

Protocol

Materials

- Hemocytometer (Neubauer, Improved Neubauer, Bürker, or Bürker-Türk)
- Trypan Blue (TB) or Erythrosin B (EB) stain for viability
- Microscope compatible with SnapCyte™ Adapter
- SnapCyte™ Adapter and App
- Cell culture medium
- Pipettes and tips
- Cell suspension

Procedure

Sample Preparation:

1. Harvest your cell culture and prepare a cell suspension in an appropriate medium.
2. Stain the cells with Trypan Blue (TB) or Erythrosin B (EB) to differentiate live and dead cells. Typically, use a 1:1 dilution (e.g., 10 μ L of cell suspension mixed with 10 μ L of stain).
3. Gently pipette 10-20 μ L of the stained cell suspension into the hemocytometer chamber. Ensure that the suspension fills the chamber evenly without overflowing.

Ensure your cell cultures are free from debris and clumps for clear imaging.



Capture and Analyze Images:

1. Set up the SnapCyte™ Adapter on your microscope and focus on the hemocytometer grid at 10X magnification. Ensure the grid lines and cells are clearly visible.
2. Open the SnapCyte™ App and create a new experiment or a new snapshot. Select the "Hemocytometer" option to enable automatic cell concentration calculations.
3. Capture images of the hemocytometer chambers using the SnapCyte™ App. For best results, capture at least 4 images of the central grid area.
4. The SnapCyte™ App will automatically analyze the images to count cells and assess viability.

When using a hemocytometer, always capture images from the same specific regions (e.g., the four corner squares). This consistency helps reduce variability between counts.



avoid areas near the edges of the hemocytometer grid where cells may appear distorted or unevenly distributed



Quantification:

Review Results:

1. Use the Image thumbnails in your experiment to review individual images and confirm cell counts. Adjust the Regions of Interest (ROIs) if necessary.

Export Data:

1. Utilize the Graph Tab in the SnapCyte™ App to visualize your data. You can choose to display data as Mean, Mean + SD, etc.
2. Export the data for further analysis or download the images for documentation and presentations.

Always visually verify that the software is accurately identifying cells and not including debris or artifacts

