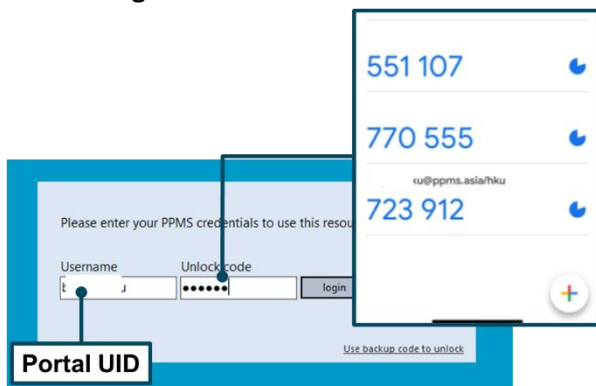




Imaging and Flow Cytometry Core

BD Aria SORP Standard Operation Protocol – Basic Operation

A. Tracker Login



B. Laser

Check if the laser needed is turned **ON**.

* Please contact our staff 30 mins in advance to turn **ON** the UV laser for you if your fluorochrome-conjugated antibodies must be activated by the UV, such as **Hoechst Blue**, **Hoechst Red** or **BUV395**. Otherwise, the UV laser is **OFF** to avoid DNA damage during sorting.

** Note that "**DAPI**" could be activated by **Violet** laser and detected by **BV421** channel.



Imaging and Flow Cytometry Core



C. User Login

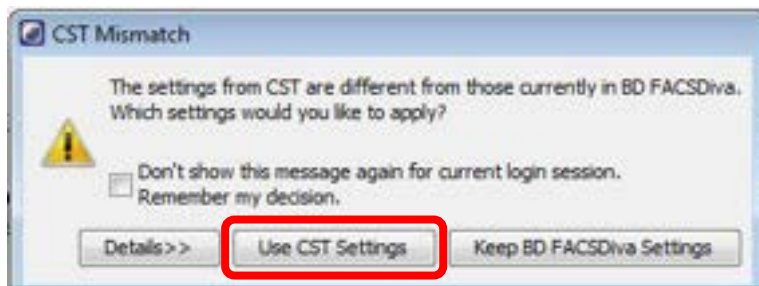
1. Login to FACSDiva Software with your username and password.
*If you do not have account, please contact our staff for assistance.



2. Click *Use CST Settings* if the window below pops up.



Imaging and Flow Cytometry Core



CPOS - Imaging and Flow Cytometry Core

Imaging and Flow Cytometry Core

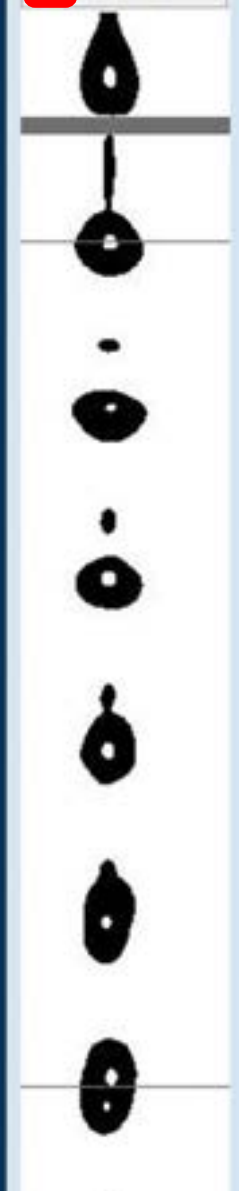
D. Stream Optimization

1. Go to 100 micron window (upper monitor)
2. Adjust the *Freq* (Starting Value 29.2) so the “neck” of a drop is formed
3. Adjust the *Ampl* (Starting Value 8.0) so the Drop 1 is close to 120 ~ 150 and Gap is close to 10
4. Turn ON the *Sweet spot*



1. 100 micron

4. Stream
 Sweet Spot



3. Ampl: 3.1

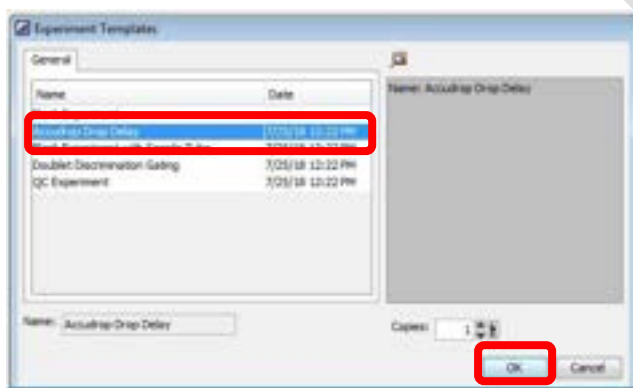
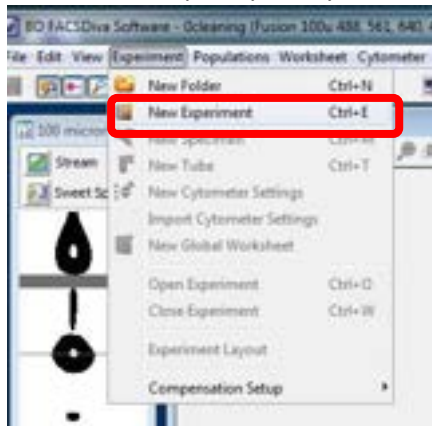
2. Freq: 29.2

| | | |
|---------|-----|-----|
| Drop 1: | 127 | 117 |
| Gap: | 10 | 9 |

Imaging and Flow Cytometry Core

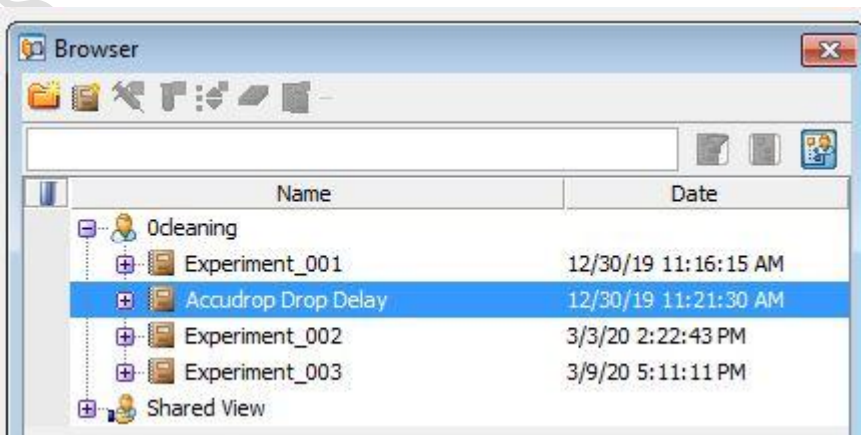
E. Accudrop Delay Assay

1. To Import Accudrop Drop Delay template, click *Experiment > New Experiment*.
Select Accudrop Drop Delay, then click *OK*



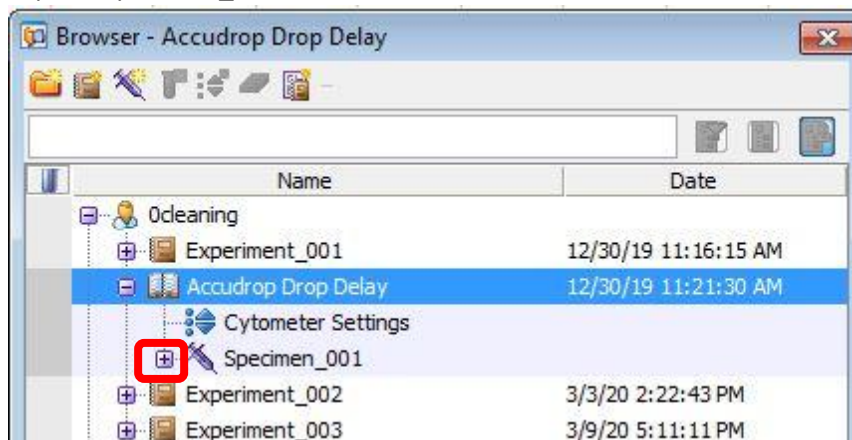
- OR -

2. To open existing Accudrop Drop Delay experiment, double click *Accudrop Drop Delay* on the Browser window

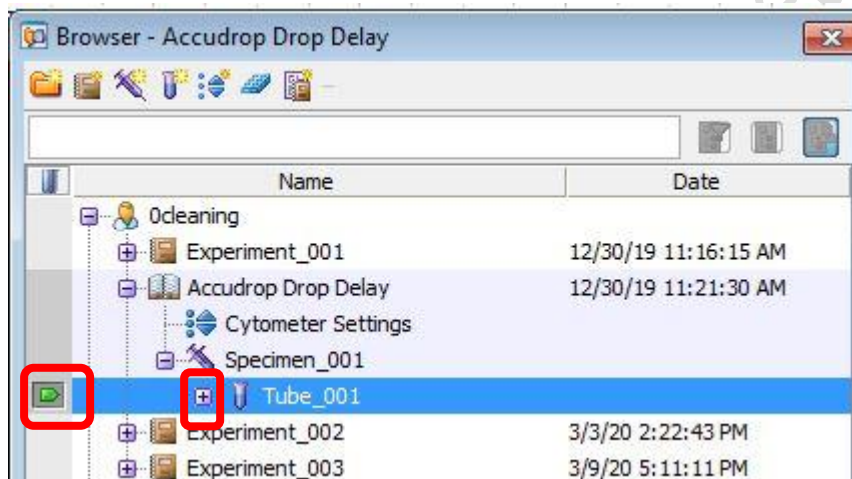


Imaging and Flow Cytometry Core

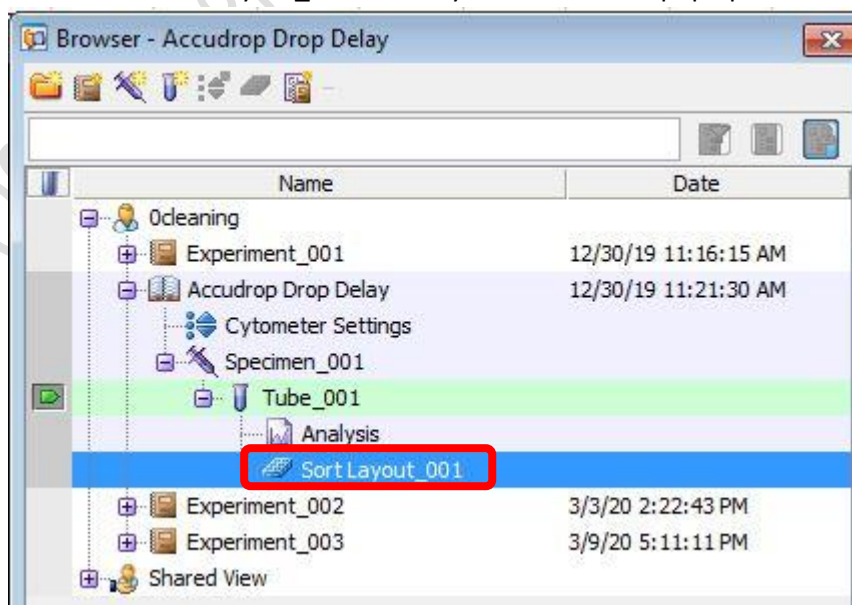
3. Expend Specimen_001



4. Click the tube pointer and expend Tube_001

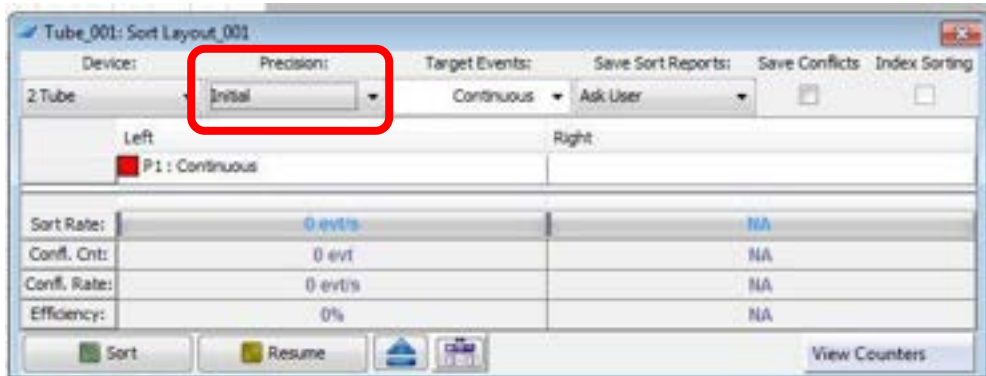


5. Double click Sort Layout_001. Sort Layout window will pop up

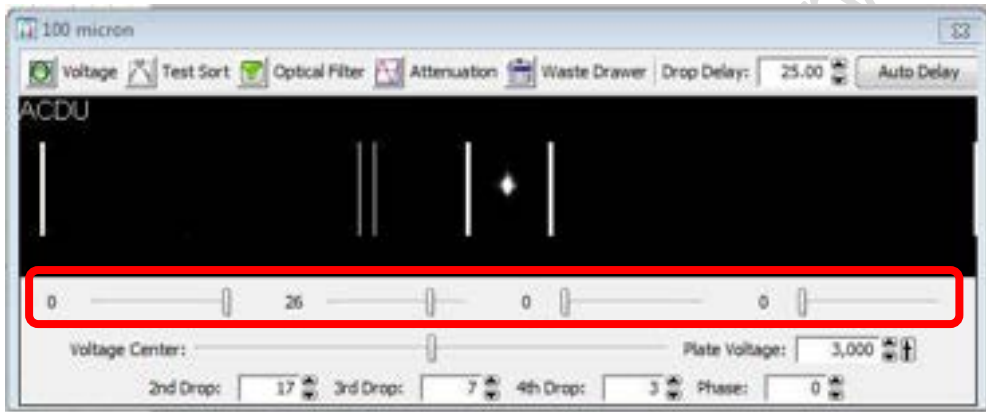


Imaging and Flow Cytometry Core

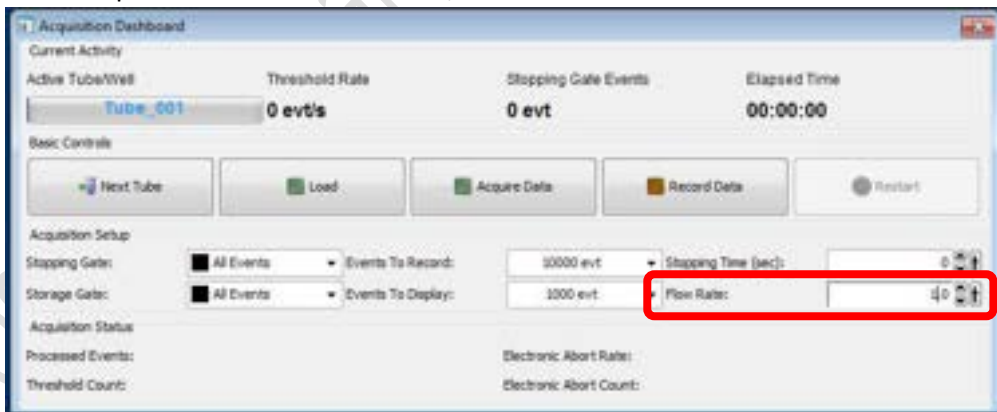
6. Go to Sort Layout Window, Select *Precision* > *Initial*



7. Go to 100 micron window (lower monitor), set the slider reading as 0 - 26 - 0 - 0

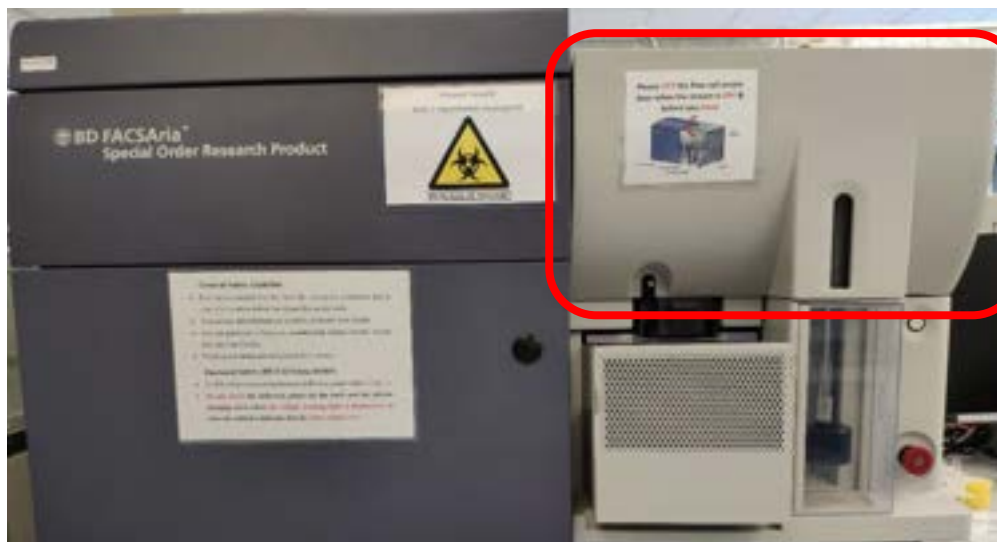


8. Go to Acquisition Dashboard window, Set *Flow rate* to 1.0



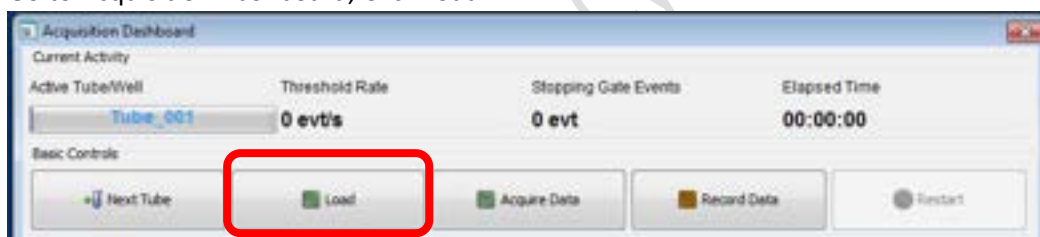
9. Close upper flow cell access door.

Imaging and Flow Cytometry Core



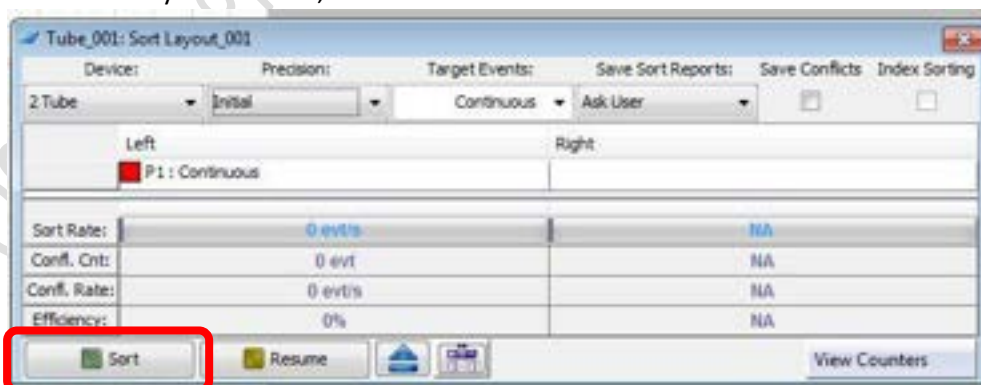
10. Load a tube of Accudrop beads (1 mL of PBS + 1 drop of stock) on the sample stage

11. Go to Acquisition Dashboard, Click *Load*



12. Adjust *Flow rate* if needed to obtain threshold rate constantly **1000-1500** events per sec

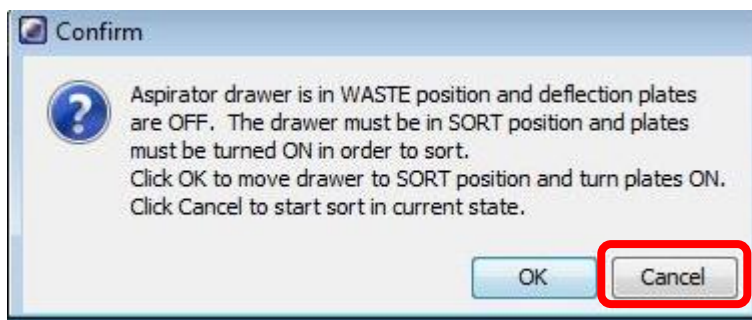
13. Go to Sort Layout window, click *Sort*



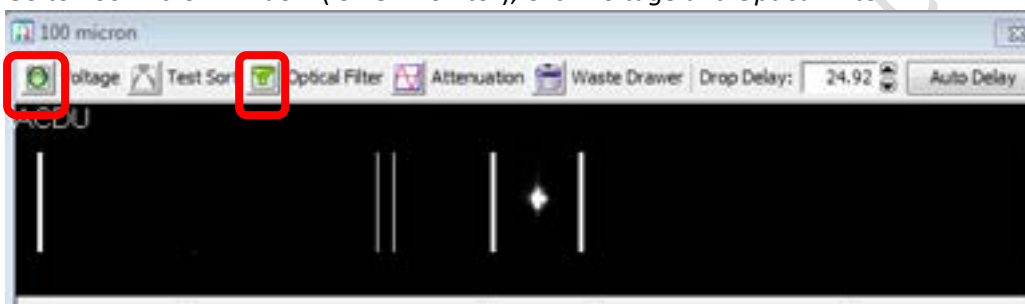


Imaging and Flow Cytometry Core

14. Click *Cancel* on the confirm window



15. Go to 100 micron window (lower monitor), Click *Voltage* and *Optical Filter*

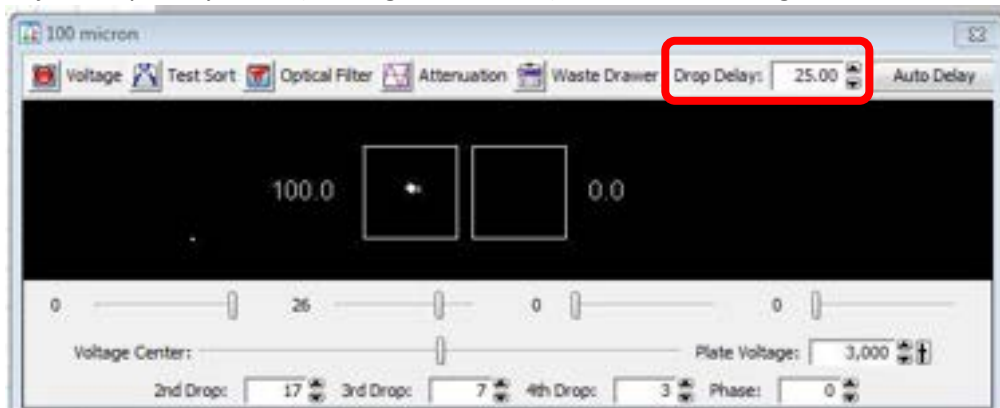


CPOS - Imaging and Flow Cytometry Core

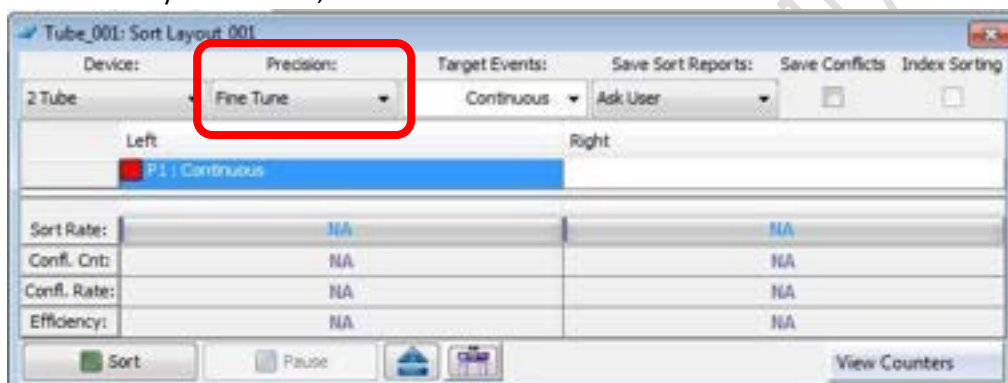


Imaging and Flow Cytometry Core

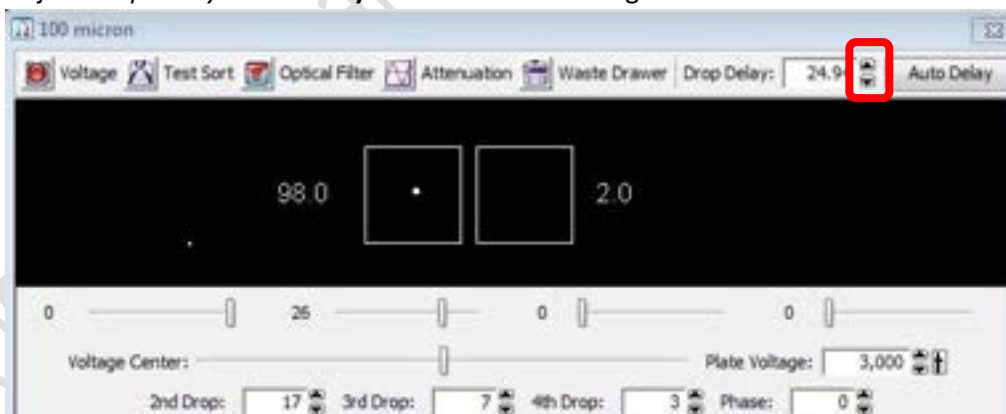
16. Adjust *Drop Delay* value (starting value: 25.00) so that the reading on the left reach 100



17. Go to Sort Layout window, Select Precision > Fine Tune



18. Adjust *Drop Delay* value **bit by bit** so that the reading on the left reach >97



Imaging and Flow Cytometry Core

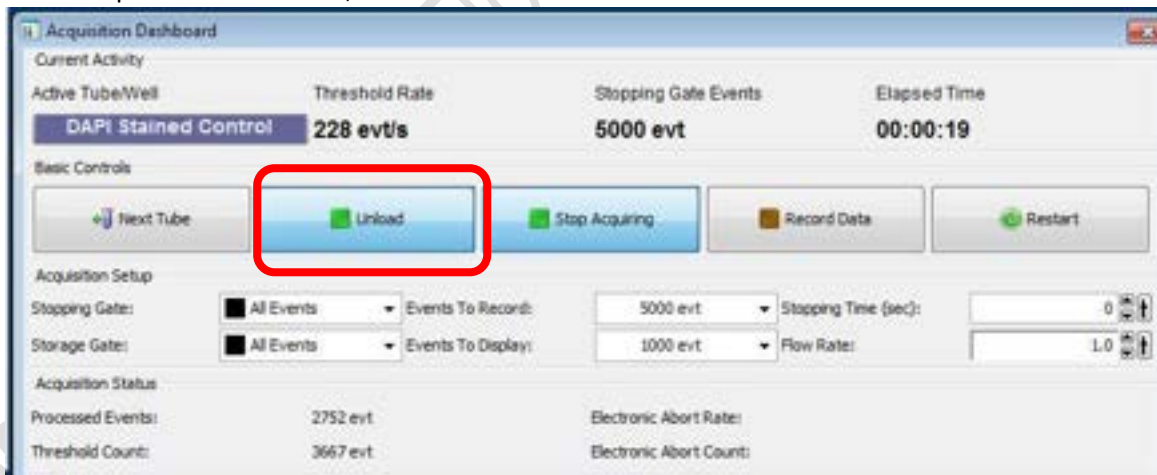
19. Go to Sort Layout Window, Click Sort again.



20. Click *Cancel* on the confirm window



21. Go to Acquisition Dashboard, Click *Unload*



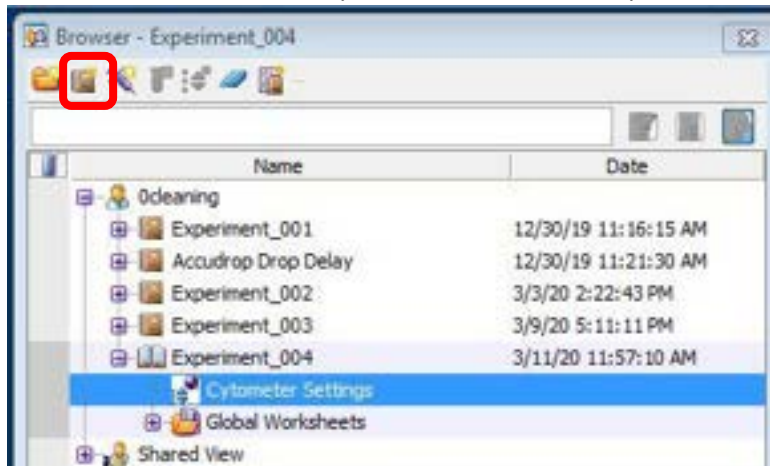
22. Return the Accudrop beads to 4 degree refrigerator

Imaging and Flow Cytometry Core

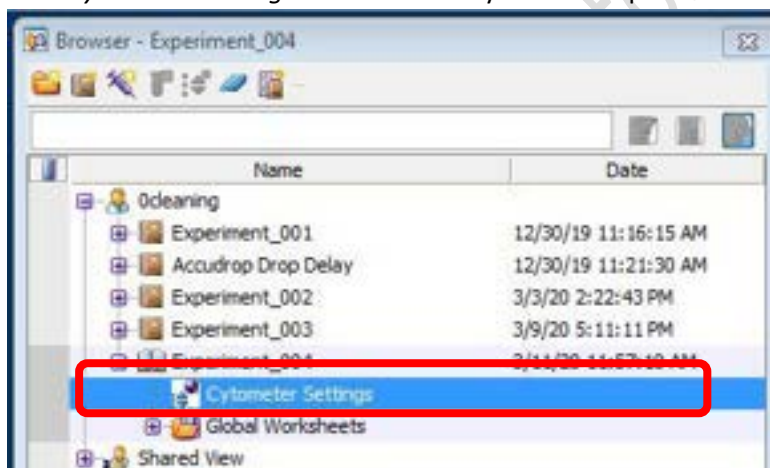
F. Experiment Setup

1. Setup New Experiment

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



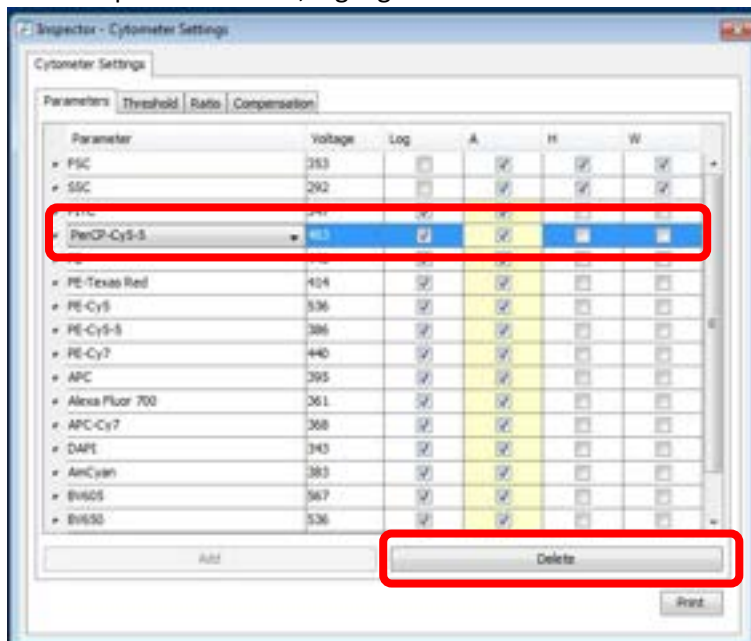
1.2 Click *Cytometer Settings* under the newly created Experiment



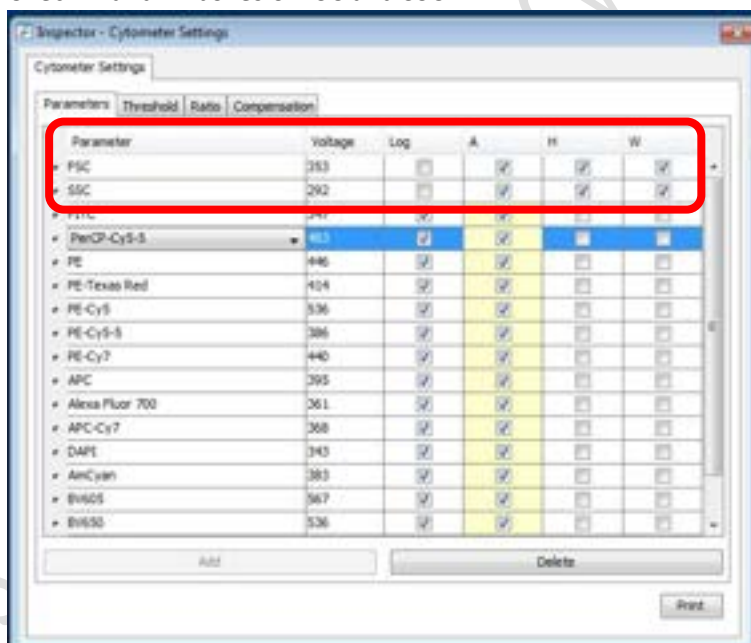
CPOS - Image

Imaging and Flow Cytometry Core

1.3 Go to Inspector Window, highlight **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**

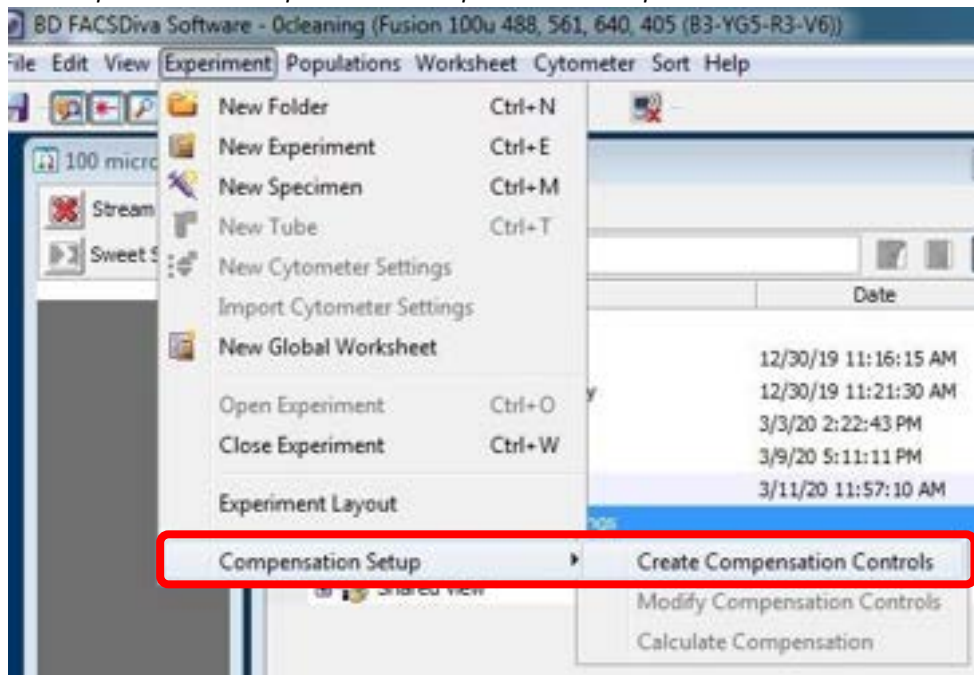
* If you are doing **cell cycle or DNA content** analysis, please keep **log box** of your DNA specific fluorescence channel **unchecked**.

2. Setup Compensation (for Multi-colour panel)



Imaging and Flow Cytometry Core

2.1 Click *Experiment > Compensation Setup > Create Compensation Control*

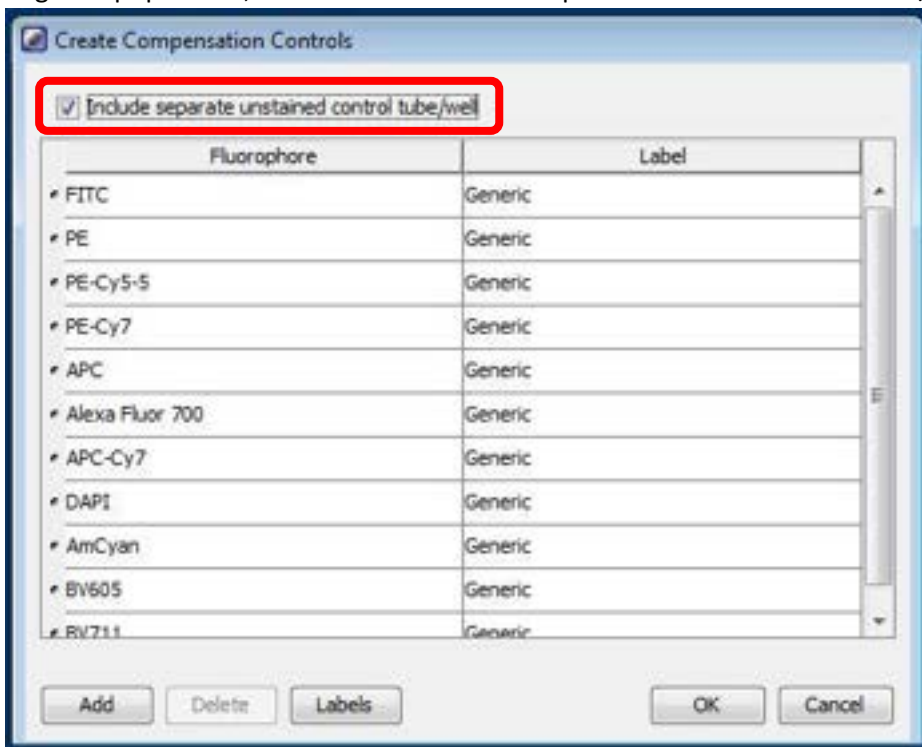


CPOS - Imaging and Flow Cy

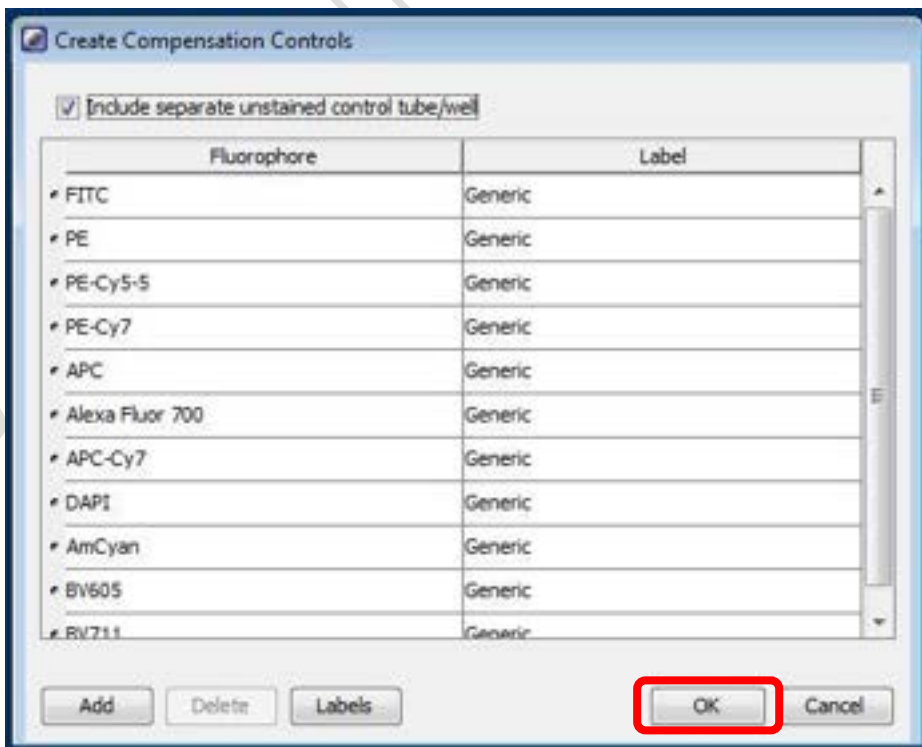


Imaging and Flow Cytometry Core

- 2.2 If any one of your single stain controls is known to be 100% positive, i.e. no negative population, check the box “Include separate unstained control tube/well”.

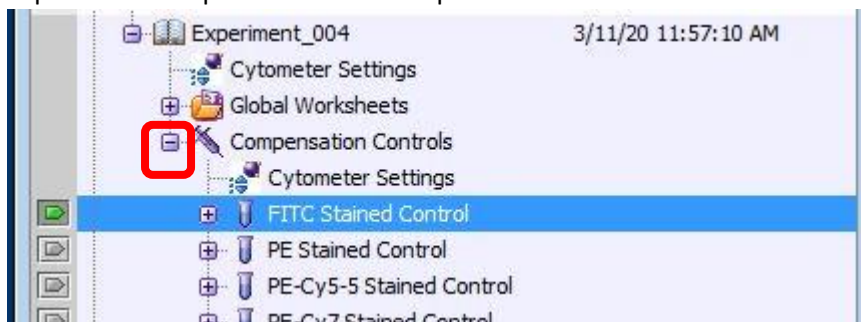


- 2.3 Click OK

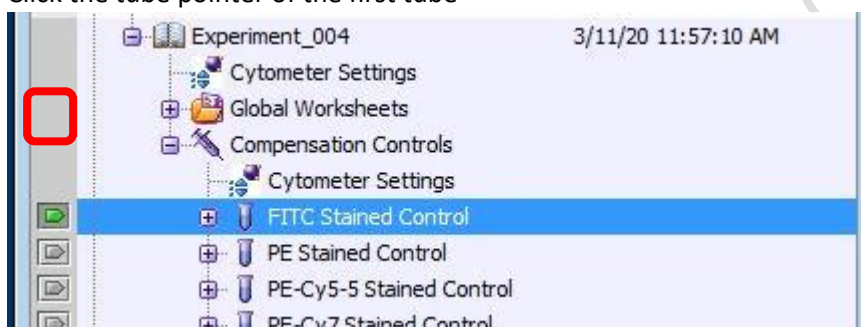


Imaging and Flow Cytometry Core

2.4 Expand the Compensation Control Specimen



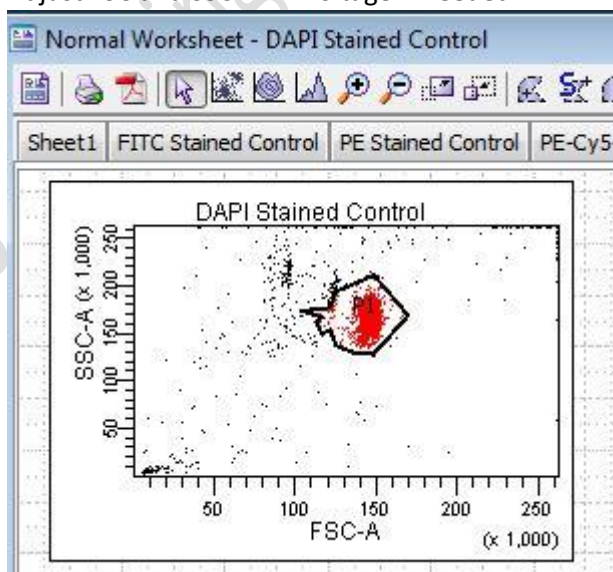
2.5 Click the tube pointer of the first tube



2.6 Load the single stain controls on the sample stage according to the tube label, i.e. run FITC single stain when the tube pointer is pointing at "FITC Stained Control"

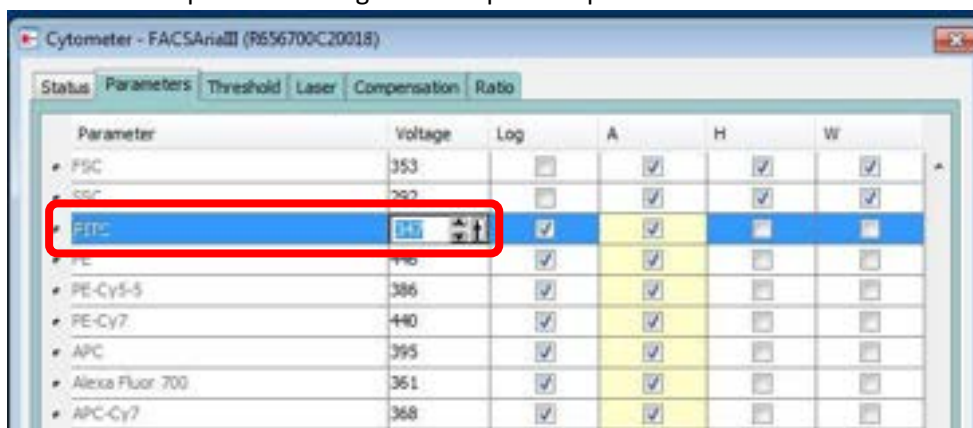
2.7 Go to Acquisition Dashboard, Click *Load*.

2.8 Go to Normal Worksheet, move the P1 gate to include major cluster. Adjust FSC and SSC PMT Voltage if needed



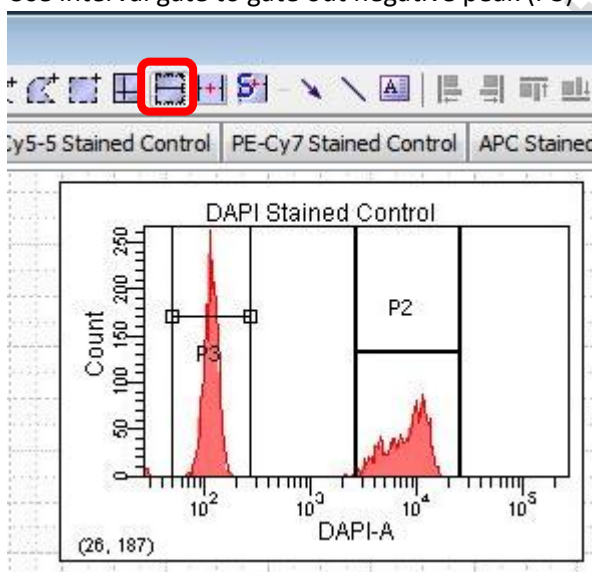
Imaging and Flow Cytometry Core

- 2.9 Go to Cytometer window, Fine tune the corresponding fluorescence PMT voltage to have best separation of negative and positive peak

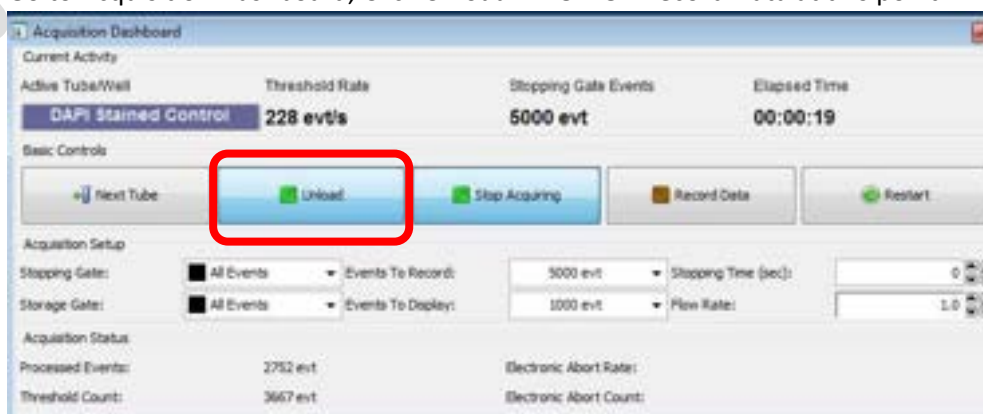


- 2.10 Move the interval gate (P2) to include the positive peak

- 2.11 Use interval gate to gate out negative peak (P3)



- 2.12 Go to Acquisition Dashboard, Click *Unload*. *DO NOT Record Data at this point.

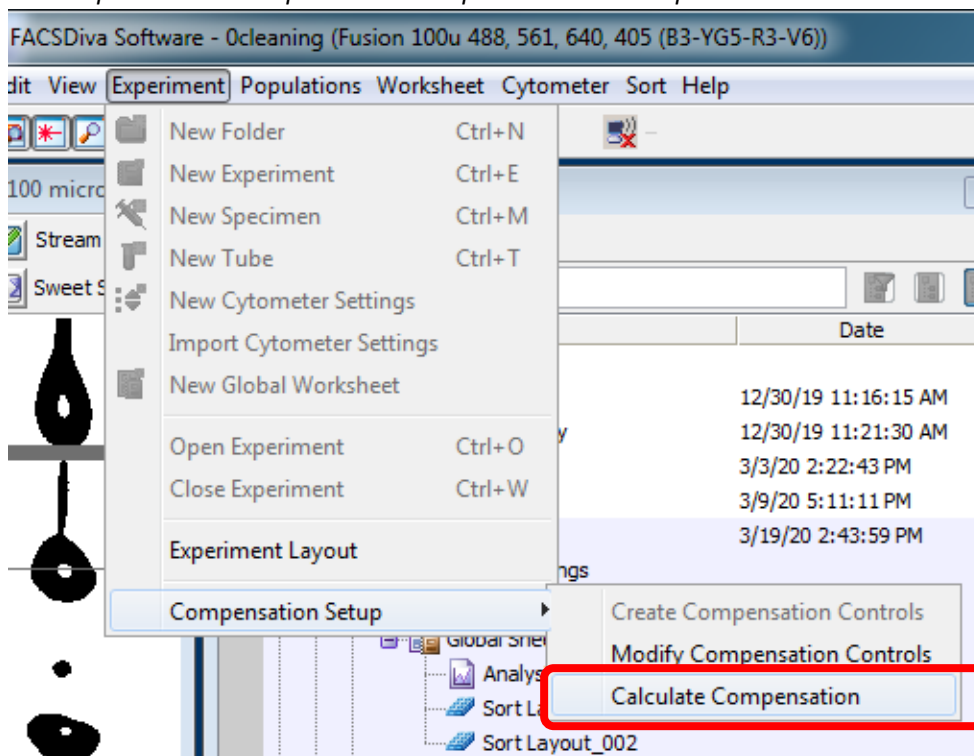


Imaging and Flow Cytometry Core

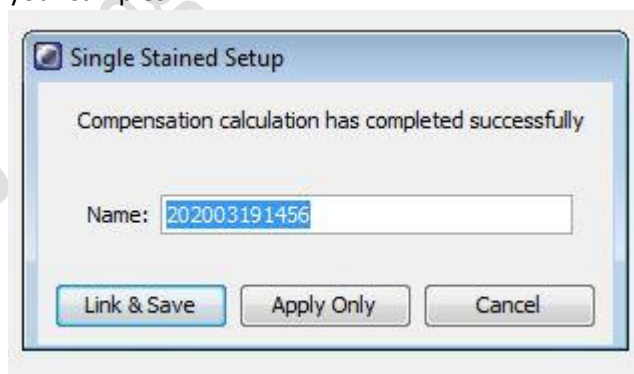
2.13 Repeat step 2.5 – 2.12 with all the single stain controls.

2.14 After optimising the PMT voltage of ALL the fluorescence channel, load each single stained control and click *Record Data* for ALL single stain controls

2.15 Click *Experiment > Compensation Setup > Calculate Compensation*



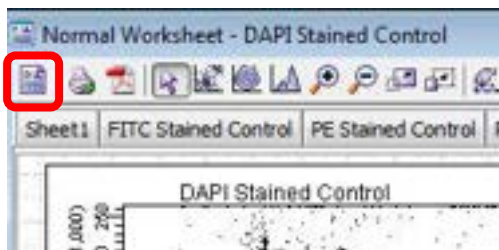
2.16 Click *Link and Save* for the most stringent practice, i.e. cannot adjust PMT voltage anymore **OR** Click *Apply Only* for some flexibility on PMT voltage adjustment of your samples.



2.17 Switch Normal worksheet to Global worksheet



Imaging and Flow Cytometry Core

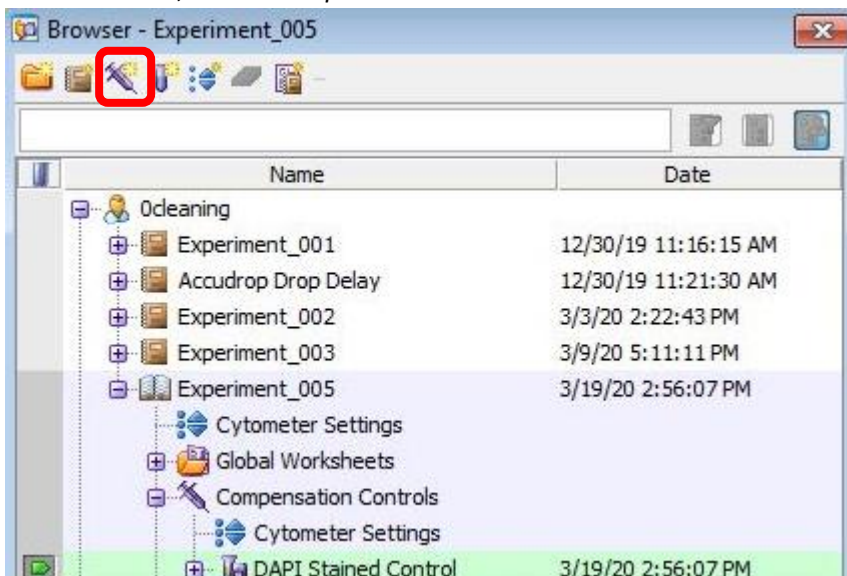


CPOS - Imaging and Flow Cytometry Core

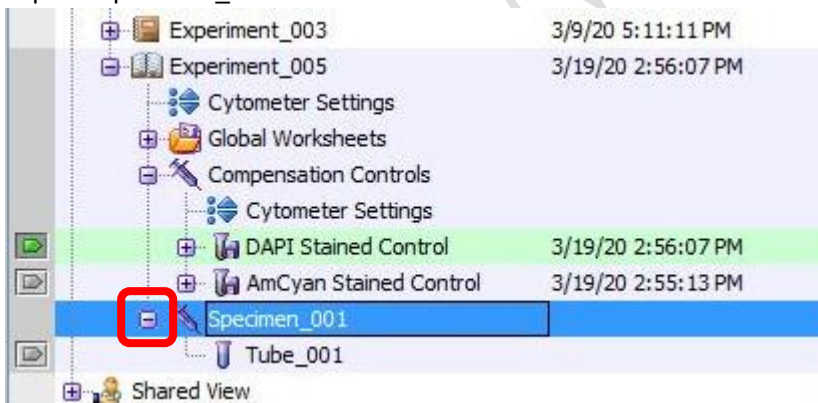
Imaging and Flow Cytometry Core

3. Setup Plots and Tables

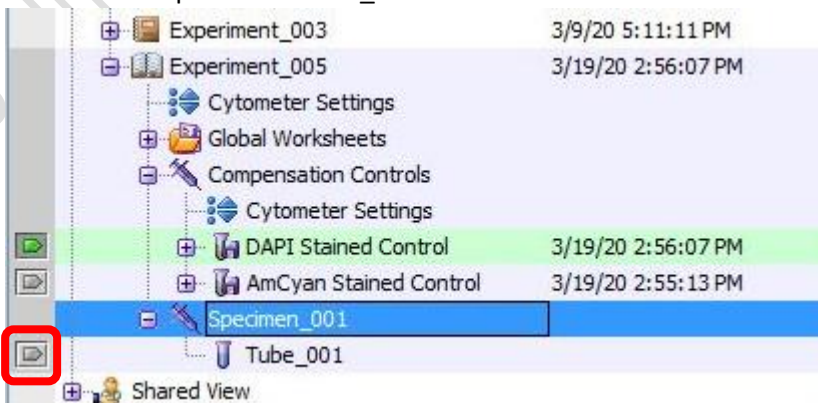
3.1 Go to Browser, Click *New Specimen icon*



3.2 Expand Specimen_001

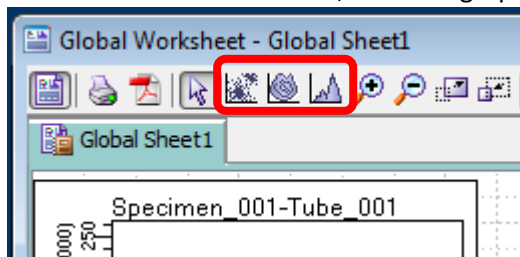





3.3 Click the tube pointer of Tube_001



Imaging and Flow Cytometry Core

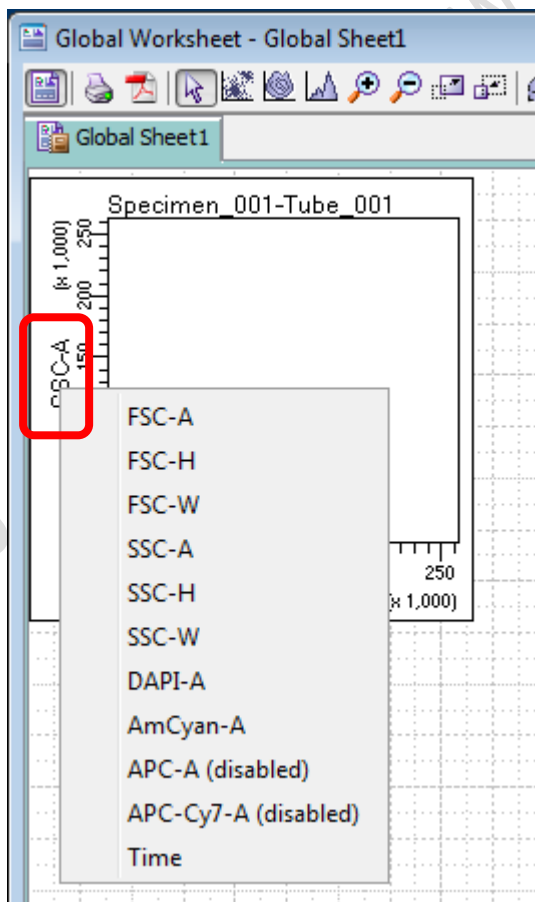
3.4 Go to Global Sheet Window, Click the graph type icon



| Icon | Type |
|---|--------------|
|  | Dot Plot |
|  | Contour Plot |
|  | Histogram |

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.

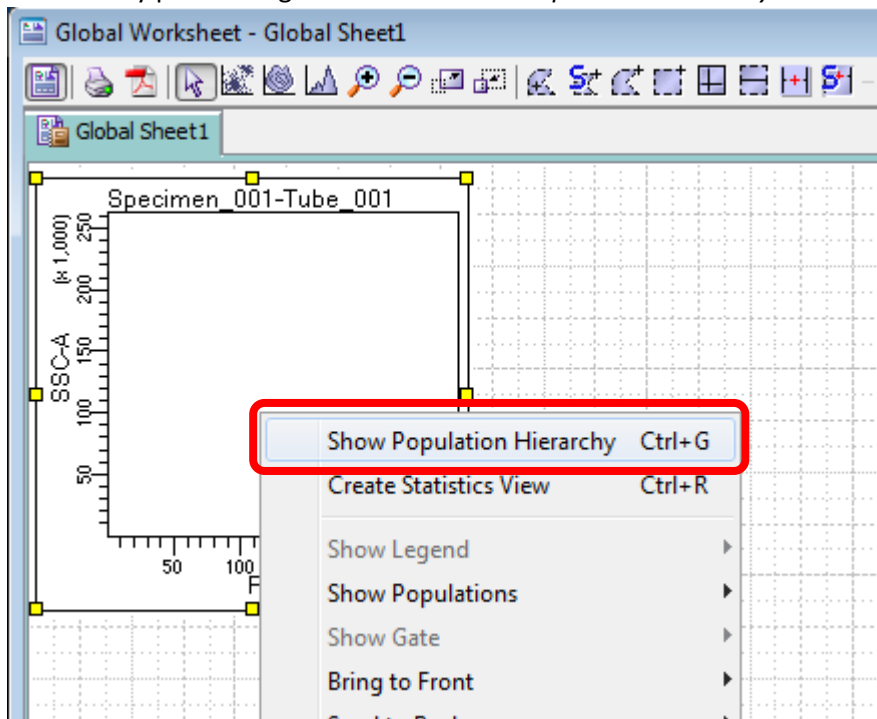


Imaging and Flow Cytometry Core

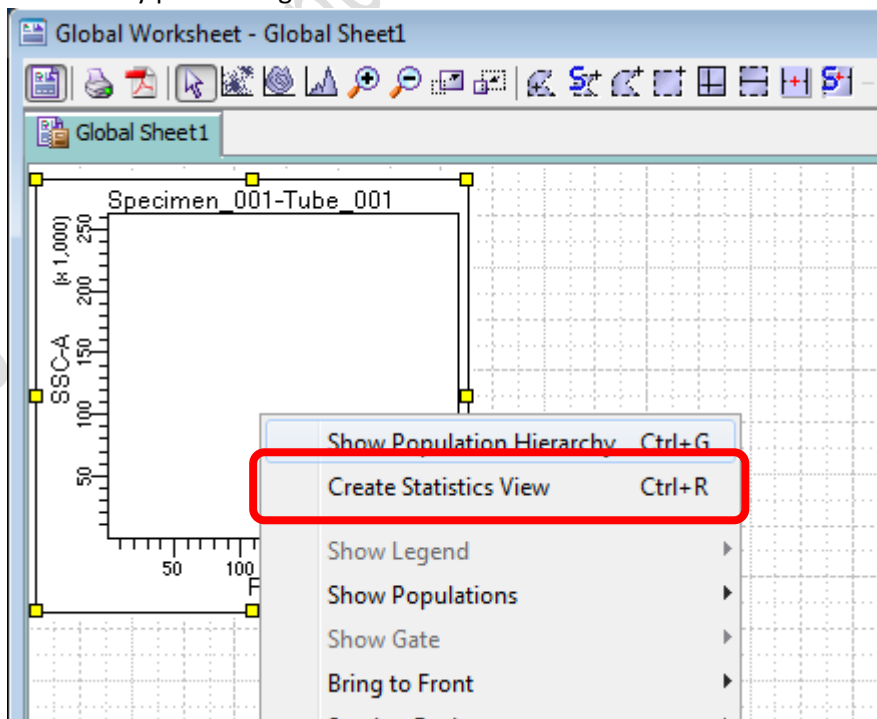
3.7 Repeat step 3.4 – 3.6 until all plots needed is created.

*** 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*



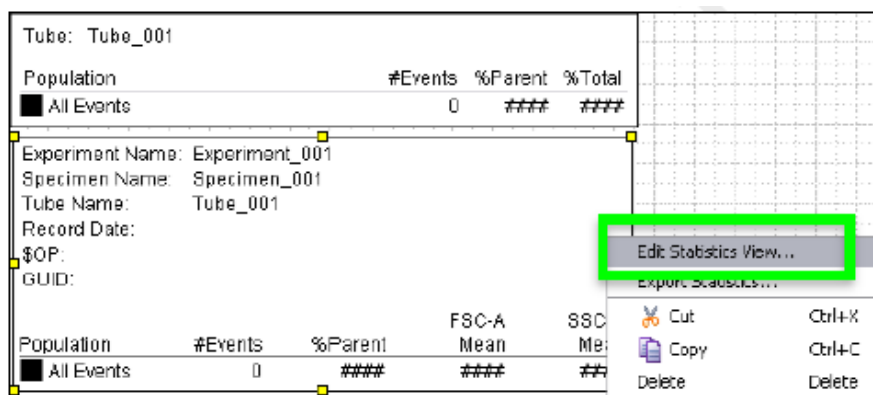
3.9 Click on any plot and right click. Click *Create Statistics View*



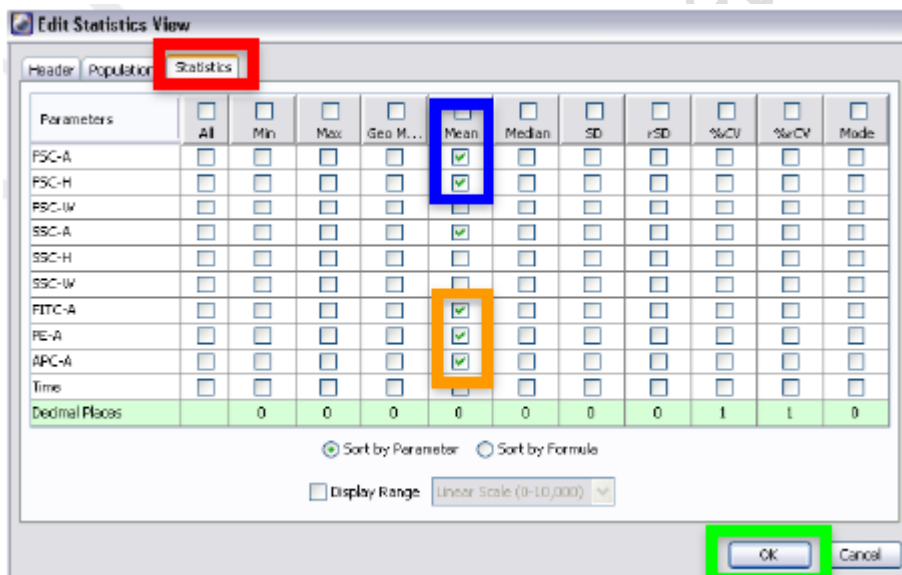


Imaging and Flow Cytometry Core

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

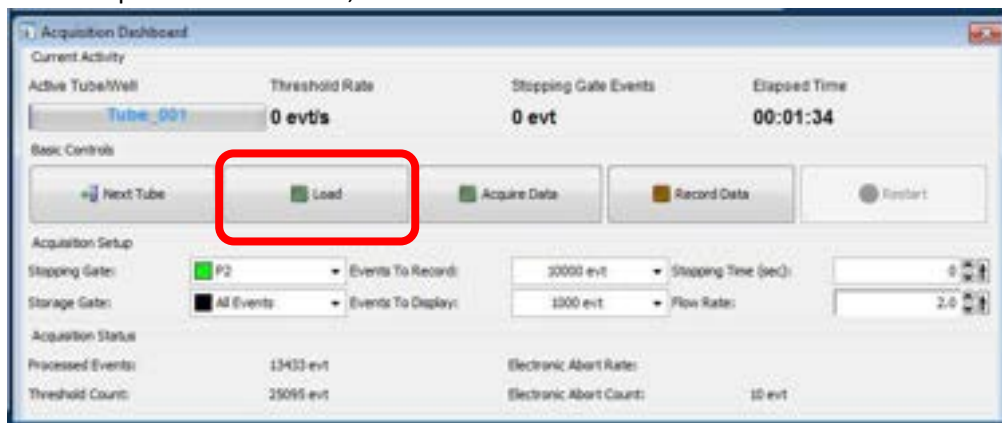


3.11 Click *Statistics* Tab, check the boxes of interested statistics and then click *OK*



Imaging and Flow Cytometry Core

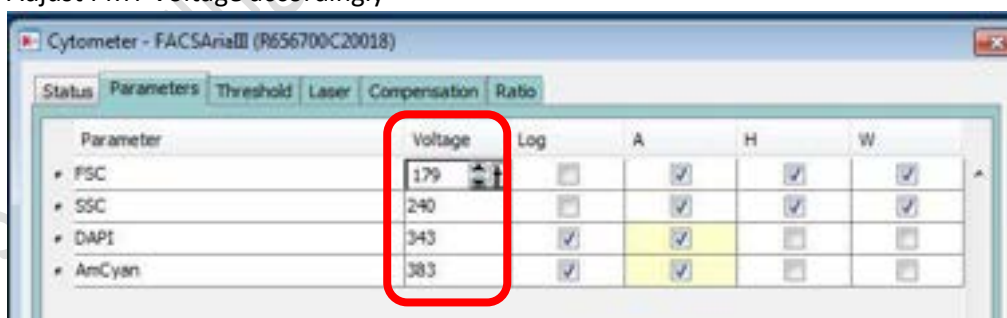
4. Sample Acquisition
 - 4.1 Load your sample on the sample stage
 - 4.2 Go to Acquisition Dashboard, Click *Load*



- 4.3 When you start seeing dots appear on the plot, Go to Cytometry window and Click *Parameters*



- 4.4 Adjust PMT Voltage accordingly

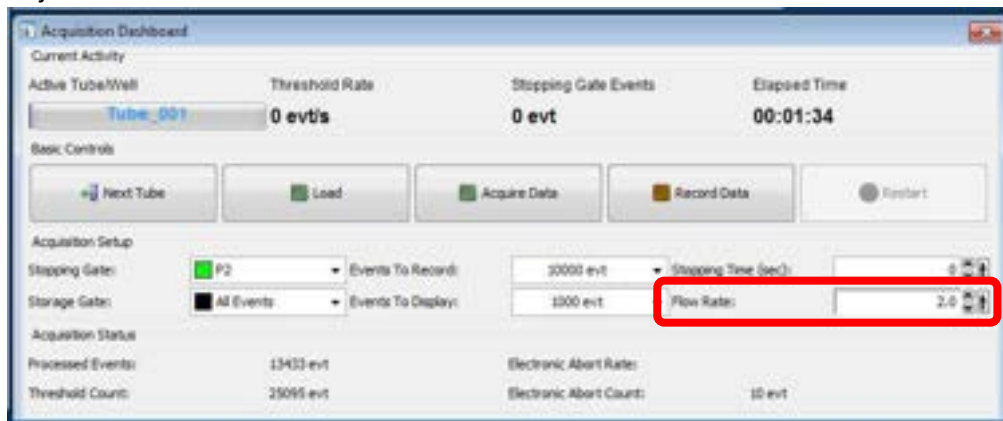


| Channel | Suggested Voltage range for mammalian cells |
|--------------|--|
| FSC | 180-300 *If you sample cell size is too big to visualise with FSC voltage 180, you may change the FSC ND filter from 1.0 to 1.5 |
| SSC | 230-330 |
| Fluorescence | 300-850 |



Imaging and Flow Cytometry Core

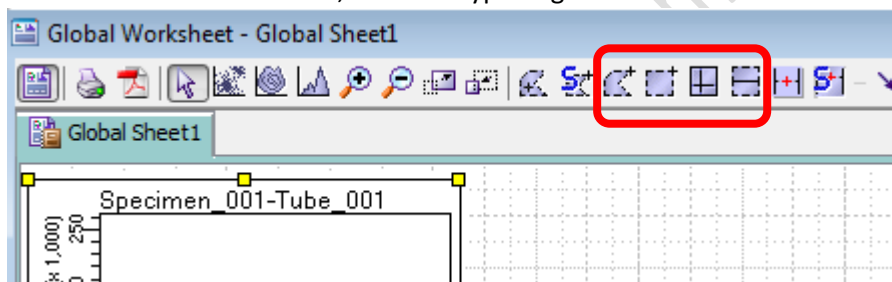
4.5 Adjust *Flow rate* if needed



* If you perform sorting, DO NOT set flow rate > 5.0 or threshold rate > 5000 event/s

5. Create Gates

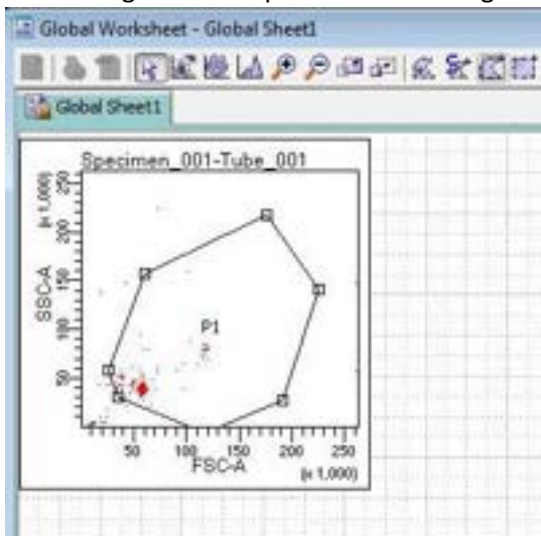
5.1 Go to Global Sheet Window, Click the type of gate needed



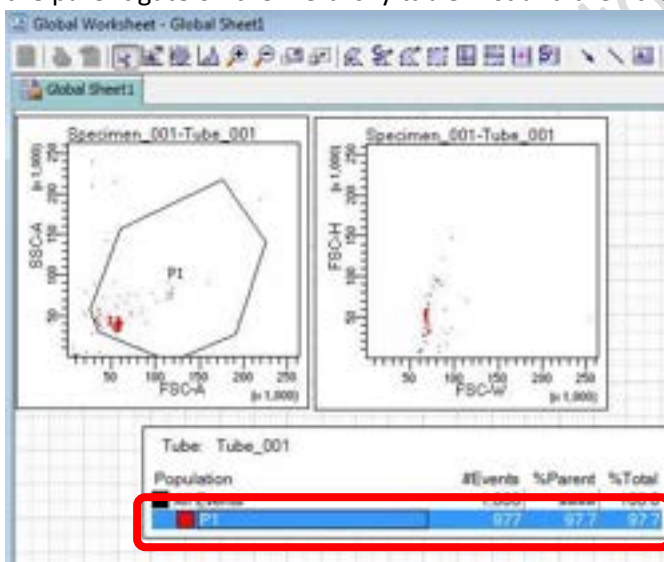
| Icon | Type |
|------|---------------------|
| | Polygon Area Gate |
| | Rectangle Area Gate |
| | Quadrant Gate |
| | Interval Gate |

Imaging and Flow Cytometry Core

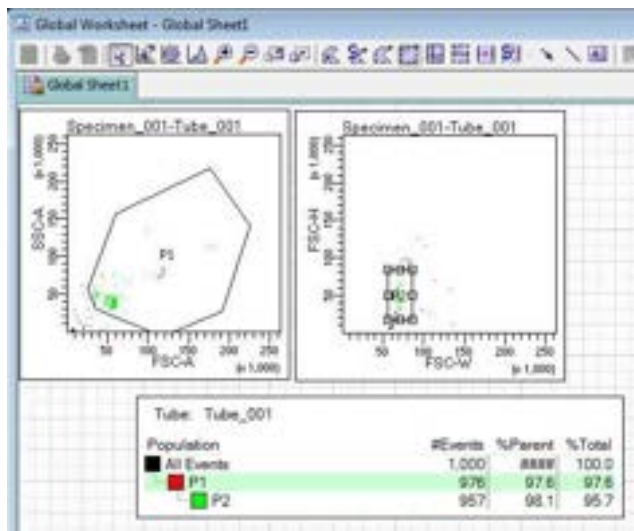
5.2 Draw the gate on the plot of interest to gate out target cluster /peak



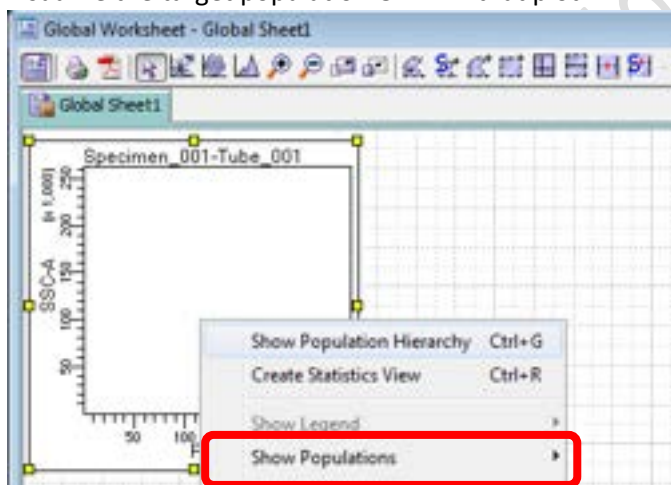
5.3 If you want to create a new population out of particular parent population, highlight the parent gate on the hierarchy table first and then create the gate.



Imaging and Flow Cytometry Core



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



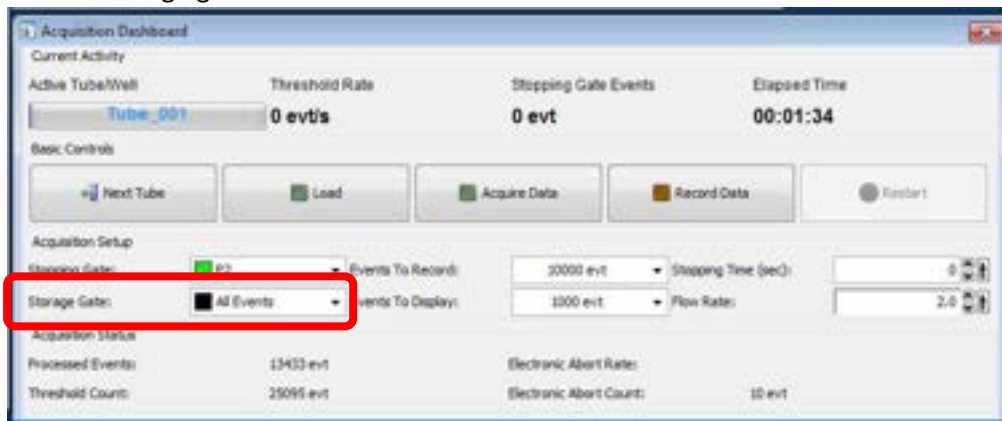
6. Data Recording

6.1 Go to Acquisition Dashboard, set Stopping gate to singlet gate or live cell gate

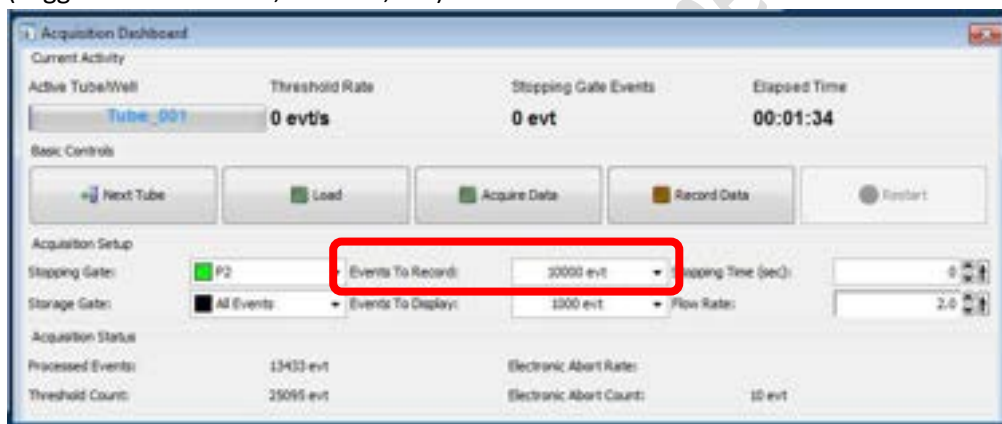


Imaging and Flow Cytometry Core

6.2 Set the Storage gate to All Events



6.3 Set Events To Record, i.e. events number out of stopping gate to be recorded (suggested number 10,000 – 50,000)



6.4 if the sample is Unload or Acquisition is stopped, Click *Load* or *Acquire Data*

6.5 Click *Record Data*

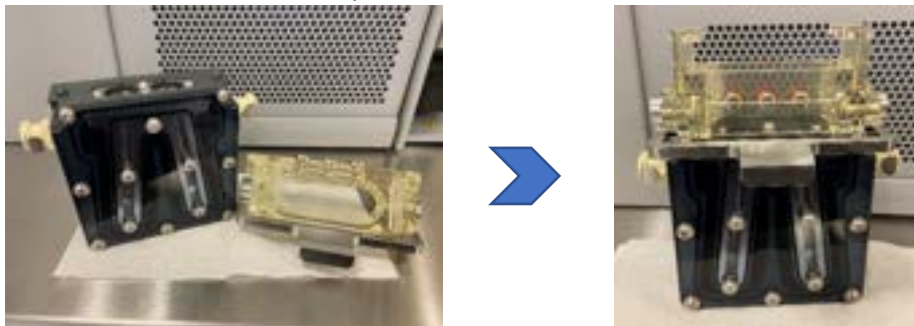
6.6 Click *Next Tube* to create a new sample

Imaging and Flow Cytometry Core

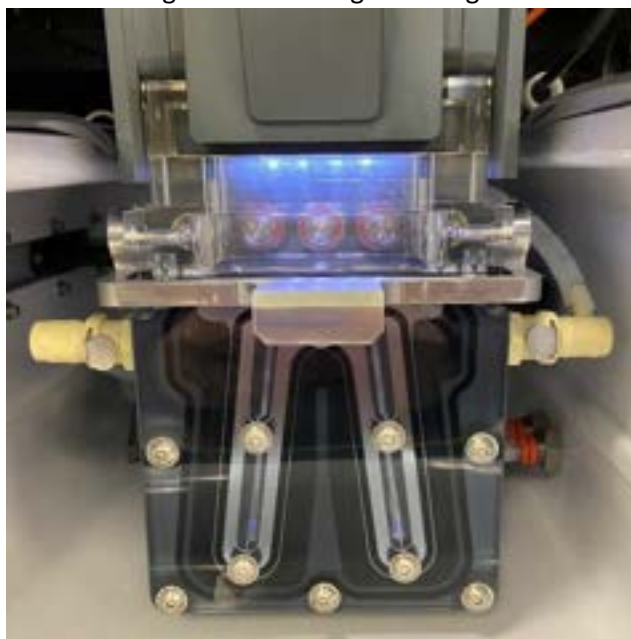
G. Sort Device Alignment

1. Tube Holder

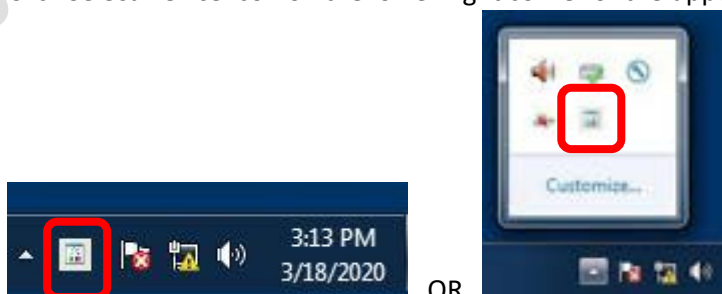
1.1 Assemble the tube holder as picture below



1.2 Put collection tubes into position. Slide the tube holder right under the sort chamber. Plug in water tubing if cooling is needed.



1.3 Click *Select Device* icon on the lower right corner of the upper monitor





Imaging and Flow Cytometry Core

1.4 Select collection device of interest



1.5 Go to 100 micron window (lower monitor). Adjust the Slider

| Device | Suggested Slider reading |
|----------------------|--------------------------|
| 2-tube 15 ml | 0 – 49 – 41 – 0 |
| 4-tube 1.5ml / 2.0ml | 71 – 30 – 25 – 68 |
| 4-tube 5 ml | 80 – 30 – 25 – 74 |

1.6 Click *Voltage*. Wait 2 seconds and then Click *Test Sort*.



1.7 Adjust the slider so the side stream dot lines within the target lines

Imaging and Flow Cytometry Core



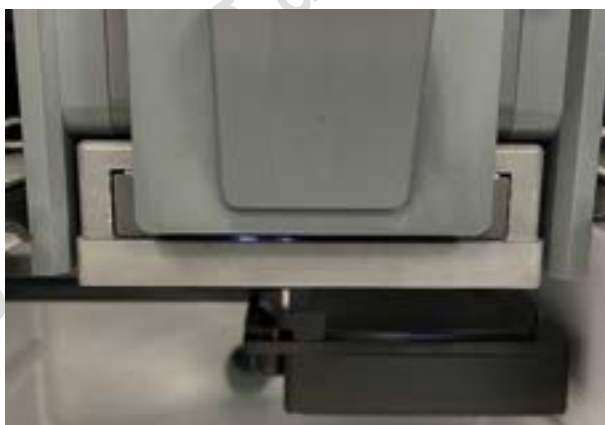
1.8 Turn off *Voltage* and *Test Sort*

2. ACDU (Culture plate)

2.1 Slide the ACDU adaptor right under the sort chamber



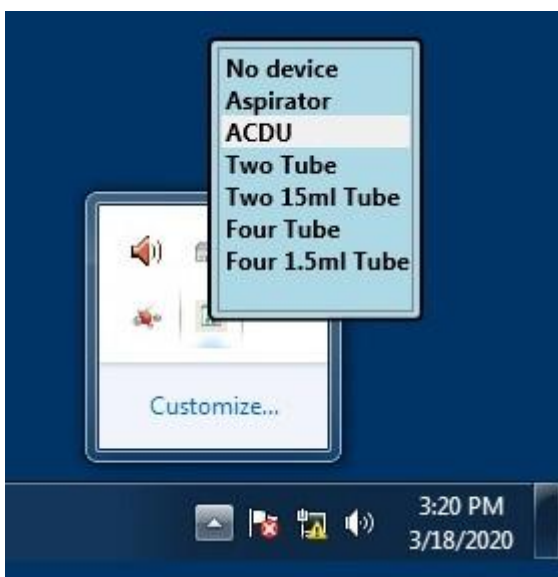
ACDU adaptor



2.2 Select the device (ACDU) on the lower right corner of the upper monitor



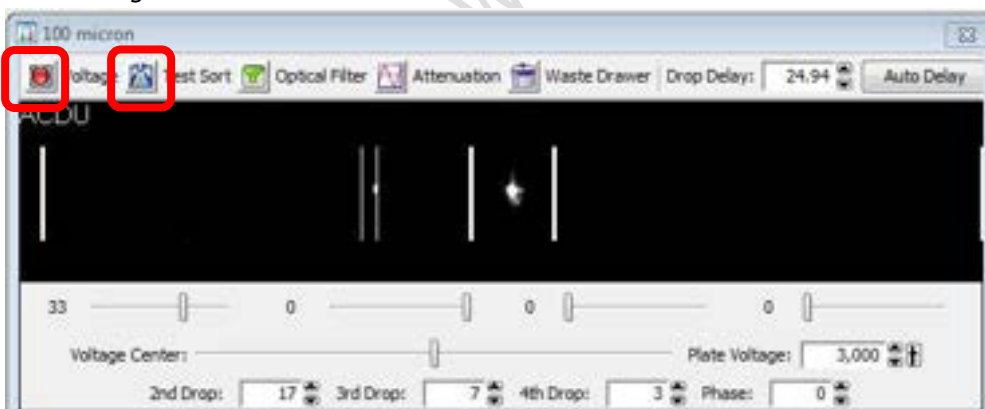
Imaging and Flow Cytometry Core



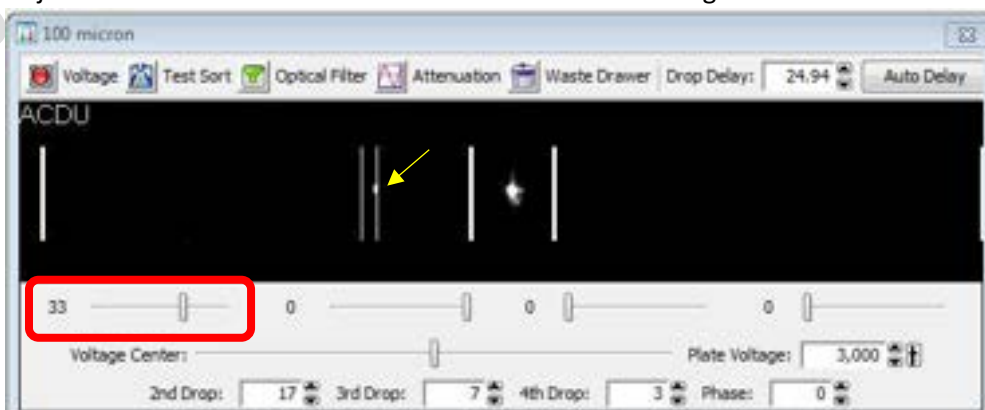
2.3 Go to 100 micron window (lower monitor). Adjust the Slider of the **far left**

| Device | Suggested Slider reading |
|---------------|--------------------------|
| 96-well plate | 33 - 0 - 0 - 0 |

2.4 Click *Voltage*. Wait 2 seconds and then Click *Test Sort*



2.5 Adjust the slider so the side stream dot lines within the target lines

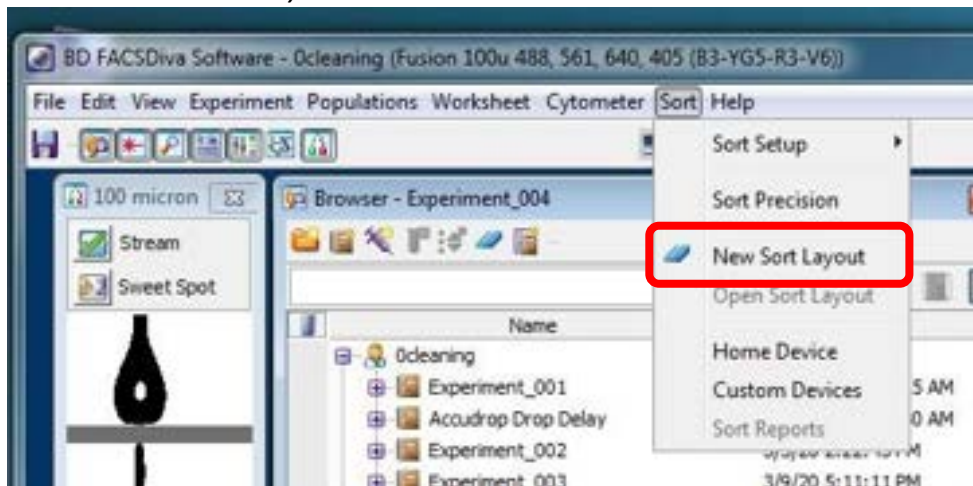




Imaging and Flow Cytometry Core

2.6 Click *Voltage* again to stop test sort

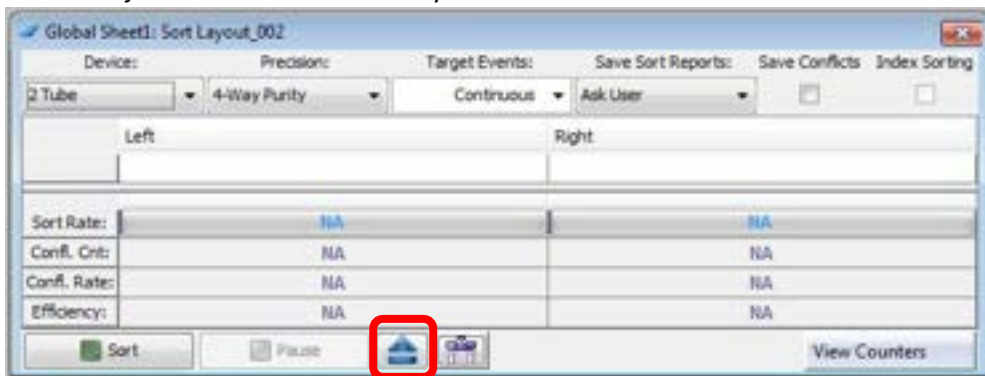
2.7 Click *Sort > New Sort Layout*



CPOS - Imaging and Flow Cytometry

Imaging and Flow Cytometry Core

2.8 Click the *Eject* button on the Sort Layout window



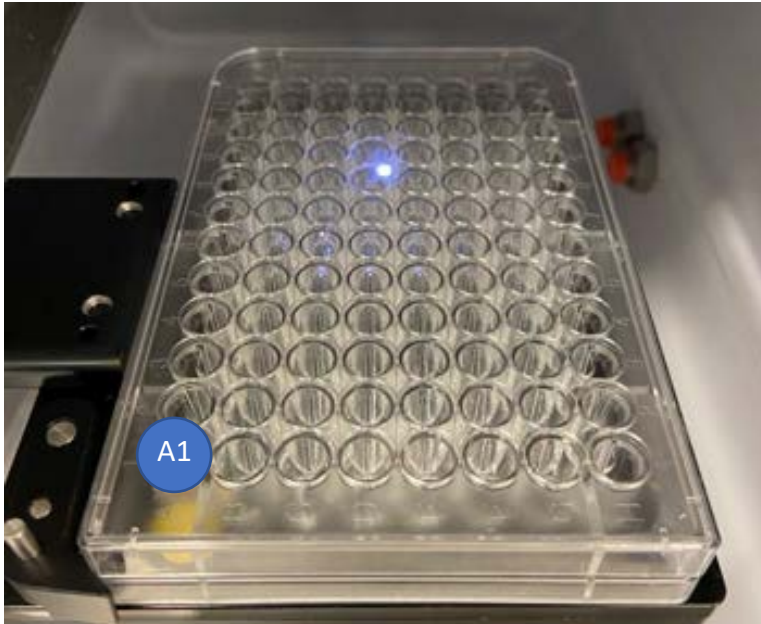
2.9 Plug in water tubing if cooling of the ACUDU stage is needed



2.10 Load a dummy plate on the ACUDU stage with A1 on the outer left corner

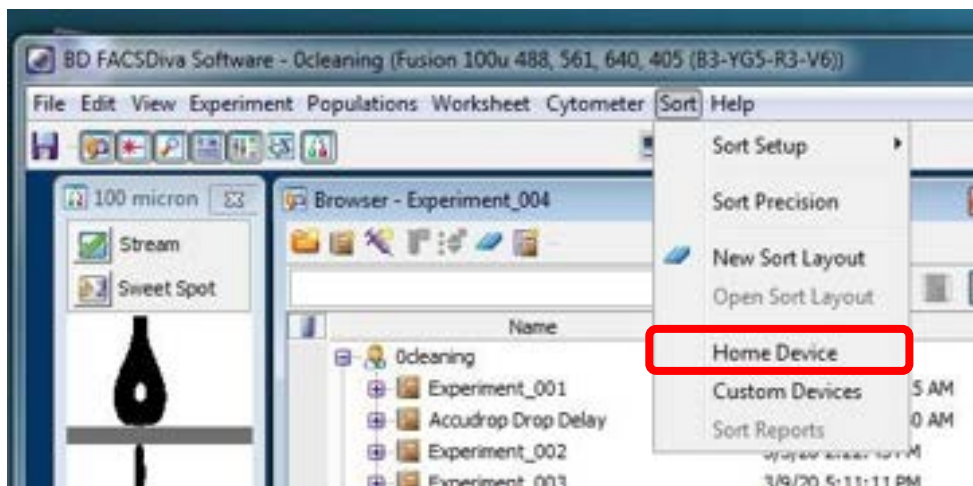


Imaging and Flow Cytometry Core

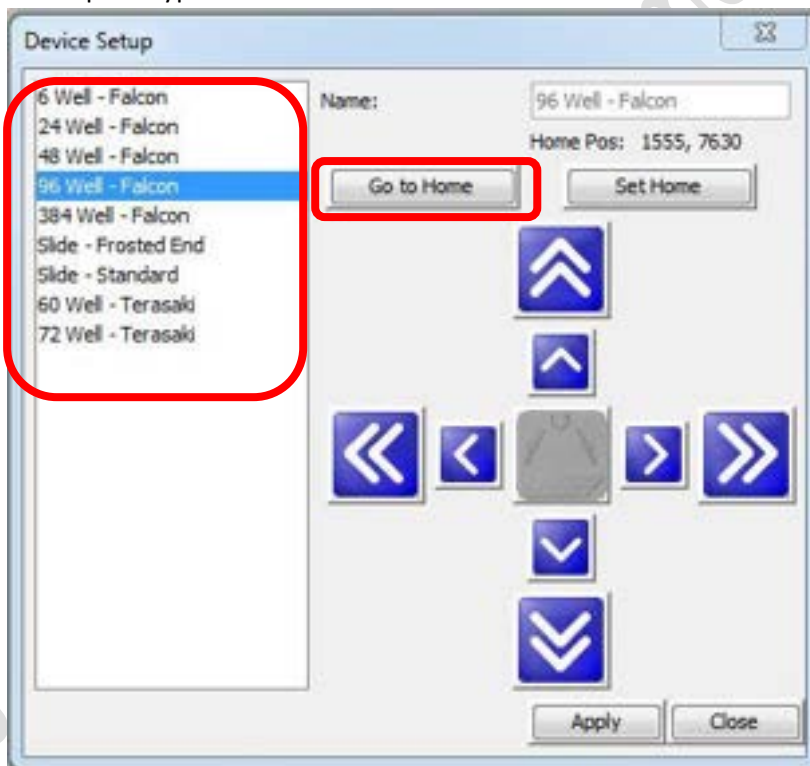


Imaging and Flow Cytometry Core

2.11 Click Sort > Home Device



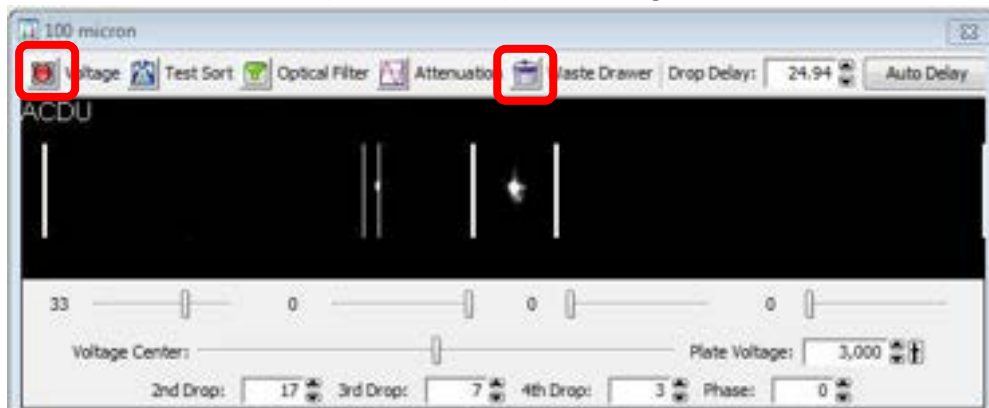
2.12 Select plate type and then click *Go to Home*



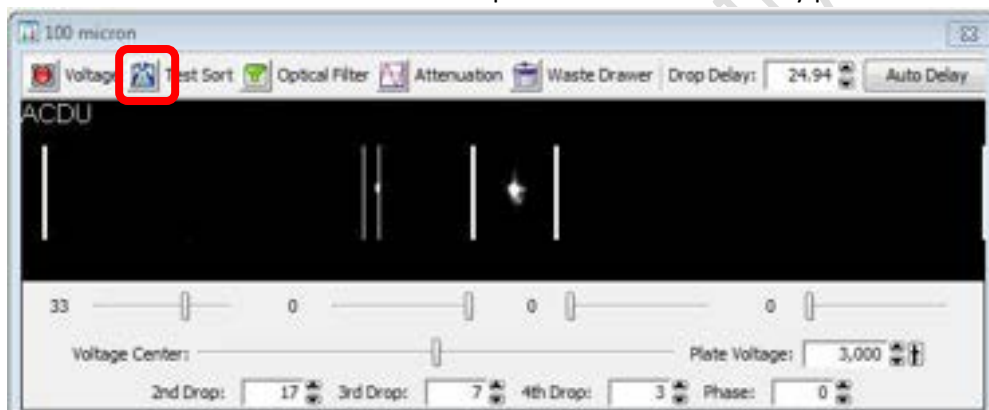


Imaging and Flow Cytometry Core

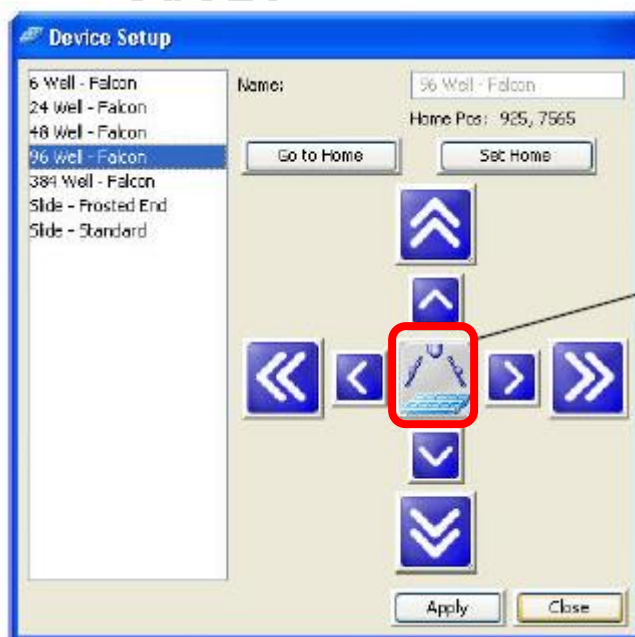
2.13 Go to 100 micron window (lower monitor), Click *Voltage* and *Waste Drawer*



2.14 Double Click *Test sort* to shot a small drop of sheath on the dummy plate cover

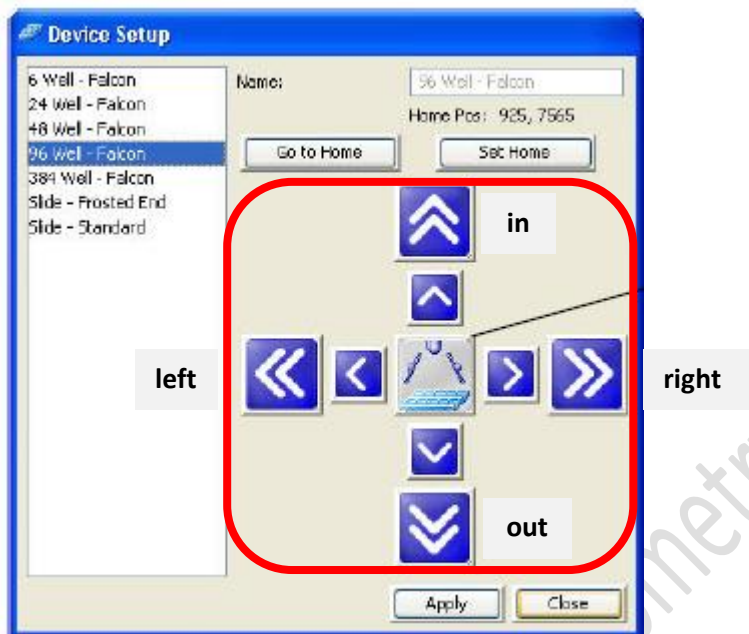


- OR -

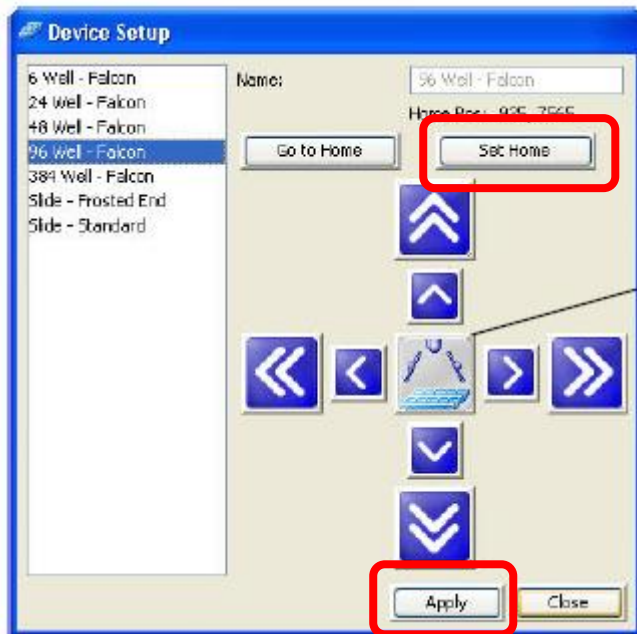


Imaging and Flow Cytometry Core

2.15 Move the stage accordingly in order to shot the drop of sheath on A1 position



2.16 Go to Home Device Window, Click *Set Home* and *Apply*

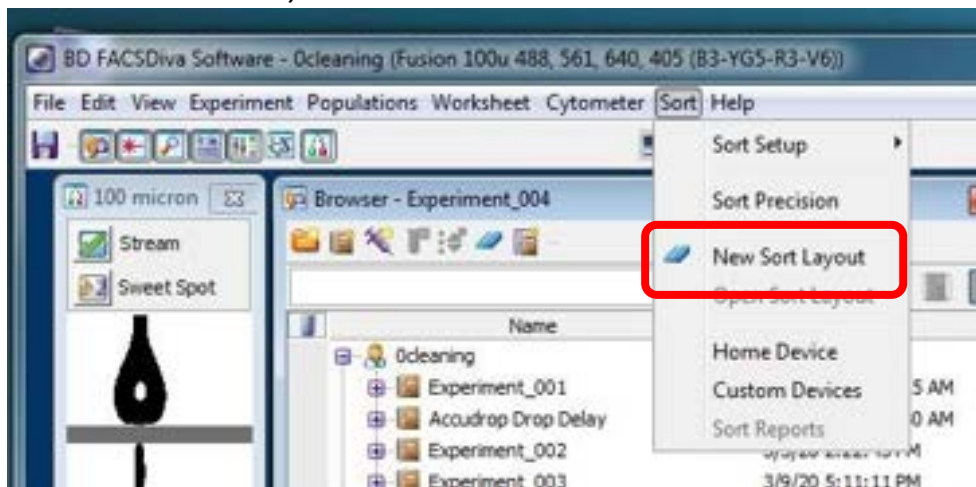


2.17 Remove Dummy plate from the stage and Load the collection plate

Imaging and Flow Cytometry Core

H. Sort Setup

1. Click Sort > *New Sort Layout*

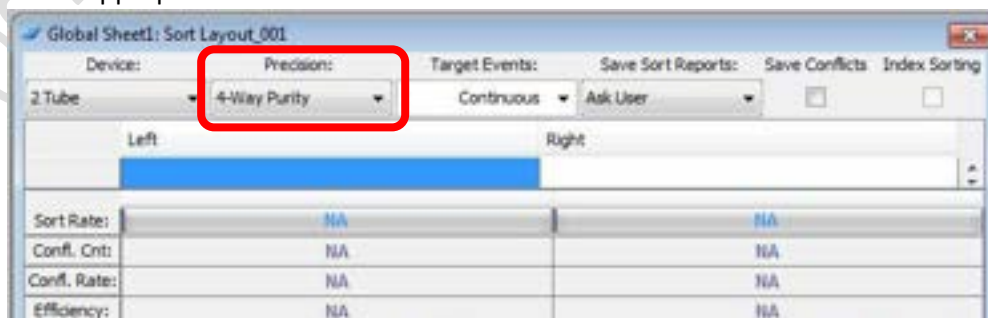


2. Select appropriate *Device*:



| Name | Supported Device |
|----------------|------------------------------------|
| 2 tube | 2-way 15 mL, 5.0 mL, 2.0 mL, 1.5mL |
| 4 tube | 4-way 5.0 mL, 2.0 mL, 1.5 mL |
| 96-well Falcon | 96-well culture plate |

3. Select appropriate *Precision*:



| Name | Suitable Application |
|--------|---|
| Purity | Sorting target population higher than 20% |
| Yield | Sorting target population less than 20% |

Imaging and Flow Cytometry Core

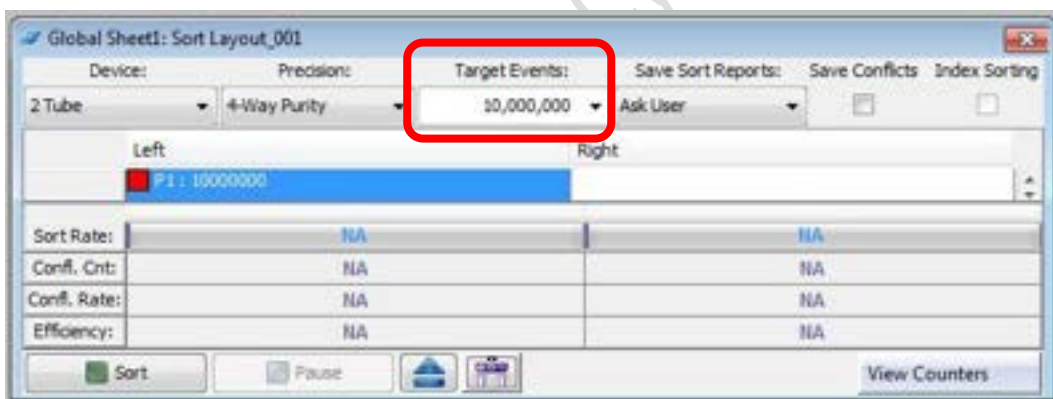
| | |
|--------------|---|
| Single Cell | Single cell sorting into 96-well plate / Single cell sequencing |
| 4-way Purity | 4-way sorting target population higher than 20% |

- Assign target population to position by *clicking the position > Add > Target gate*



- Input Target Events (sorting will stop when the sorted cell number reached the target event) for each target population if needed.

Select *Continuous* for unlimited number.



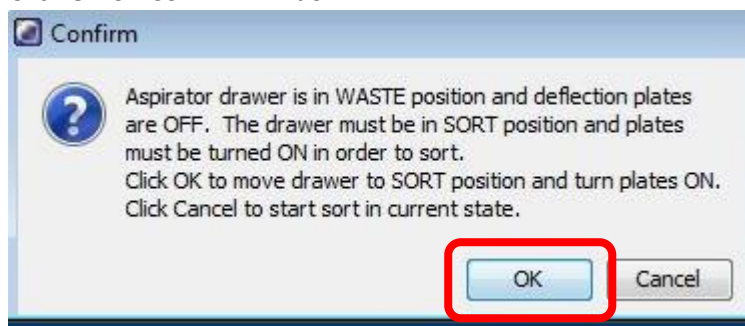
- Load your Sample onto the sample stage.
- Go to Acquisition Dashboard window, click *Load*

Imaging and Flow Cytometry Core

8. Go to Sort layout window, Click *Sort*



9. Click *OK* on Confirm window

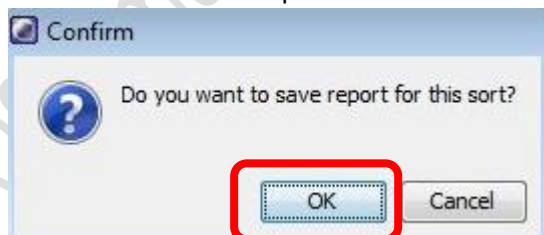


10. During the sort keep **monitoring Threshold Rate and Drop 1 value**

11. Click *Pause* if you wish to pause the sort and replace new collection tube. Click *Resume* after finish replacement.

12. Click *Sort* to end the sort

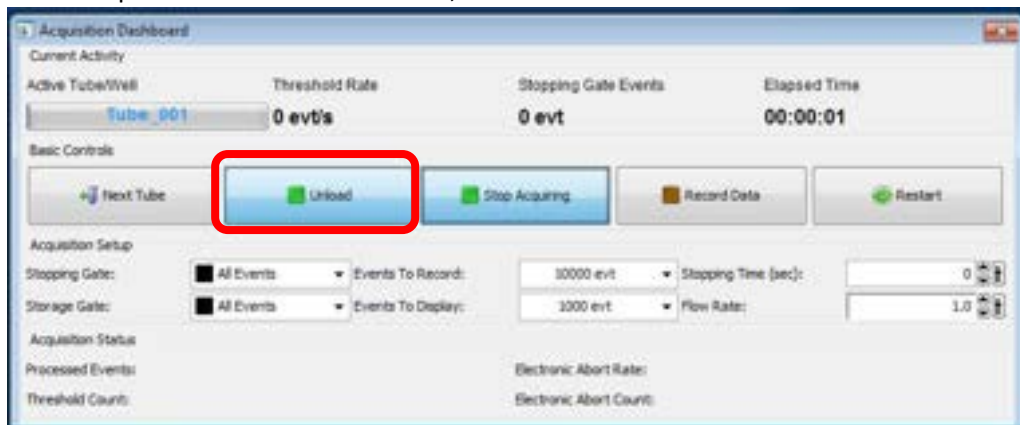
13. Click *OK* to save sort report





Imaging and Flow Cytometry Core

14. Go to Acquisition Dashboard window, click *Unload*.



CPOS - Imaging and Flow Cytometry



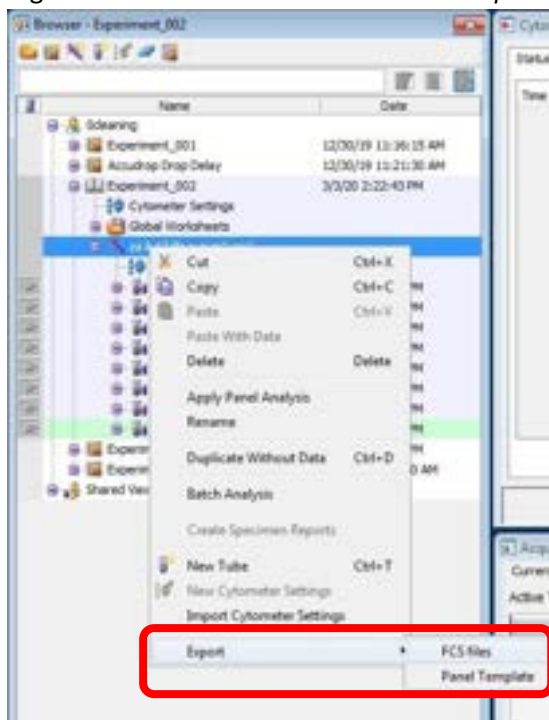
Imaging and Flow Cytometry Core

I. 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*



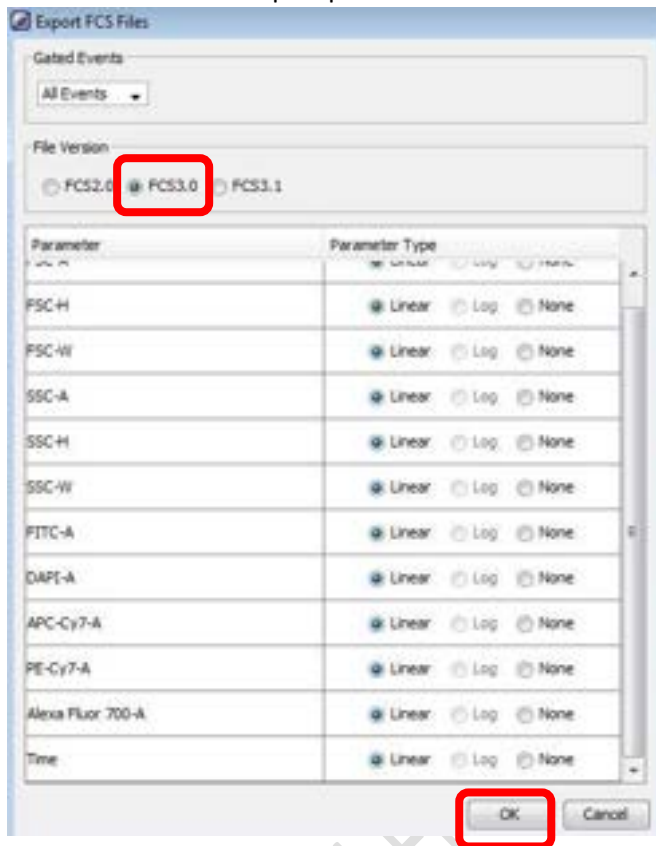
Cytometry Core

CPOS - Imaging and



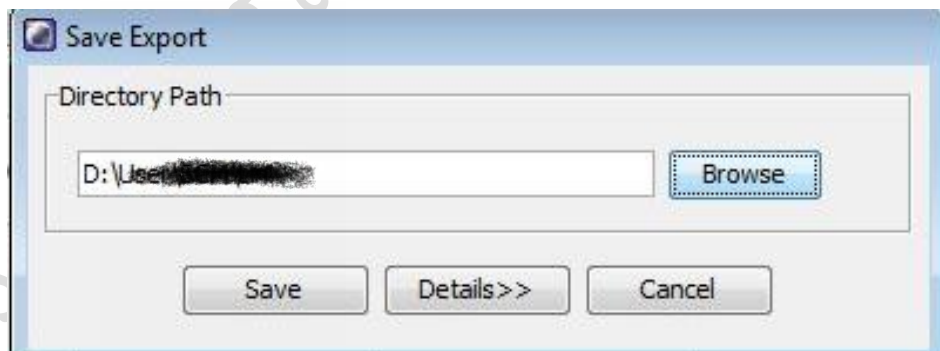
Imaging and Flow Cytometry Core

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*

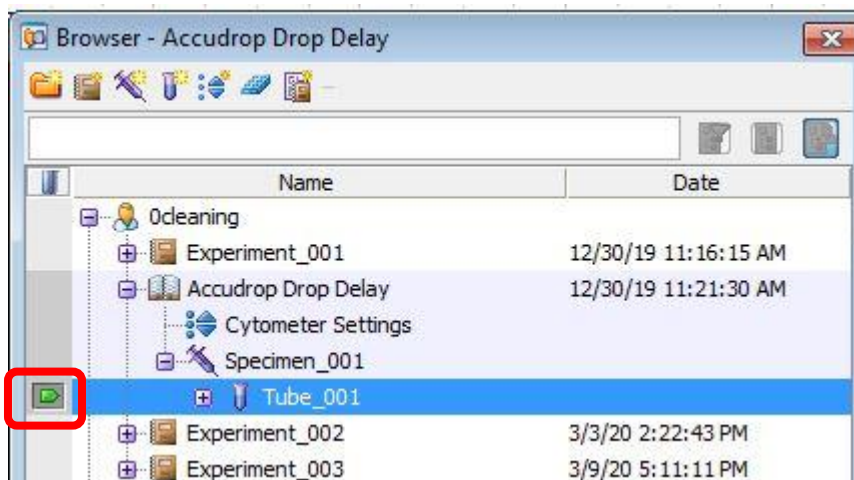


2. PDF file

2.1 To export pdf of single tube, Click the tube pointer of the selected tube



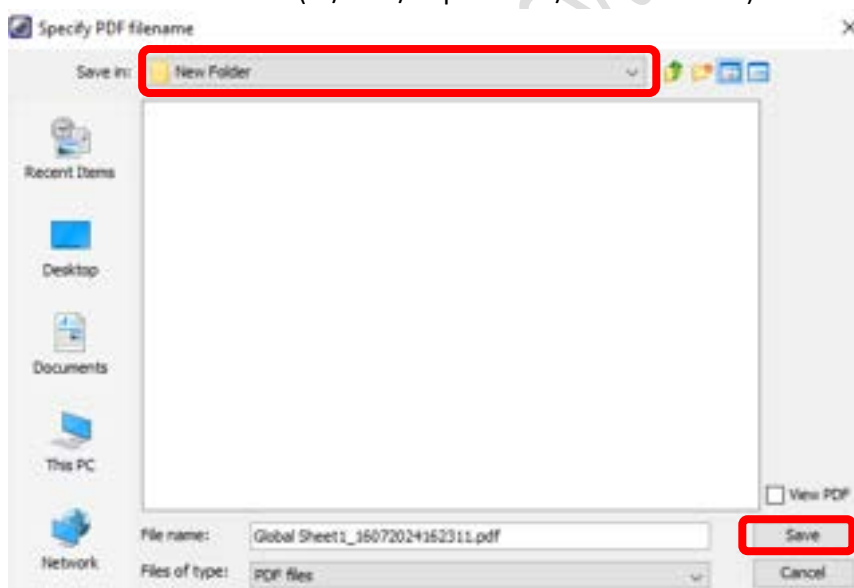
Imaging and Flow Cytometry Core



2.2 Click PDF icon



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

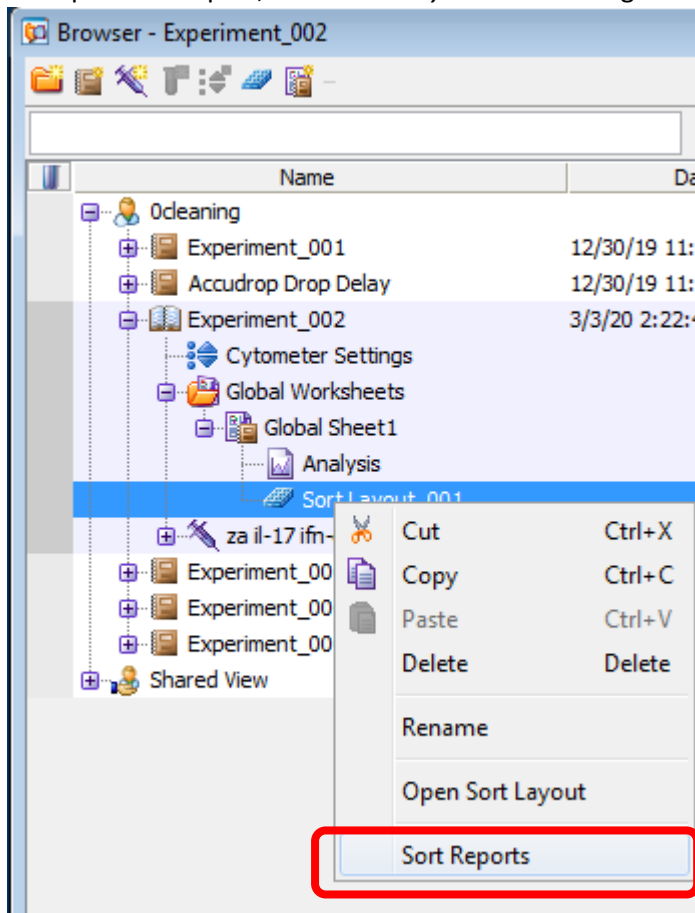


3. Sort Report

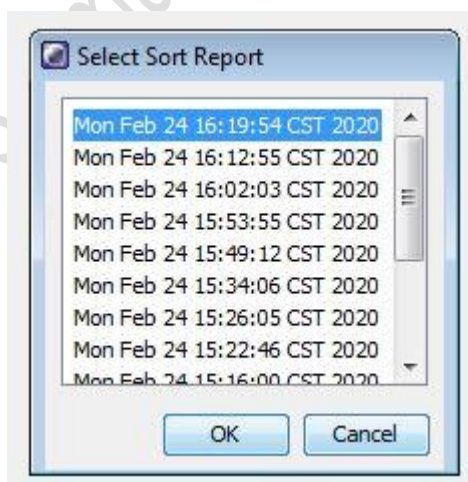


Imaging and Flow Cytometry Core

3.1 To export sort report, select *Sort Layout* and then right click. Click *Sort Reports*

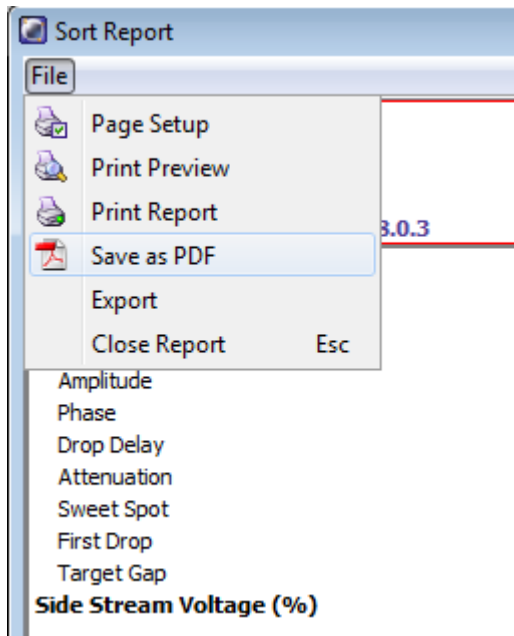


3.2 Select the sort from the list and then click *OK*



Imaging and Flow Cytometry Core

3.3 Click *File > Save As PDF*



3.4 Choose the destination (D:/User/Department/Personal Folder) and click *Save*

J. Cleaning

1. **TURN OFF** the *Sweet Spot*

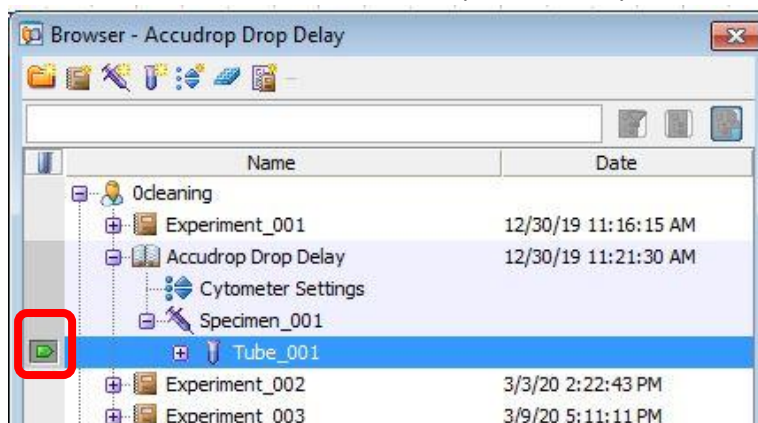


2. Open the upper flow cell access door of the system.



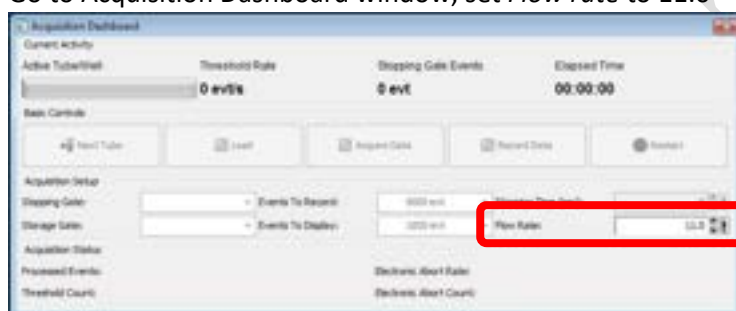
Imaging and Flow Cytometry Core

3. Go to Browser window, click the tube pointer of any tube

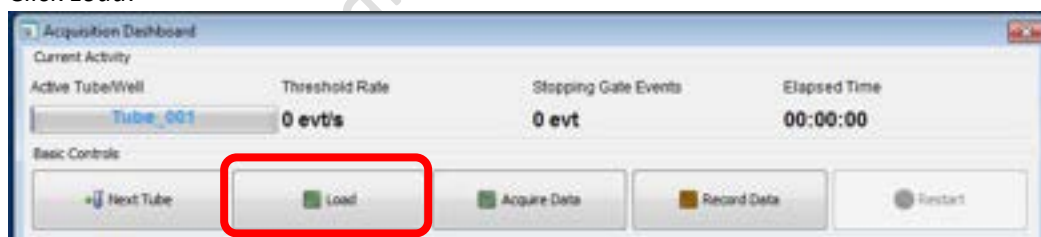


4. Load a tube of 2 mL of cleaning solution No. 1 (FACSClean) on the sample stage

5. Go to Acquisition Dashboard window, set *Flow rate* to 11.0



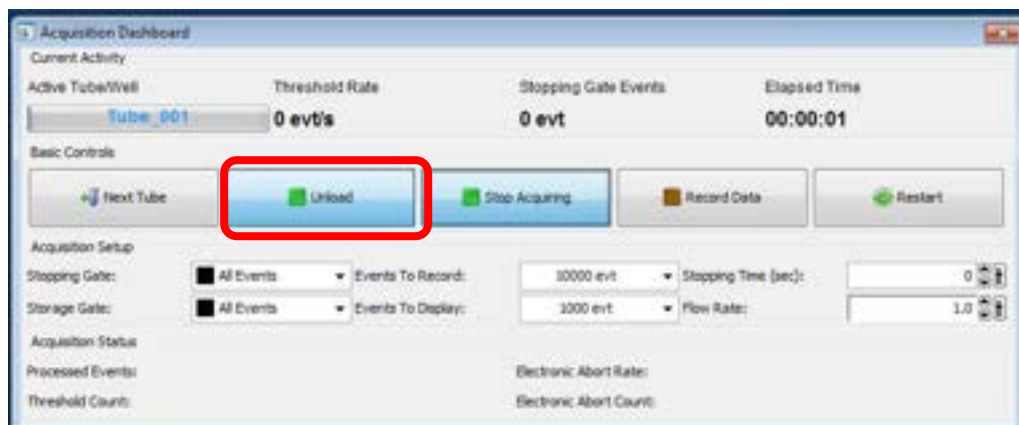
6. Click *Load*.



7. Acquire the solution for 5 minutes.

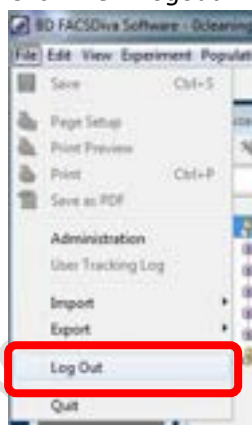
Imaging and Flow Cytometry Core

8. Click *Unload*.



9. Load a tube of 2 mL of cleaning solution No.2 (FACSRinse) on the sample stage
10. Repeat step 6 - 8. If Propidium Iodide (PI) is used, No.2 solution should be washed for 10 mins.
11. Load a tube of 2 mL of cleaning solution No.3 (MilliQ water) on the sample stage
12. Repeat step 6 - 8.
13. **User Logout**

Click *File > Logout*



14. Log out Tracker before leave

