

## CytoFLEX SRT Experimental setup form - Initial setup

Updated: 06-20-2023

### **Auburn University Flow Cytometry and High-Speed Cell Sorting Laboratory**

**Location:** 250A Greene Hall, Auburn, AL 36849

**Contact:** [rzw0041@auburn.edu](mailto:rzw0041@auburn.edu) (Rie Watanabe, Operator & Manager) [birdric@auburn.edu](mailto:birdric@auburn.edu) (Dr. Bird, Director)

**Lab phone#:** 334-844-2711

**Operation hours:** 800am-445pm (M-F)

**\*\*Please read through this document and contact us with any questions before you submit sorting request\*\***

#### **Scheduling:**

- Sorting can be started between the hours of 900am – 330pm. If sorting outside of the regular hours is required, the request must be discussed and approved before the submission.
- A minimum of 30 minutes before or between sorts is required to clean the machine
- A minimum of 1 hour is required before or between sorts is required if aseptic cleaning is requested
- For scheduling, email [rzw0041@auburn.edu](mailto:rzw0041@auburn.edu). Sorting must be scheduled a minimum of 1 working day before the requested appointment time to ensure the scheduling

#### **Sample requirement and preparation tips:**

- Minimum volume is 0.5ml even if this volume does not give the ideal cell concentration
- Desired concentration of cells is  $1\sim 10 \times 10^6$  cells/mL (after filtration)
- Samples are supposed to be single cell suspension only (NO CLUMPS). Should be filtered through cell strainer to make good single cell suspension (increases sort efficiency!) and prevent clogging
- Recommended sorting buffers are 1xPBS (free of  $\text{Ca}^{2+}/\text{Mg}^{2+}$ ), Hank's. Buffers can be supplemented with BSA (0.1-1%) or dialyzed FBS (1-5%). Addition of EDTA (up to 5mM) or DNaseI (25-50ug/ml, along with  $\text{MgCl}_2$ ) may help preventing cells clumping via DNA released from dead cells
- Always prepare the unstained control. Cell numbers don't have to be as high as your sorting samples but it is essential to set the gates
- If your samples are stained with more than 1 fluorochrome, prepare samples stained with each fluorochrome individually. This is for compensation purposes

#### **Sample and Collection containers:**

- Prepare samples in polypropylene 5ml (12x75mm) round bottom tubes suit our instrument best
- Sorted cells can be collected in 15ml conical tube, 5ml (12x75mm) tube, or 96-well plate
- Prepare your collection containers pre-loaded with receiving solutions (media etc)

#### **Expectations for cell sorting:**

- Sorting time depends on your objective and sample quality. If you need more information, see Appendix or contact us
- Expect the final yield is about 75% of the starting number of desired cells. For example,  $1 \times 10^7$  cells with 30% GFP(+) would yield  $2.25 \times 10^6$  GFP(+) cells ( $10^7 * 0.3 * 0.75$ ). Rare event sort (1% or less) can produce a lower yield, as low as 50%. It will be diluted as well since sorted cells takes up sheath

**Submitter information**

<b>Date:</b>	<b>Start/Finish:</b>	
<b>Researcher:</b>	<b>Contact information</b>	<b>Email:</b>
<b>Lab/PI:</b>		<b>Phone:</b>
<b>Department/College/Institution:</b>		
<b>Billing contact:</b>		
<b>If outside of the Auburn University, Street address and City:</b>		

**Sample information (check all applied)**

*Cell type:*  Cell line  Primary cells  Other \_\_\_\_\_

*Species:*  Human  Non-human primate  Mouse  Rat  Other \_\_\_\_\_

*Tissue Origin:*  Bone Marrow  Peripheral Blood  Neural  Other \_\_\_\_\_

*Adhesiveness:*  Non-adherent  Adherent

*Mode of Disaggregation:*  Trituration  Trypsin  Collagenase  Other \_\_\_\_\_

*Size of cell strainers used:*  70um  50 um  30um  Other \_\_\_\_\_

*Estimated cell size:*  <10um  10-20um  20-70um  70um<

*Number of starting cells:* \_\_\_\_\_

*Volume of cell suspension at start (ml):* \_\_\_\_\_

*Solutions cells suspended:*  PBS  HEPES  Hanks  BSA \_\_\_%  Other \_\_\_\_\_

*Do samples potentially contain infectious agents?*  Yes  No

→If yes, what infectious agents? \_\_\_\_\_

*Do samples contain recombinant or synthetic nucleic acids?*  Yes  No

→If yes, how was your nucleic acid introduced to your samples?  Transfection  Viral vector

→If yes, do those genes encode any recombinant proteins with potential oncogenic properties?  Yes  No

*What is the appropriate Biosafety level of the samples? (See appendix A)*  BL1  BL2  BL2-enhanced

*BUA accession number by IBC, Auburn University or your institution* \_\_\_\_\_

**---Flow lab use only**-----

BUA confirmed	
Sorting BUA level	
AU flow lab sorting experiment #	
Comment	

**Sorting channel information (select all applied)**

Light scatter: FSC SSC

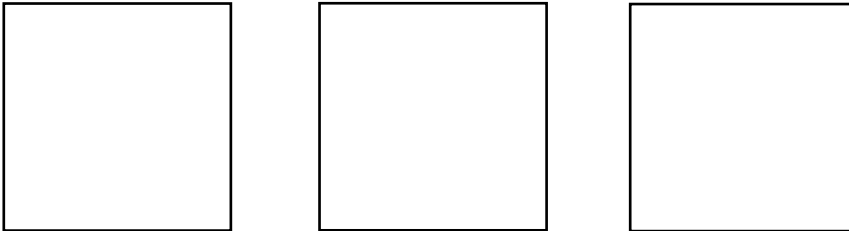
Blue laser (488nm): B525-FITC B690-PC5.5

Yellow laser (561nm): Y585-PE Y610-mCherry Y675-PC7 Y710-PC5.5 Y780

Red laser (638nm): R660-APC R712-PCA700 R780

Violet laser (405nm): V450-PB V525-KrO V610 V660 V780

**Sorting Schema (please show fluorescent patterns, gating parameters, and hierarchy of target cells and subsets)**



**Sample information (for reservoir position and sorting mode, see Appendix B and C)**

Tube ID	Sample	Position*, reservoir	Sorting mode**	Initial vol	Target population
Example	Ex	<input type="checkbox"/> Plate <input checked="" type="checkbox"/> L2: <input checked="" type="checkbox"/> 15ml <input type="checkbox"/> 5ml <input type="checkbox"/> L1: <input type="checkbox"/> 5ml <input type="checkbox"/> R1: <input type="checkbox"/> 5ml <input checked="" type="checkbox"/> R2: <input type="checkbox"/> 15ml <input checked="" type="checkbox"/> 5ml	<input type="checkbox"/> Single cell <input type="checkbox"/> P <input checked="" type="checkbox"/> P1-2 <input type="checkbox"/> Enr <input type="checkbox"/> P <input type="checkbox"/> P1-2 <input type="checkbox"/> Enr <input type="checkbox"/> P <input type="checkbox"/> P1-2 <input type="checkbox"/> Enr <input checked="" type="checkbox"/> P <input type="checkbox"/> P1-2 <input type="checkbox"/> Enr	_____ uL ___ 2 ___ mL _____ mL _____ mL ___ 1 ___ mL	_____ Middle of GFP(+) _____ _____ Top 5% of GFP(+)
1		<input type="checkbox"/> Plate <input type="checkbox"/> L2: <input type="checkbox"/> 15ml <input type="checkbox"/> 5ml <input type="checkbox"/> L1: <input type="checkbox"/> 5ml <input type="checkbox"/> R1: <input type="checkbox"/> 5ml <input type="checkbox"/> R2: <input type="checkbox"/> 15ml <input type="checkbox"/> 5ml	<input type="checkbox"/> Single cell <input type="checkbox"/> P <input type="checkbox"/> P1-2 <input type="checkbox"/> Enr <input type="checkbox"/> P <input type="checkbox"/> P1-2 <input type="checkbox"/> Enr <input type="checkbox"/> P <input type="checkbox"/> P1-2 <input type="checkbox"/> Enr <input type="checkbox"/> P <input type="checkbox"/> P1-2 <input type="checkbox"/> Enr	_____ uL _____ mL _____ mL _____ mL _____ mL	_____ _____ _____ _____ _____
2		<input type="checkbox"/> Plate <input type="checkbox"/> L2: <input type="checkbox"/> 15ml <input type="checkbox"/> 5ml <input type="checkbox"/> L1: <input type="checkbox"/> 5ml <input type="checkbox"/> R1: <input type="checkbox"/> 5ml <input type="checkbox"/> R2: <input type="checkbox"/> 15ml <input type="checkbox"/> 5ml	<input type="checkbox"/> Single cell <input type="checkbox"/> P <input type="checkbox"/> P1-2 <input type="checkbox"/> Enr <input type="checkbox"/> P <input type="checkbox"/> P1-2 <input type="checkbox"/> Enr <input type="checkbox"/> P <input type="checkbox"/> P1-2 <input type="checkbox"/> Enr <input type="checkbox"/> P <input type="checkbox"/> P1-2 <input type="checkbox"/> Enr	_____ uL _____ mL _____ mL _____ mL _____ mL	_____ _____ _____ _____ _____

If needed, add rows

**Checklist on your sorting day**

- Cells suspended in sorting solution, prepared in round bottom 12x75mm tube, 5ml (Ex. Falcon 352058)
- Receiving tubes pre-filled with necessary amount of media or transporting solution (15ml or 5ml tube)
- Transporting container (Need to meet the requirement of your BUA)

## Appendix

### A. Examples of Biosafety level of samples

	<b>BL1 exempt</b>	<b>BL1</b>	<b>BL2</b>
<b>Origin of cells</b>	murine or other non-human, non-primate species	Most, excluding human, non-human primates	human, non-human primates, others
<b>Infectious agent</b>	NO	No	Yes (BL2)
<b>Recombinant nucleic acid</b>	NO	Yes (txfn, TG)	Yes
<b>Viral vector</b>	NO	No	Yes
<b>BUA number</b>	not required	<b>required</b>	<b>required</b>
<b>PPE</b>	Recommended	recommended	<b>required</b>

### B. Differences in reservoir position (5-71)

<b>Position</b>	<b>Reservoir accommodated</b>	<b>Benefit, suitable for</b>
Plate	Multi-well plates	Single-cell cloning
L2 (outer left)	15ml, 5ml	Provides maximum purity. Recommended for precious/rare events to be sorted in.
L1 (inner left)	5ml	Macro-particles ( $\geq 15\mu\text{m}$ ) are recommended to go here
R1 (inner right)	5ml	Macro-particles ( $\geq 15\mu\text{m}$ ) are recommended to go here
R2 (outer right)	15ml, 5ml	Provides maximum purity. Recommended for precious/rare events to be sorted in.

### C. Differences in sorting mode (2-47)

<b>Mode</b>	<b>Definition and major applications</b>	
Single cell	Definition	Having only one event per drop is the most important aspect of the sort.
	Application	single cell cloning (plate/slide sorting)
P: Purity	Definition	The purity of the sort is the most important. Can gate the % of cells wanted.
	Application	Harvesting top XX% positive cells
P1-2: Purity 1-2	Definition	Same with "Purity" but more inclusive than Purity mode.
	Application	Get sorting done a little faster (sacrificing purity slightly).
Enr:Enrich	Definition	Recovery is the most important aspect of the sort.
	Application	Collecting all positive events even it contains negative events in or near the drop.