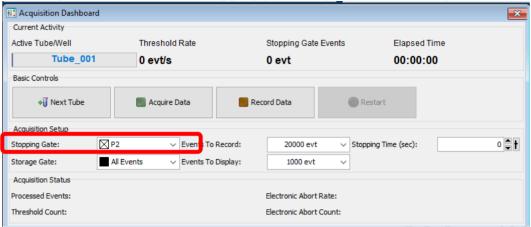


4. Click on target plot and right click, Click *Show Population > Target population* to visualize the target population ONLY in that plot.

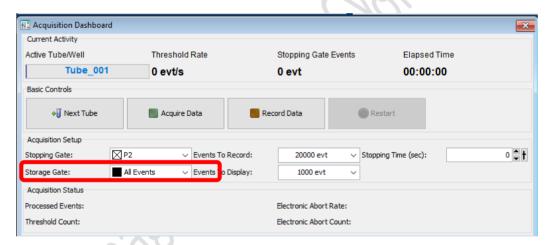


F. Data Recording

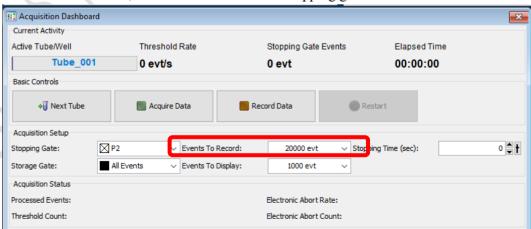
1. Go to Acquisition Dashboard, set **Stopping gate** to singlet gate or live cell gate



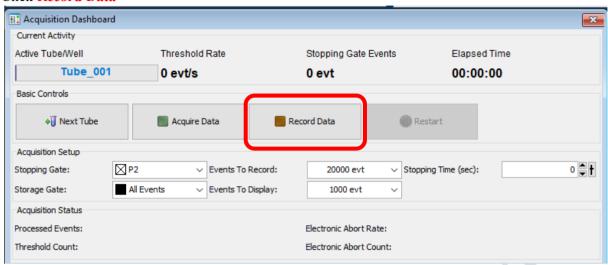
2. Set the Storage gate to All Events



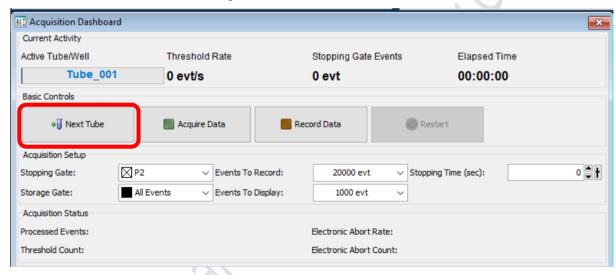
3. Set Events To Record, i.e. events number out of stopping gate to be recorded



4. Click Record Data

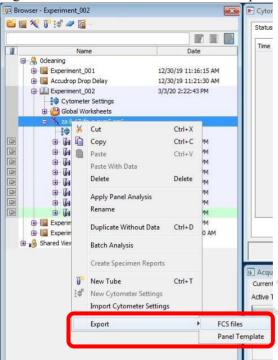


5. Click **Next Tube** to create a new sample

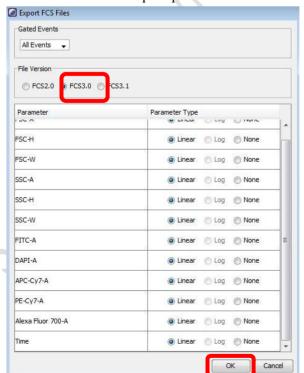


G. Data Export

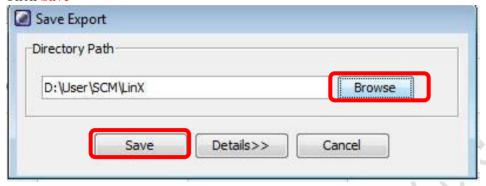
- 1. FCS file
 - 1.1 Go to Browser window, Select the Tubes / Specimen of interest.
 - 1.2 Right Click over the selection and click *Export* > *FCS file*



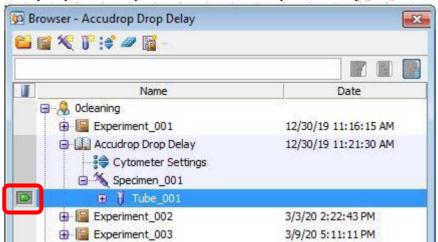
1.3 Select FCS 3.0 and keep all parameters Linear. Click OK



- 1.4 Click **Browse** to choose the destination (D:/User/Department/Personal Folder)
- 1.5 Click Save



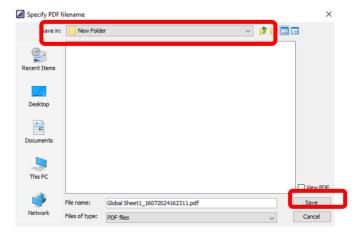
- 2. PDF file
 - 2.1 To export pdf of multiple tubes, Click the tube pointer of any tubes



2.2 Click PDF icon



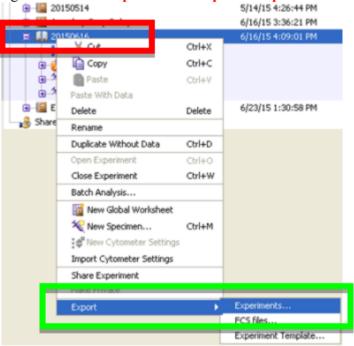
2.3 Choose the destination (D:/User/Department/PersonalFolder) and click Save



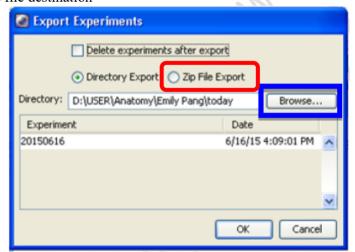


3. Experiment

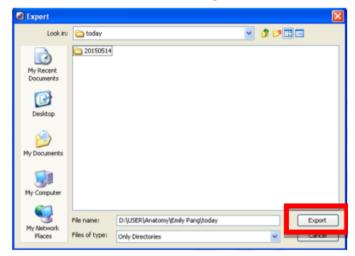
3.1 Right click on the **Experiment > Export > Experiments**



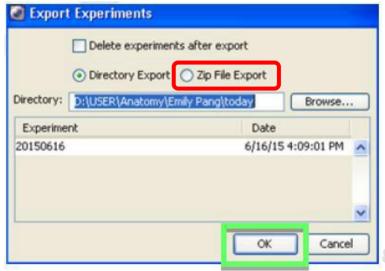
3.2 Choose **Zip File Export** and click **Browse** to select the file destination



Select the file destination and click **Export**.

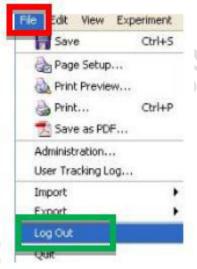


3.4 Click **OK** to export (select Zip File Export)



H. Software Log out

1. To log out of FACSDiva software, go to File and click Log Out.





I. System Cleaning

*Cleaning procedure will not interfere with the data export and analysis.

- 1. Prepare 3 mL of each cleaning solution (Solution 1: FACSClean; Solution 2: FACSRinse; Solution 3: MilliQ water)
- 2. Press HIGH (60 uL/min) and RUN (Sample Injection) on fluidics control panel
- 3. Load the tube with Solution 1 on the SIP with the support arm on the side for 1 minute
- 4. Move the support arm under the tube and run for another 4 minutes
- 5. Repeat step 3 and 4 with Solution 2 and Solution 3*If PI stain is used, please clean the system 9 minutes instead of 4 for Solution 2
- 6. After cleaning with Solution 3, press LOW (12 uL/min) and STANDBY (Flow Stop)