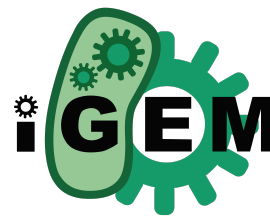


May 08, 2019

## iGEM Calibration Protocol - Flow Cytometry Cell Size

DOI

[dx.doi.org/10.17504/protocols.io.2pegdje](https://dx.doi.org/10.17504/protocols.io.2pegdje)



Jacob Beal<sup>1</sup>, Cheryl Telmer<sup>1</sup>, Richard Tennant<sup>1</sup>, Paul Rutten<sup>1</sup>

<sup>1</sup>iGEM Measurement Committee

iGEM Measurement

Tech. support email: [pauljrutten@gmail.com](mailto:pauljrutten@gmail.com)



**Paul Rutten**

The University of Oxford

OPEN  ACCESS



**DOI:** [dx.doi.org/10.17504/protocols.io.2pegdje](https://dx.doi.org/10.17504/protocols.io.2pegdje)

**Protocol Citation:** Jacob Beal, Cheryl Telmer, Richard Tennant, Paul Rutten 2019. iGEM Calibration Protocol - Flow Cytometry Cell Size. [protocols.io](https://protocols.io) <https://dx.doi.org/10.17504/protocols.io.2pegdje>

**License:** This is an open access protocol distributed under the terms of the [Creative Commons Attribution License](https://creativecommons.org/licenses/by/4.0/), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited

**Protocol status:** In development

**Created:** May 07, 2019

**Last Modified:** August 07, 2019

**Protocol Integer ID:** 22982



## Abstract

Conversion from forward scatter to  $E_{\mu m}$  is not a linear function, so data cannot be converted simply by multiplying with a scaling factor. We thus recommend use of software tools for processing data with size calibration.

This protocol can be combined with bead-based fluorescence calibration.

## Materials

### MATERIALS

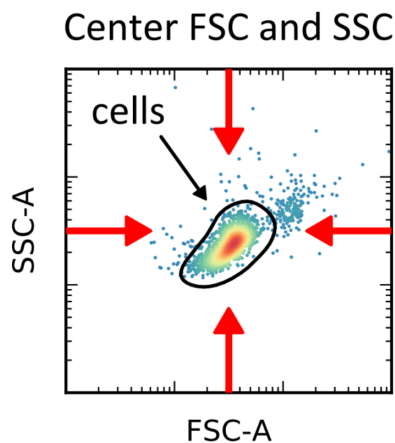
 Size calibration beads **SpheroTech Catalog #PPS-6K**

- Experimental samples: transformed per your desired experimental procedure
- Flow cytometer with a forward scatter channel (typically a 488nm excitation and a 488nm/10nm emission filter). Measurements on this channel will be converted to equivalent  $\mu m$  diameter ( $E_{\mu m}$ ).
- **TASBE Flow Analytics** - Analysis software

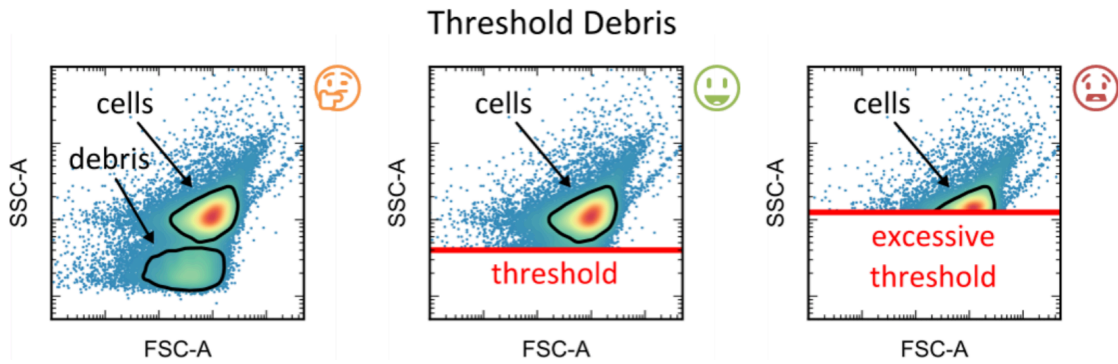
## Before start

Read through this entire protocol carefully before you start your experiment and prepare any materials you may need. See the "Results" section for an example of a completed data analysis spreadsheet.

- 1 Prepare and culture experimental samples according to your desired experimental procedure.
- 2 Prepare experimental samples as needed for running through your flow cytometer.
- 3 Follow SpheroTech directions for preparation for PPS-6K bead sample.
- 4 Measure all samples in flow cytometer
  - Using the bead sample, adjust FSC channel voltage so that the top bead peak is below detector saturation.
  - Using the negative process control sample, adjust forward-scatter and side-scatter voltages to place the strong cell peak as close to the center of the detector range as possible (see example illustrated below).



- Instrument gating should be set to ensure that no cell events are discarded (see example illustrated below).



- Collect at least 10,000 events per sample.
- 5 Compute the  $E_{\mu\text{m}}/\text{a.u.}$  scaling function from bead sample using **TASBE Flow Analytics** to create a TASBE color model.
  - 6 Convert experimental data to  $E_{\mu\text{m}}$  using **TASBE Flow Analytics** analysis functions.