VERSION 2

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Calibration Protocol - Plate Reader Abs600 (OD) Calibration with Microsphere Particles V.2

Version 1 is forked from [Calibration Protocol - Particle Standard Curve with Microspheres](#)

ACS Synthetic Biology

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ABSTRACT

You will prepare a dilution series of monodisperse silica microspheres and measure the Abs₆₀₀ in your plate reader.

The size and optical characteristics of these microspheres are similar to cells, and there is a known amount of particles per volume. This measurement will allow you to construct a standard curve of particle concentration which can be used to convert 600 nm absorbance measurements into an estimated equivalent number of cells.

ATTACHMENTS

[iGEM Data Analysis Template - Particle Standard Curve - v1.xlsx](#)

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Protocol status: Working
We use this protocol and it's working

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26497

GUIDELINES

For a full set of calibrations, you should run two protocols: this Abs600 (OD) calibration with microspheres, and the fluorescence calibration curve with fluorescein.

Before beginning these protocols, please ensure that you are familiar with the measurement modes and settings of your instrument. For all of these calibration measurements, you must use the same plates and volumes that you will use in your cell-based assays. You must also use the same settings (e.g., filters or excitation and emission wavelengths) that you will use in your cell-based assays. If you do not use the same plates, volumes, and settings, the calibration will not be valid.

Make sure to record all information about your instrument to document your experiment. If your instrument has variable temperature settings, the instrument temperature should be set to room temperature (approximately 20-25 C) for all measurements.

MATERIALS

MATERIALS

☒ 96 well plate **Contributed by users**

☒ double distilled water (ddH₂O) **Contributed by users**

☒ 300µl Silica beads **Contributed by users**

STEP MATERIALS

☒ 300µl Silica beads **Contributed by users**

☒ ddH₂O **Contributed by users**

300 µL Silica beads are provided in the iGEM Measurement Kit. The 96-well plate should preferably be black with a clear flat bottom.

BEFORE START INSTRUCTIONS

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

Prepare the Microsphere Stock Solution

- 1 Obtain the tube labeled "Silica Beads" from the Measurement Kit and vortex vigorously for 30 seconds.

☒ 300µl Silica beads **Contributed by users**

Note

Microspheres should NOT be stored at 0°C or below, as freezing affects the properties of the microspheres. If you believe your microspheres may have been frozen, please contact the iGEM Measurement Committee for a replacement (measurement@igem.org).

- 2 Immediately pipet 100 μ L microspheres into a 1.5 mL eppendorf tube
- 3 Add 900 μ L of ddH₂O to the microspheres
ddH₂O Contributed by users
- 4 Vortex well. This is your Microsphere Stock Solution

Prepare the serial dilution of microspheres

- 5 Accurate pipetting is essential. Serial dilutions will be performed across columns 1-11. **Column 12 must contain ddH₂O only.**

Initially you will setup the plate with the microsphere stock solution in column 1 and an equal volume of 1x ddH₂O in columns 2 to 12.

You will perform a serial dilution by consecutively transferring 100 μ L from column to column with good mixing.

- 6 Add 100 µl of ddH₂O into wells A2, B2, C2, D2....A12, B12, C12, D12
- 7 Vortex the tube containing the stock solution of microspheres vigorously for 10 seconds
- 8 Immediately add 200 µl of microspheres stock solution into A1
- 9 Transfer 100 µl of microsphere stock solution from A1 into A2
- 10 Mix A2 by pipetting up and down 3x and transfer 100 µl into A3
- 11 Mix A3 by pipetting up and down 3x and transfer 100 µl into A4

- 12** Mix A4 by pipetting up and down 3x and transfer 100 µl into A5
- 13** Mix A5 by pipetting up and down 3x and transfer 100 µl into A6
- 14** Mix A6 by pipetting up and down 3x and transfer 100 µl into A7
- 15** Mix A7 by pipetting up and down 3x and transfer 100 µl into A8
- 16** Mix A8 by pipetting up and down 3x and transfer 100 µl into A9
- 17** Mix A9 by pipetting up and down 3x and transfer 100 µl into A10
- 18** Mix A10 by pipetting up and down 3x and transfer 100 µl into A11
- 19** Mix A11 by pipetting up and down 3x and transfer 100 µl into liquid waste

Note

Take care not to continue serial dilution into column 12

20 Repeat dilution series for rows B, C, D

21 **IMPORTANT!**

Re-Mix (pipette up and down) each row of your plate **immediately before** putting in the plate reader! (This is important because the beads begin to settle to the bottom of the wells within about 10 minutes, which will affect the measurements.)

Note

Take care to mix gently and avoid creating bubbles on the surface of the liquid

Measure OD

22 Measure OD₆₀₀ of all samples in instrument. Disable any path length correction setting on your instrument, if it has one.

If you will be using your data in conjunction with measurements from the [Fluorescence standard curve](#) protocol, make sure you use the same instrument settings for both protocols.

23 Record the data in your notebook. Please note your standard curve should still work well even if a few of your measurements are saturating the instrument

24 Import data into this Excel sheet:

iGEM Data Analysis Template - Particle Standard Curve - v1.xlsx

Congratulations!

25 You have now completed this calibration protocol