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## iGEM Calibration Protocol - Flow Cytometry Cell Size

Forked from [iGEM Calibration Protocol - Flow Cytometry Cell Size](#)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

1 Works for me

[dx.doi.org/10.17504/protocols.io.5ncg5aw](https://doi.org/10.17504/protocols.io.5ncg5aw)

iGEM Measurement

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### 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

NAME ▾	CATALOG # ▾	VENDOR ▾
<a href="#">Size calibration beads</a>	PPS-6K	SpheroTech

### MATERIALS TEXT

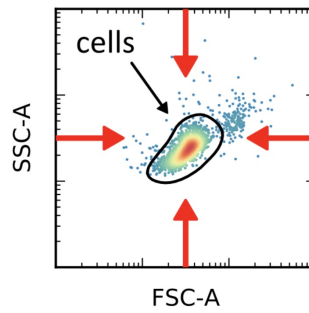
- Experimental samples: transformed per your desired experimental procedure
- Flow cytometer with a forward scatter channel (typically a 488nm excitation and a 488nm/10nm emission filter). Measurements on this channel will be converted to equivalent  $\mu m$  diameter ( $E_{\mu m}$ ).
- [TASBE Flow Analytics](#)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

### BEFORE STARTING

Read through this entire protocol carefully before you start your experiment and prepare any materials you may need.

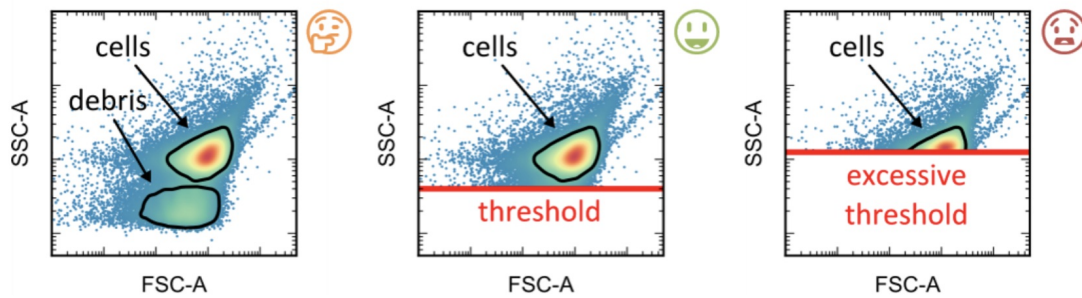
- 1 Prepare and culture experimental samples according to your desired experimental procedure.
- 2 Prepare experimental samples as needed for running through your flow cytometer.
- 3 Follow SpheroTech directions for preparation for PPS-6K bead sample.
- 4 Measure all samples in flow cytometer
  - Using 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).

## Center FSC and SSC



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

## Threshold Debris



- Collect at least 10,000 events per sample.

5 Compute the  $E_{\mu m/a.u.}$  scaling function from bead sample using [TASBE Flow Analytics](#) to create a TASBE color model.

6 Convert experimental data to  $E_{\mu m}$  using [TASBE Flow Analytics](#) analysis functions.



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