BD Aria SORP Standard Operation Protocol – Basic Operation

A. Laser

- 1. Check if the laser needed is turned ON.
 - * Please turn **ON** the UV laser 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. Please contact our staff to confirm your UV laser setting.



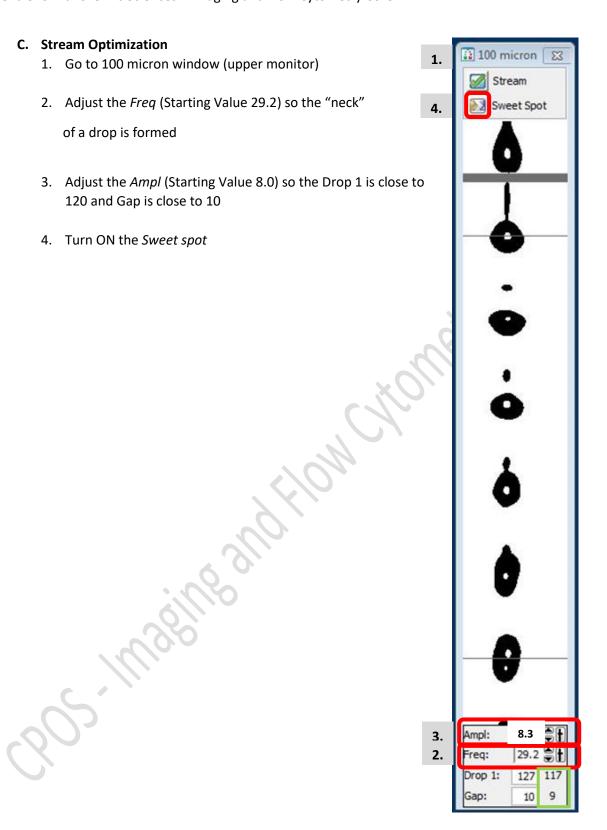
B. User Login

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



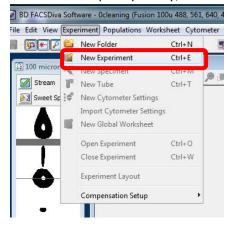
2. Click Use CST Settings if the window below pops up.



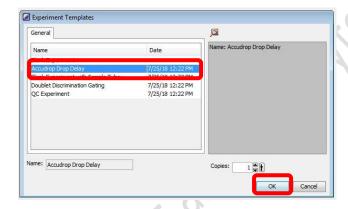


D. Accudrop Delay Assay

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

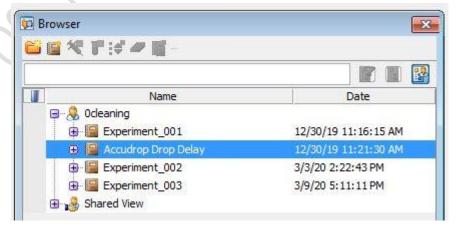




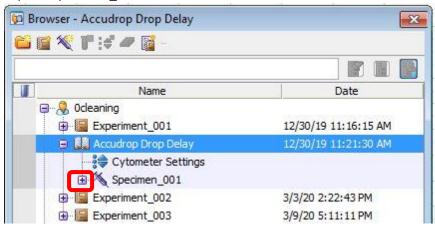


- OR -

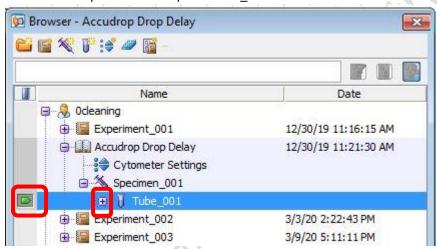
2. To open existing Accudrop Drop Delay experiment, double click *Accudrop Drop Delay* on the Browser window



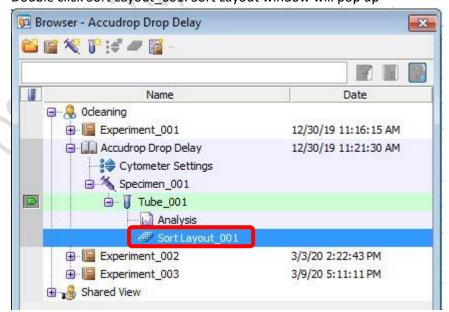
3. Expend Specimen_001



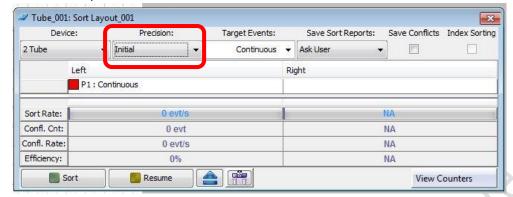
4. Click the tube pointer and expend Tube_001



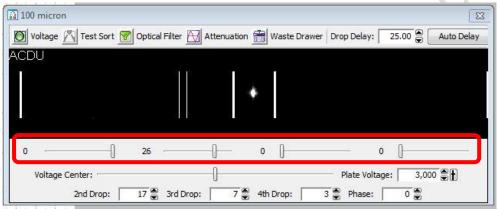
5. Double click Sort Layout_001. Sort Layout window will pop up



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



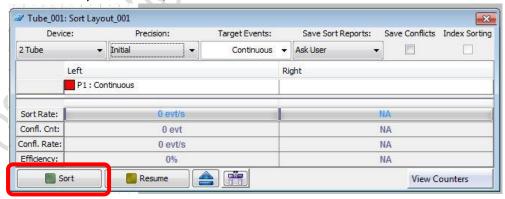
9. Close upper flow cell access door.



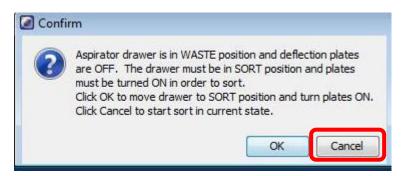
- 10. Load a tube of Accudrop beads (1 mL of PBS + 1 drop of stock) on the sample stage
- 11. Go to Acquisition Dashboard, Click Load



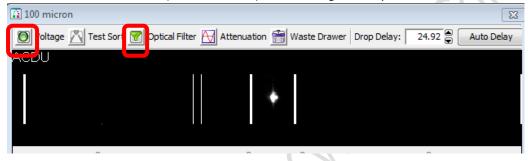
- 12. Adjust Flow rate if needed to obtain threshold rate constantly 1000-1500 events per sec
- 13. Go to Sort Layout window, click Sort



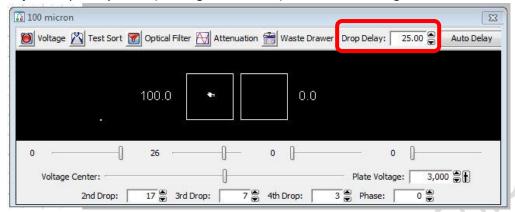
14. Click Cancel on the confirm window



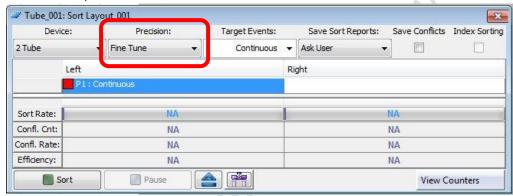
15. Go to 100 micron window (lower monitor), Click Voltage and Optical Filter



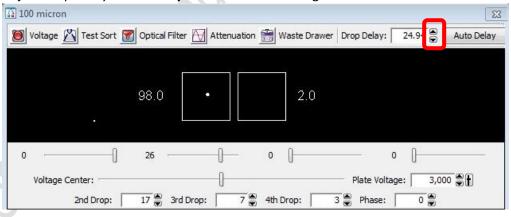
16. Adjust Drop Delay value (starting value: 25.00) so that the reading on the left reach 100



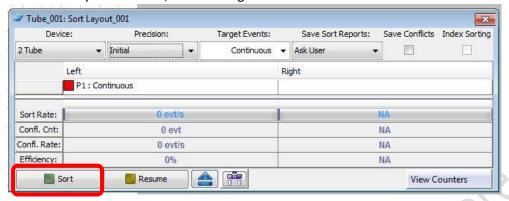
17. Go to Sort Layout window, Select Precision > Fine Tune



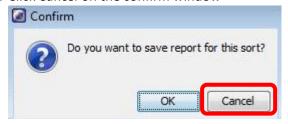
18. Adjust Drop Delay value bit by bit so that the reading on the left reach >97



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



20. Click Cancel on the confirm window



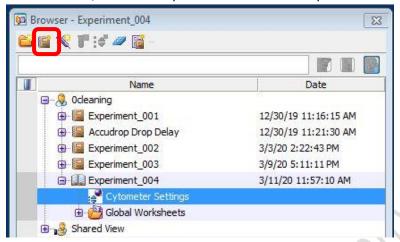
21. Go to Acquisition Dashboard, Click Unload



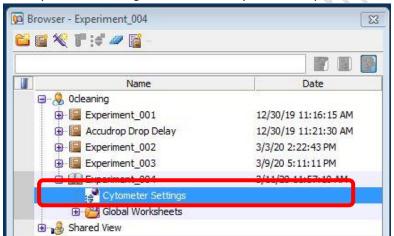
22. Return the Accudrop beads to 4 degree refrigerator

E. Experiment Setup

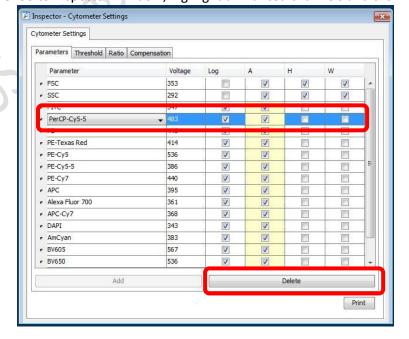
- 1. Setup New Experiment
 - 1.1 Go to Browser, Click New Experiment icon. A new experiment will be created



1.2 Click Cytometer Settings under the newly created Experiment



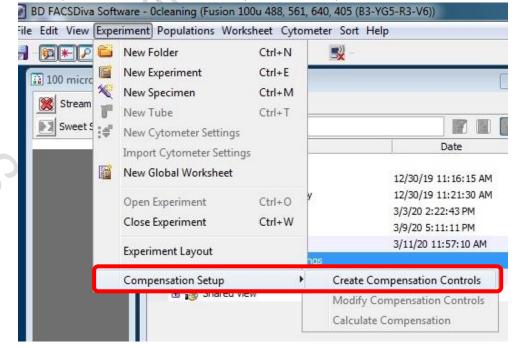
1.3 Go to Inspector Window, highlight unwanted channels and click Delete.



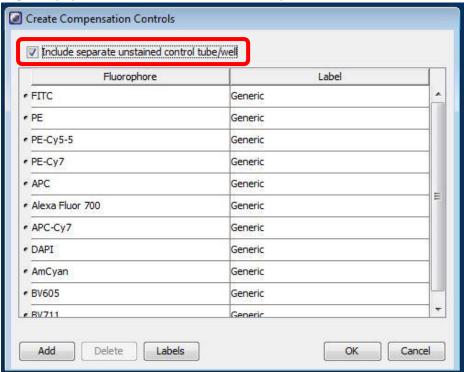
🗾 Inspector - Cytometer Settings Cytometer Settings Parameters Threshold Ratio Compensation Parameter Voltage Log FSC 353 SSC 292 1 PerCP-Cy5-5 V 446 V PE-Texas Red 414 # PE-Cy5 536 1 V PE-Cy5-5 386 PE-Cy7 • APC 395 Alexa Fluor 700 361 APC-Cy7 368 1 1 DAPI 343 AmCyan 383 1 567 • BV650 536 7 Add Delete Print

1.4 Check H and W boxes of FSC and SSC

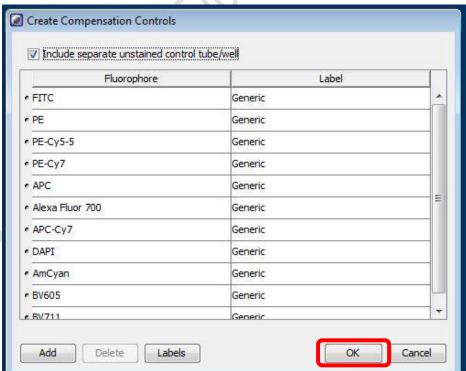
- 1.5 Keep *Log* boxes of FSC and SSC **unchecked**
- 1.6 Keep Log boxes of all fluorescence channels checked
 - * If you are doing **cell cycle or DNA content** analysis, please keep **log box** of your DNA specific fluorescence channel **unchecked**.
- 2. Setup Compensation (for Multi-colour panel)
 - 2.1 Click Experiment > Compensation Setup > Create Compensation Control



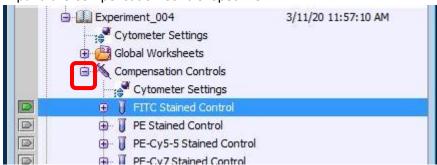
2.2 If any one of your single stain controls is known to be 100% positive, i.e. no negative population, check the box "Include separate unstained control tube/well".



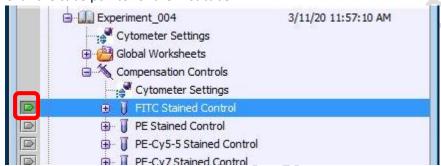
2.3 Click OK



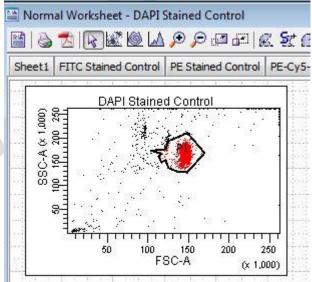
2.4 Expand the Compensation Control Specimen



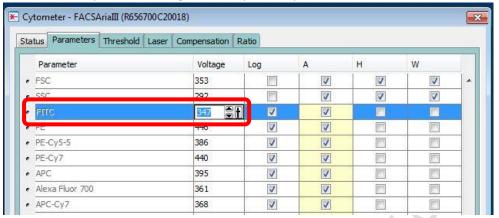
2.5 Click the tube pointer of the first tube



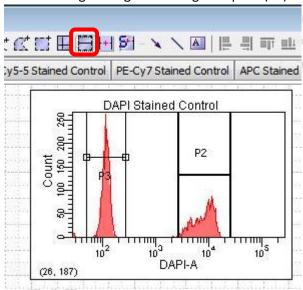
- 2.6 Load the single stain controls on the sample stage according to the tube label, i.e. run FITC single stain when the tube pointer is pointing at "FITC Stained Control"
- 2.7 Go to Acquisition Dashboard, Click Load.
- 2.8 Go to Normal Worksheet, move the P1 gate to include major cluster. Adjust FSC and SSC PMT Voltage if needed



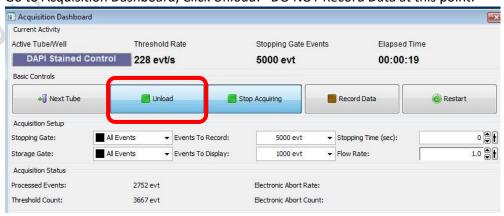
2.9 Go to Cytometer window, Fine tune the corresponding fluorescence PMT voltage to have best separation of negative and positive peak



- 2.10 Move the interval gate (P2) to include the positive peak
- 2.11 Use interval gate to gate out negative peak (P3)

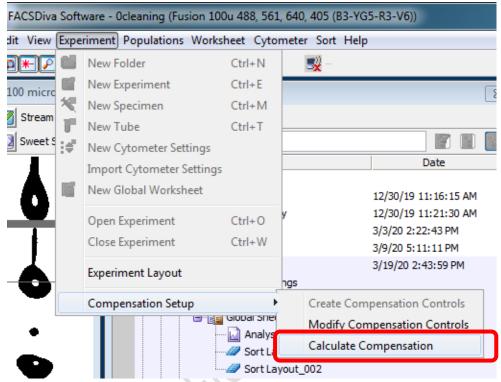


2.12 Go to Acquisition Dashboard, Click Unload. *DO NOT Record Data at this point.

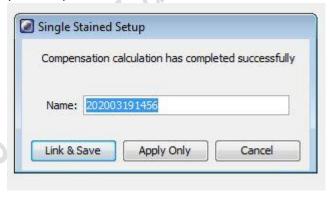


2.13 Repeat step 2.5 - 2.12 with all the single stain controls.

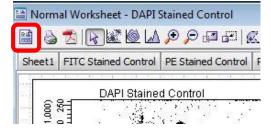
- 2.14 After optimising the PMT voltage of ALL the fluorescence channel, load each single stained control and click *Record Data* for ALL single stain controls
- 2.15 Click Experiment > Compensation Setup > Calculate Compensation



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.

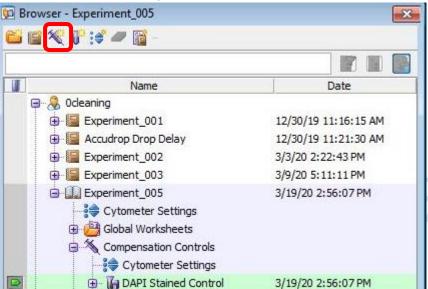


2.17 Switch Normal worksheet to Global worksheet

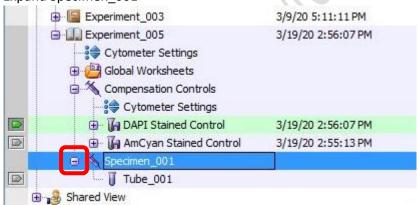


3. Setup Plots and Tables

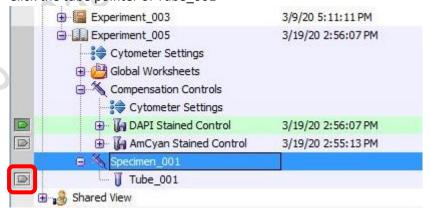
3.1 Go to Browser, Click New Specimen icon



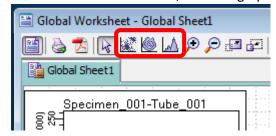
3.2 Expand Specimen 001



3.3 Click the tube pointer of Tube 001

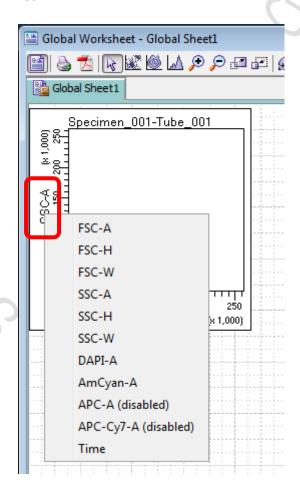


3.4 Go to Global Sheet Window, Click the graph type icon



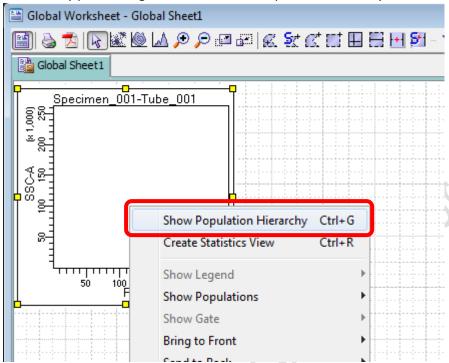
Icon	Туре
	Dot Plot
	Contour Plot
	Histogram

- 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.

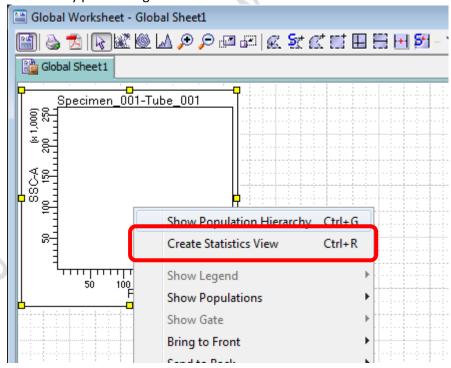


- 3.7 Repeat step 3.4 3.6 until all plots needed is created.
 - * Essential Plots: FSC-A VS SSC-A; FSC-H VS FSC-W; SSC-H VS SSC-W

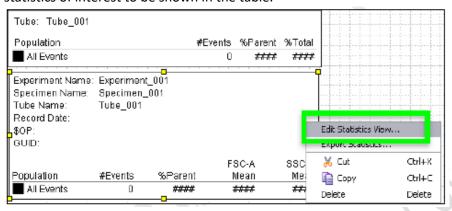
3.8 Click on any plot and right click. Click Show Population Hierarchy



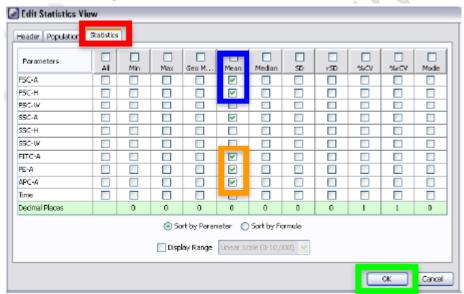
3.9 Click on any plot and right click. Click Create Statistics View



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.



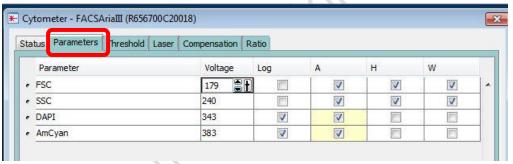
3.11 Click Statistics Tab, check the boxes of interested statistics and then click OK



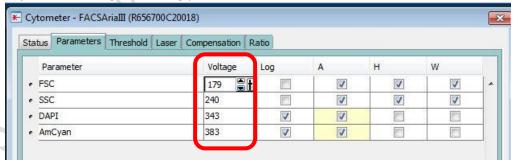
- 4. Sample Acquisition
 - 4.1 Load your sample on the sample stage
 - 4.2 Go to Acquisition Dashboard, Click Load



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

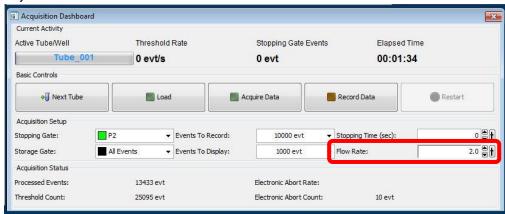


4.4 Adjust PMT Voltage accordingly



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

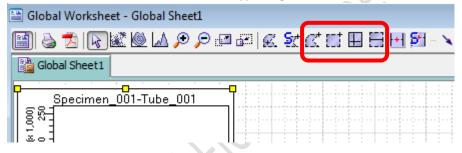
4.5 Adjust Flow rate if needed



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

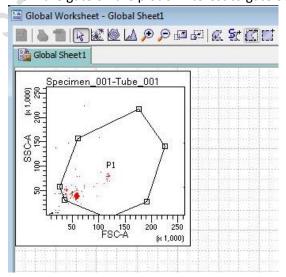
5. Create Gates

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

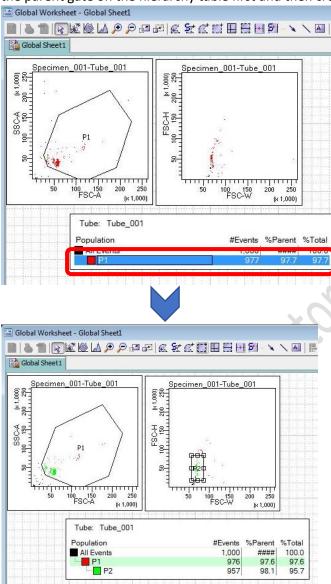


Icon	Туре
Œ	Polygon Area Gate
ET 6 0	Rectangle Area Gate
	Quandrant Gate
	Interval Gate

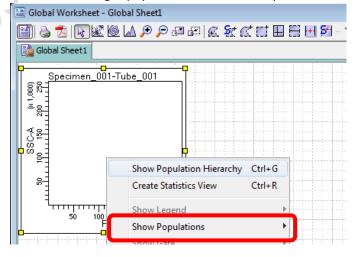
5.2 Draw the gate on the plot of interest to gate out target cluster /peak



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.



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

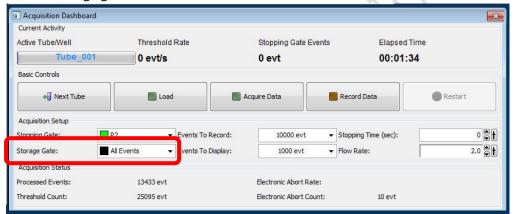


6. Data Recording

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



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

F. Sort Device Alignment

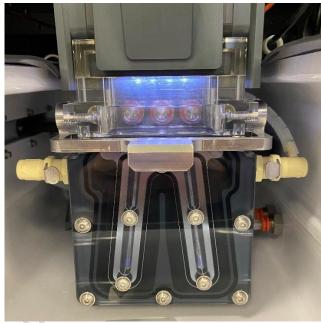
- 1. Tube Holder
 - 1.1 Assemble the tube holder as picture below





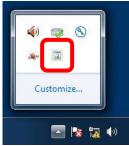


- 1.2 Put dummy tubes into the position
- 1.3 Slide the tube holder right under the sort chamber. Plug in water tubing if cooling is needed.

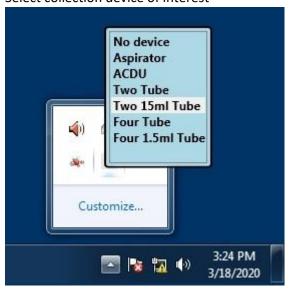


1.4 Click Select Device icon on the lower right corner of the upper monitor





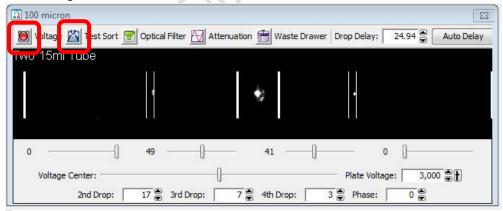
1.5 Select collection device of interest



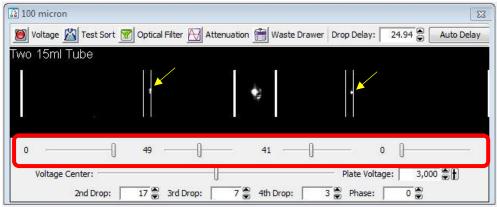
1.6 Go to 100 micron window (lower monitor). Adjust the Slider

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

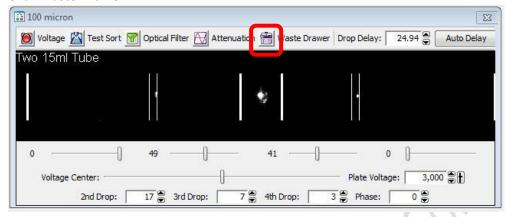
1.7 Click Voltage. Wait 2 seconds and then Click Test Sort.



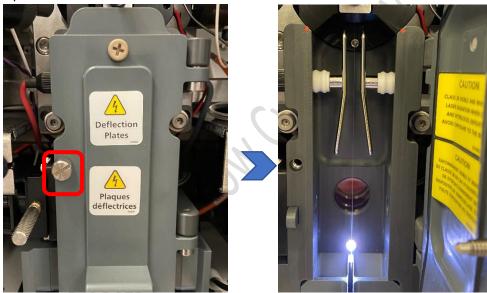
1.8 Adjust the slider so the side stream dot lines within the target lines



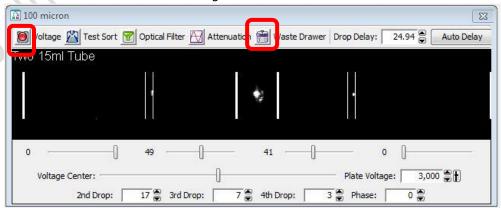
1.9 Click Waste Drawer



1.10 Open the sort chamber door



- 1.11 Check if the side streams can enter the dummy tubes. Adjust the slider if needed.
- 1.12 Click Waste Drawer and then Voltage



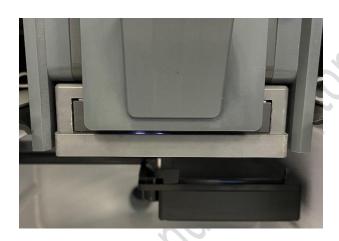
- 1.13 Close the sort chamber door
- 1.14 Slide out the tube holder and Put collection tubes into position

2. ACDU (Culture plate)

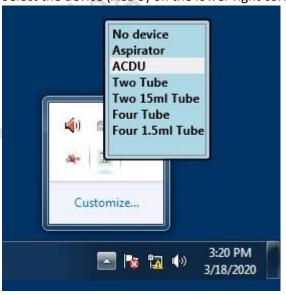
2.1 Slide the ACDU adaptor right under the sort chamber



ACDU adaptor



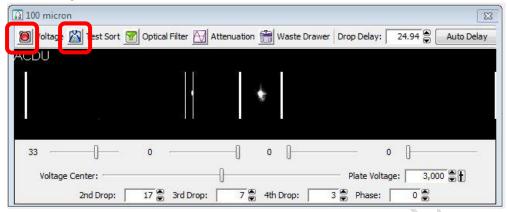
2.2 Select the device (ACDU) on the lower right corner of the upper monitor



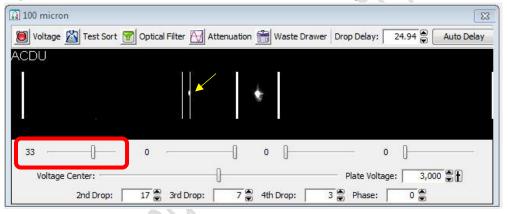
2.3 Go to 100 micron window (lower monitor). Adjust the Slider

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

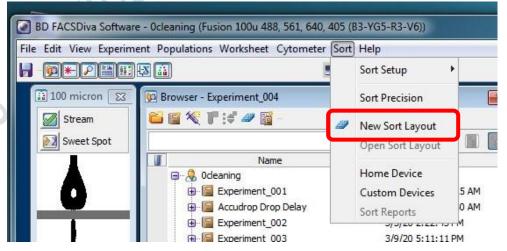
2.4 Click Voltage. Wait 2 seconds and then Click Test Sort



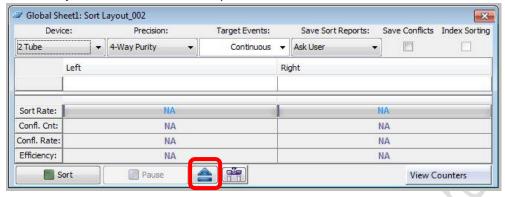
2.5 Adjust the slider so the side stream dot lines within the target lines



- 2.6 Click Voltage again to stop test sort
- 2.7 Click Sort > New Sort Layout



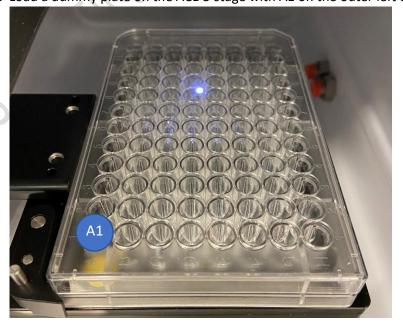
2.8 Click the Eject button on the Sort Layout window



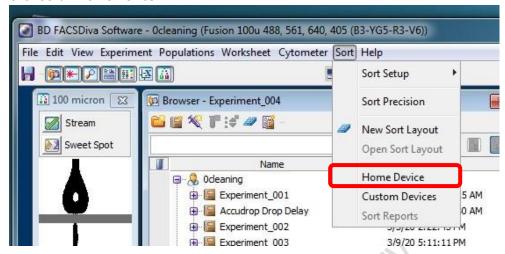
2.9 Plug in water tubing if cooling of the ACDU stage is needed



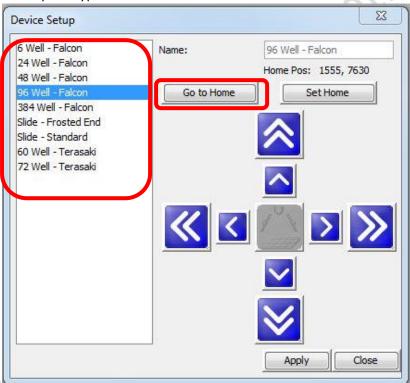
2.10 Load a dummy plate on the ACDU stage with A1 on the outer left corner



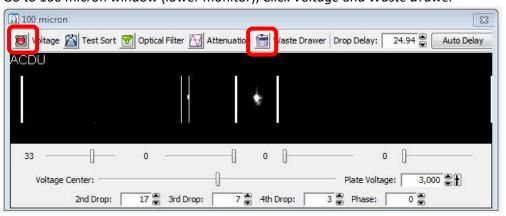
2.11 Click Sort > Home Device



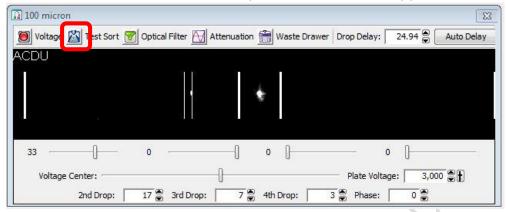
2.12 Select plate type and then click Go to Home



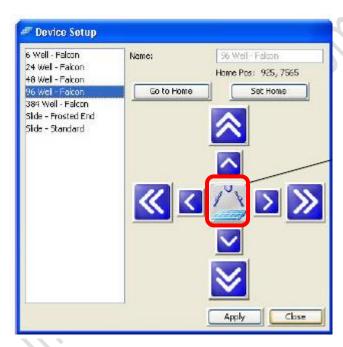
2.13 Go to 100 micron window (lower monitor), Click Voltage and Waste Drawer



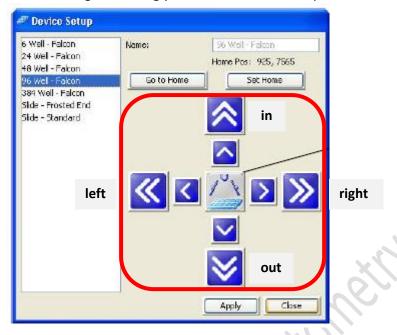
2.14 Double Click Test sort to shot a small drop of sheath on the dummy plate cover



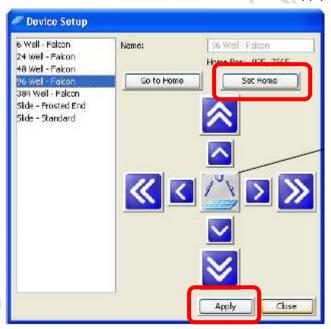
- OR -



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



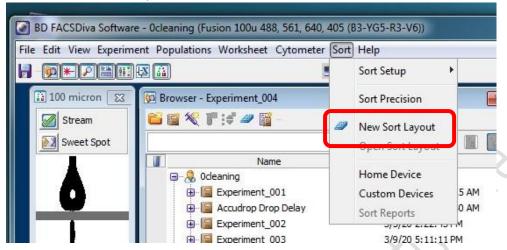
2.16 Go to Home Device Window, Click Set Home and Apply



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

G. Sort Setup

1. Click Sort > New Sort Layout

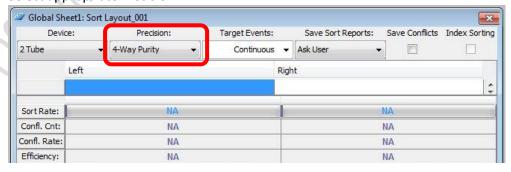


2. Select appropriate *Device*:



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*:



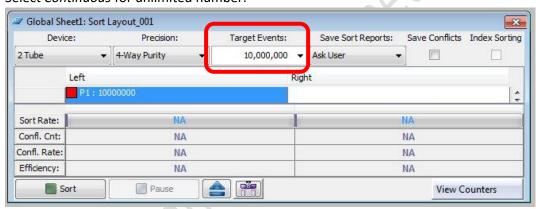
Name	Suitable Application
Purity	Sorting target population higher than 20%
Yield	Sorting target population less than 20%
Single Cell	Single cell sorting into 96-well plate / Single cell sequencing
4-way Purity	4-way sorting target population higher than 20%

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

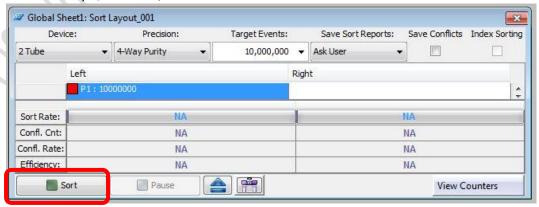


5. Input Target Events (sorting will stop when the sorted cell number reached the target event) for each target population if needed.

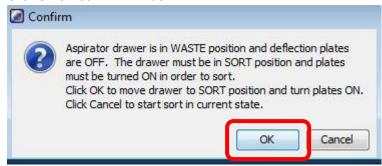
Select Continuous for unlimited number.



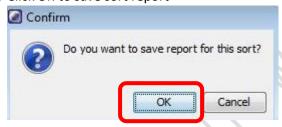
- 6. Load your Sample onto the sample stage.
- 7. Go to Acquisition Dashboard window, click Load
- 8. Go to Sort layout window, Click Sort



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

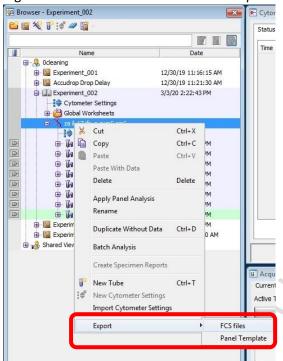


14. Go to Acquisition Dashboard window, click Unload.

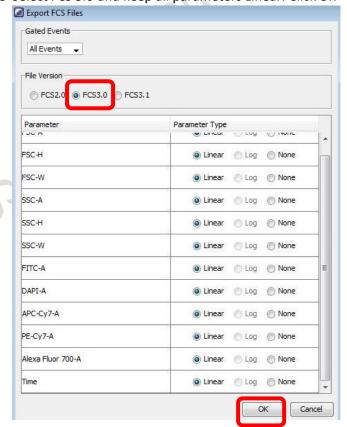


H. Data Export

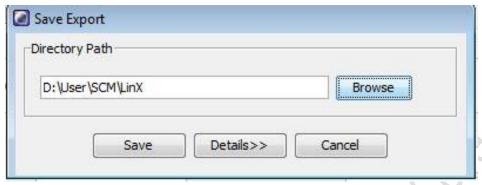
- 1. FCS file
 - 1.1 Go to Browser window, Select the Tubes / Specimen of interest.
 - 1.2 Right Click over the selection and click Export > FCS file



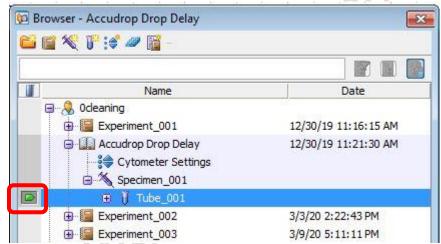
1.3 Select FCS 3.0 and keep all parameters Linear. Click OK



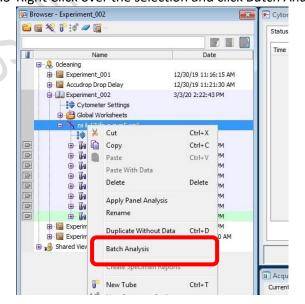
- 1.4 Click Browse to choose the destination (D:/User/Department/Personal Folder)
- 1.5 Click Save



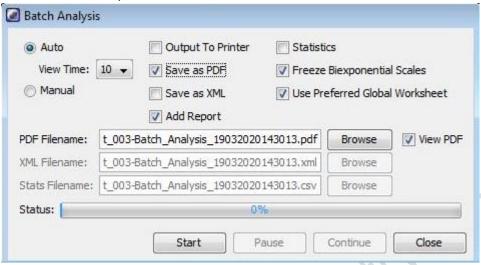
- 2. PDF file
 - 2.1 To export pdf of multiple tubes, Click the tube pointer of any tubes



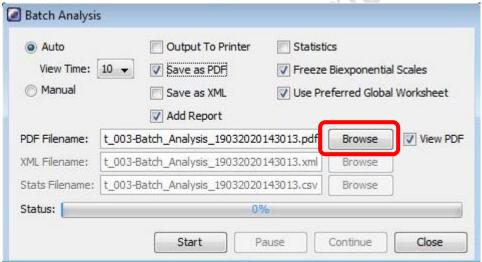
- 2.2 Select the wanted tubes/ Specimen.
- 2.3 Right Click over the selection and click Batch Analysis



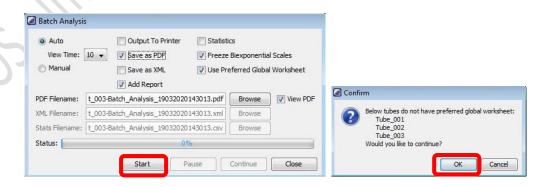
2.4 Check the boxes as picture below



2.5 Click Browse to choose the destination (D:/User/Department/Personal Folder)

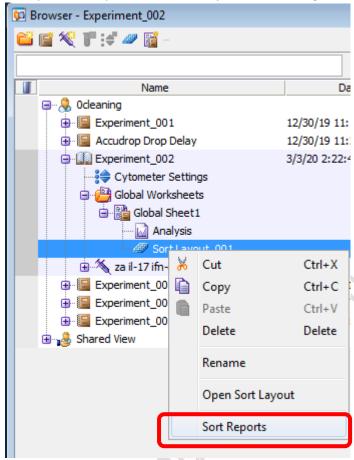


2.6 Click Start and then OK.

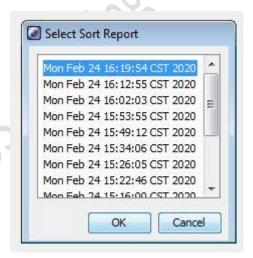


3. Sort Report

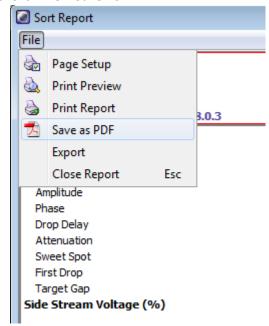
3.1 To export sort report, select Sort Layout and then right click. Click Sort Reports



3.2 Select the sort from the list and then click OK



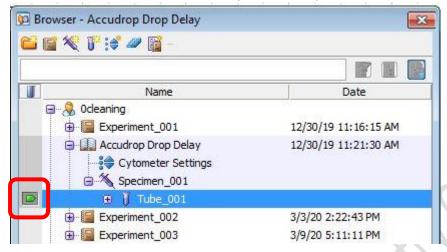
3.3 Click File > Save As PDF



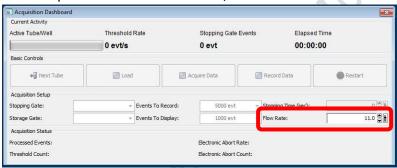
3.4 Choose the destination (D:/User/Department/Personal Folder) and click Save

I. Cleaning

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



- 2. Load a tube of 2 mL of cleaning solution No. 1 (FACSClean) on the sample stage
- 3. Go to Acquisition Dashboard window, set Flow rate to 11.0



4. Click Load.



5. Acquire the solution for **5 minutes**.

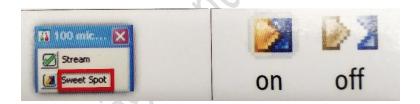
6. Click Unload.



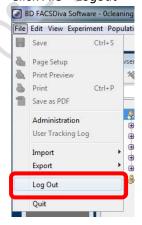
- 7. Load a tube of 2 mL of cleaning solution No.2 (FACSRinse) on the sample stage
- 8. Repeat step 3-5. If Propidium Iodide (PI) is used, No.2 solution should be washed for 10 mins.
- 9. Load a tube of 2 mL of cleaning solution No.3 (MilliQ water) on the sample stage
- 10. Repeat step 3-5.

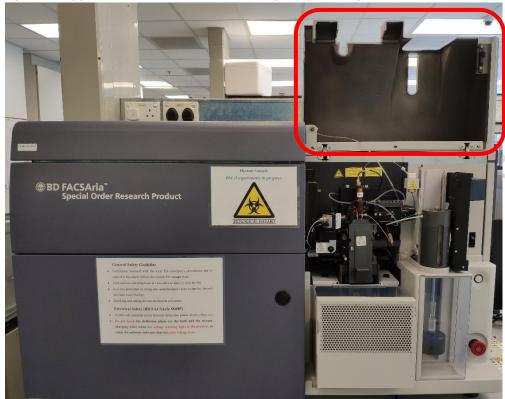
J. User Logout

1. TURN OFF the Sweet Spot



2. Click File > Logout





3. Open the upper flow cell access door of the system before you leave.