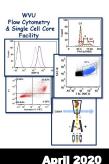
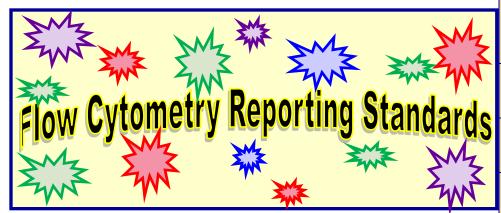
# WVU FLOW CYTOMETRY & SINGLE CELL CORE FACILITY



### Newsletter Volume 6, issue 4

# Minimal Information about a Flow Cytometry Experiment: MIFlowCyt Standards



Over the last few years, the ability to reproduce published data has been demonstrated in many cases to be very difficult in part due to a lack of essential experimental details in the published papers. As grant funding agencies and journals start to ask for more detailed information, it seems like this is a good time to discuss the <u>Minimal Information about a</u> <u>Flow Cytometry Experiment (MIFlowCyt) standards</u> developed by the International Society for the Advancement of Cytometry (ISAC) and endorsed by the Data Interoperability Steering Committee of the Division of Allergy, Immunology, and Transplantation within the National Institute of Allergy and Infectious Diseases (NIAID).

### What is the Purpose of MIFlowCyt Standards?

The purpose of the MIFlowCyt standards is to provide a framework for recording and reporting information about a flow cytometry experiment. It is basically a way to promote consistent annotated information about the biological and technical issues surrounding a flow experiment. MIFlowCyt does this by specifying requirements for data content in a standardized format.

### Who came up with the MIFlowCyt Standards?

An international group of flow cytometry experts got together along with bioinformaticians, computational statisticians, software developers, instrument manufacturers, and clinical and basic research scientists to develop these standards.

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| Facility Location:<br>2160 HSCN  |   |  |  |  |
| <u>Phone:</u><br>304-293-6273  |   |  |  |  |
| <u>email:</u><br>flowcore@hsc.wvu.edu  |   |  |  |  |
| Hours of operation:<br>9:30 am to 5:00 pm, M-F   |   |  |  |  |
| After hours access is availa-<br>ble for experienced users<br>with prior approval from<br>Dr. Kathy Brundage |   |  |  |  |
| Contact Dr. Brundage at:<br>kbrundage@hsc.wvu.edu  |   |  |  |  |

### Why do we need reporting standards?

There are many reasons to have flow cytometry reporting standards. First, with high throughput, high multi-parameter flow cytometry data being generated on a daily basis the computational means for handling and analyzing the data is having a hard time keeping up. Second, the lack of standardization in reporting limits collaborations and independent validation. Third, standardization makes meta-analysis possible. Finally, lack of a standard reporting framework minimizes the value of existing flow cytometry data due to the poor annotation of the data.

### What are the MIFlowCyt Standards?

The MIFlowCyt Standards are clearly laid out in a 2008 Cytometry Part A paper by Lee et al (Lee, JA, et al. MiFlowCyt: The Minimum Information About a Flow Cytometry Experiment. Cytometry PartA Vol.73A pp926-930). The paper gives a nice overview of the standards and why each type is included. In the paper's supplemental data, the authors give an example experiment, descriptive outline of the standards and a required info summary. This paper with supplements will be added to the WVU Flow Cytometry & Single Cell Core Facility website under the Useful Links tab for easy access.

The table below shows the different components that make up the MIFlowCyt Standards. The standards are grouped into 4 catagories; Experimental Overview, Flow Samples, Data Analysis, and Instrument Details. In the paper and supplemental materials the authors use the words shall, shall if relevant, and should to denote whether a standard is an absolute requirement (shall), not generally applicable but maybe in some cases (shall if relevant), or recommended but not required (should).

| Experiment Over-<br>view | Purpose/goal/hypothesis<br>Experimental Variables<br>Conclusions<br>Quality Control                      |  |
|--------------------------|--|--|
| Flow Samples             | Material<br>Source/Organism/location<br>Treatment<br>Reagent/analyte/detector/reporter                   |  |
| Data Analysis            | List-mode data<br>Compensation<br>Gating<br>Descriptive statistics                                       |  |
| Instrument Details       | Instrument Identification<br>Fluidics Configuration<br>Optical Configuration<br>Electronic Configuration |  |

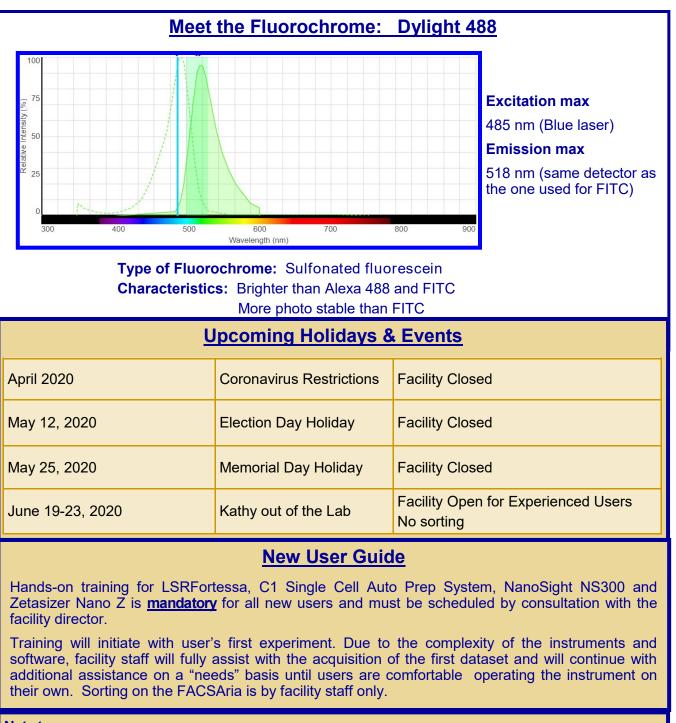
The Experimental Overview component is designed to explain the specific purpose or objective of the experiment. It shall include a summary of the experimental design, results and interpretation of the results. Under the Quality Control subsection, information about replicates, calibrations, control assays etc shall be included.

The Flow Samples components contains the information about the samples, including sample source, sample type, any treatment they under went and how they were prepared. Critical for interpretation, a list of reagents including the protein being detected and fluorescent label antibody used (include clone name, company and fluorescent label) shall be included. Also in this section, a description of process being evaluated i.e. apoptosis, cell cycle shall be included.

The third category is Data Analysis.

For this category, FCS file type as well as the type of compensation performed shall be described. A table of the compensation matrix is also required. A clear description of the gating strategy going all the way back to the all events shall also be included.

The final category is Instrument Details. In this category the instrument manufacturer and model number is sufficient unless the instrument has been uniquely modified. For the WVU FCSCCF, the LSRFortessa (analytical flow cytometry) is a Special Order Research Product (SORP) so the components of this category will have to be included. The FACSAria III (sorter) on the other hand is not a SORP instrument so only the manufacturer and model number is required. To make things easy for users of the core, a standard Instrument Details document for the LSRFortessa and FACSAria III will be put together. Users will be able to find the pdf for each instrument on the flow core website.



### Note to users:

Please remember to acknowledge the support of the HSC Research Office and NIH grants that support the WVU Flow Cytometry & Single Cell Core in all your publications. The grant numbers are listed below:

TME CoBRE grant: P20GM121322

WVCTS grant: GM104942 important if you used the Miltenyi AutoMACS pro (installed 6/29/18) WV InBRE grant: GM103434

Fortessa S10 grant: OD016165

NanoSight NS 300 use Stroke CoBRE grant GM109098 and WVCTS grant GM104942 ZetaSizer NanoZ use Stroke CoBRE grant GM109098 and WVCTS grant GM104942

# **Flow Cytometers in the Facility**

# FACSAria III Cell Sorter

**Operator: Facility Staff** Lasers: 488 nm solid state 561 nm solid state 633 nm solid state 407 nm solid state **Detection Parameters: Forward Scatter** Side Scatter **Simultaneous detection** of 13 fluorochromes **Applications: Cell Sorting (Aseptic)** Single Cell Sorting **Cell Phenotyping Cell Viability Cell Cycle Analysis** 

# <image>

## LSR Fortessa

### **Operator: User of Facility Staff**

Lasers:

405 nm solid state 488 nm solid state 561 nm solid state 628 nm solid state 0etection Parameters: Forward Scatter Side Scatter Simultaneous detection of 17 fluorochromes Applications: Cell phenotyping Cell Viability Cell Cycle analysis Apoptosis Assays



http://flowcore.wvu.edu

# Other Instrumentation Available in the Facility

| AutoMACS Pro Magnetic Bead separator  | gentleMACS Octo Dissociator with Heaters   |  |  |  |  |  |
|---|--|--|--|--|--|--|
| Operator: User  | Operator: User   |  |  |  |  |  |
| Application:  | Application:   |  |  |  |  |  |
| Single extracellular marker cell sorting<br>Depletion/negative cell sorting   | Dissociation of tissues into single cell<br>suspension for culture or flow cytometry<br>assays<br>Homogenizes tissues for downstream molecular<br>biology applications |  |  |  |  |  |
| C1 Single Cell Auto Prep System   | MSD Multi-Array Platform   |  |  |  |  |  |
| Operator: User or Staff   | Operator: User   |  |  |  |  |  |
| Application:  | Applications:  |  |  |  |  |  |
| Uses microfluidics, to separate cells into individual compartments, isolate RNA from the single cells, and generate cDNA for downstream genomic applications. | Detection of cytokines, cell signaling proteins<br>Multiplexed assay design: (1-10 analytes/plate)<br>Detection range: 1 – 10,000 pg/ml                                |  |  |  |  |  |
| Downstream applications:  | Sample volumes: 25 μl or less  |  |  |  |  |  |
| RNA seq   | Assay Time: 4—6 hours depending on analytes<br>being detected  |  |  |  |  |  |
| DNA seq<br>PCR  | being detected   |  |  |  |  |  |
| Format: 96 or 384 chambers per chip   |  |  |  |  |  |  |
| Nanosight NS 300  | Zetasizer Nano Z   |  |  |  |  |  |
| Operator: User or Staff<br>Application:   | Operator: User or Staff  |  |  |  |  |  |
| Determines the size and concentration of particles 10 nm to 1 microns in size   | Application:<br>Measures the zeta potential of particles in a solu-<br>tion using laser Doppler micro-electrophoresis  |  |  |  |  |  |
| Equipped with 4 lasers (405 nm, 488 nm, 532 and 642) to detect fluorescently labeled particles  |  |  |  |  |  |  |
|   | Advanta  |  |  |  |  |  |

| Fee Schedule                    |                          |                            |                         |  |  |
|---------------------------------|--------------------------|----------------------------|-------------------------|--|--|
| Instrument                      | Operator                 | For WVU &<br>NIOSH Users   | For Non-WVU<br>Users    |  |  |
| AutoMACS Pro                    | Facility Staff or User   | \$4.50 /<br>separation     | \$6.85 /<br>separation  |  |  |
| C1 Single Cell Auto Prep System | Facility Staff           | \$210/plate                | \$320/plate             |  |  |
|                                 | User                     | No Cost                    | \$115/plate             |  |  |
| FACSAria III                    | Analysis: Facility Staff | \$52.50/h                  | \$80/h                  |  |  |
|                                 | Analysis: User           | \$34.65/h                  | \$53/h                  |  |  |
|                                 | Sorting                  | \$77.70/h                  | \$120/h                 |  |  |
|                                 | Sorting Setup            | \$19.43/sort               | \$30/sort               |  |  |
| gentleMACS                      | Facility Staff or User   | \$10.50/sample             | \$16/sample             |  |  |
| LSRFortessa                     | Facility Staff           | \$52.50/h                  | \$80/h                  |  |  |
|                                 | User                     | \$34.65/h                  | \$53/h                  |  |  |
| MSD QuickPlex SQ120             | Facility Staff or User   | \$10.50/h                  | \$16/h                  |  |  |
| NanoSight NS300                 | Facility Staff           | \$61.00/h                  | \$93/h                  |  |  |
|                                 | User                     | \$42.50/h                  | \$65/h                  |  |  |
| Zetasizer Nano Z                | Facility Staff           | \$25/sample +<br>\$52.50/h | \$39/sample +<br>\$80/h |  |  |
|                                 | User                     | \$25/sample                | \$39/sample +<br>\$16/h |  |  |



Internal WVU user :

Click here to login or register using your institute login and password.

Not a WVU user?

Login using iLab credentials

If you don't have an account, please register for an iLab account.

The facility uses iLAB scheduling/billing software from Agilent to manage the use of the facility's instrumentation. If you would like to use the instruments housed in the facility please use the URL shown below to register as a WVU User and to login to reserve an instrument.

https://wvu.corefacilities.org/account/login