Evaluation of a fluorescent approach to implement monoclonality assurance using calcein-AM viability dye

Carola Mancini, PhD, Sarmad al-Bassam, PhD Molecular Devices, LLC, Munich, Germany

## Summary

Assessing monoclonality is key to the establishment of a cell line, and evidence of monoclonality is required by regulatory agencies to get a biopharmaceutical drug to the marketplace. The traditional and most accepted approach involves the visualization of microplate wells using transmitted white light (brightfield) on day 0 to confirm the presence of a single cell. However, the definitive identification of a single cell on the initial day of cell seeding is not without challenge, as cellular debris and well artifacts can be easily mistaken for cells. Here, we demonstrate an optimized workflow using the fluorescence reagent, calcein-AM (CAM), in conjunction with a fluorescence-capable CloneSelect™ Imager (CSI-FL) that shows similar viability to label-free conditions while simultaneously providing high assurance of clonality. In this workflow, we determine a concentration of CAM that is ideal for detection of single cells on a CloneSelect™ Imager-FL while minimizing cytotoxic effects on clonal outgrowth. We describe guidelines for establishing an optimal concentration of viability dye for different cell types.

# **Results**

## Determining the optimal CAM concentration for single cell detection

A critical parameter in developing a reliable labeling protocol is ensuring cells are adequately labeled for detection. Three concentrations of CAM (0.5  $\mu$ M, 1  $\mu$ M, 5  $\mu$ M) were compared under identical staining conditions. We observed that the majority of cells are detectable at 5  $\mu$ M and 1  $\mu$ M CAM, but not at 0.5  $\mu$ M. Note that the concentrations depend on specific cell type and staining protocol, but a similar dilution series can be performed to determine optimal staining coverage (Fig. 2A–F).



# Introduction

The development of cell lines that express a specific protein of interest is critical to the generation of biologics, and regulatory agencies require evidence of monoclonality in order to get a biologic to the marketplace. The traditional approach involves visualization of wells using transmitted white light on day 0 to confirm the presence of a single cell (Fig. 1A) However, this approach is not without challenge, as cellular debris and well artifacts can be easily mistaken for cells. Consequently, cell line developers typically evaluate cells at the colony state and trace back the origins of the colony to confirm monoclonality. An alternative method involves fluorescence labeling of the cell population prior to seeding single cells (Fig 1B). Here we use the viability dye calcein-AM (CAM), a molecule that fluoresces green only after translocating the membrane of living cells, to more easily automate single cell detection while simultaneously establishing clonality more conclusively. We outline the differences between a transmitted white light (WL) workflow and one involving both WL and fluorescence (FL).

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#### A Label-free Monoclonality Workflow

### FluorescenceMonoclonality Workflow







Analyze

Day 0 WL + FL

Review plate scan and assess overall outgrowth (WL + FL)

Report

No. of the second

WL

FL

CloneSelect Imager FL



**Figure 2.** Finding a suitable concentration of CAM to optimize detection at the single cell level. High-density wells containing cells were stained with 5  $\mu$ M (A), 1  $\mu$ M (B) and 0.5  $\mu$ M (C) CAM. (D–F) Fluorescence scan of plate using CSI-FL. Cells were isolated by limiting dilution. (G) Number of cell per well count for the 5, 1, 0.5  $\mu$ M CAM cell staining and comparison of theoretical values from limiting dilution at 0.5 cells per well. Comparison of single cell counts (dashed line).

### **Evaluating the effect of CAM on colony outgrowth**

Fluorescent probes that stain living cells are notorious for having deleterious effects on cell viability. As such, we tested the effect of CAM at different concentrations (0.5  $\mu$ M, 1  $\mu$ M, and 5  $\mu$ M) on outgrowth after limiting dilution. A decrease in the percentage of outgrowth when using 5  $\mu$ M CAM was observed. In comparison, the 0.5  $\mu$ M, 1  $\mu$ M, and label-free control show no discernible differences in the number of raw single-colony outgrowth (Fig. 3A–E).

Scan plate WL at Day 0 and subsequent days



Analyze



Day 14 WL Review plate scan and assess overall outgrowth (WL)



 Image: Day 14 WL
 Day 3
 Day 2
 D

Confirm monoclonality and export well data for regulatory submission



Monoclonality confirmed on Day 0

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Figure 1. Comparison of the transmitted white light (A) and fluorescence workflow (B) to assess monoclonality.



5 µM calcein AM



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**Figure 3.** Measuring the effect of different CAM concentrations on viability. (A) Day 10 clonal outgrowth of representative plates from labelfree, 0.5  $\mu$ M, 1  $\mu$ M, and 5  $\mu$ M CAM stained cells before limiting dilution. (B) Quantification of raw outgrowth of single colony-containing wells. (C) Representative images of single colony-containing wells at day 6. (D) Quantification of colony size (surface area) shows that colonies derived from cells stained with 5  $\mu$ M CAM are smaller in size compared to label-free cells. (E) Quantification of single colony wells derived from single cells shows consistent data to raw outgrowth percentages outlined in (B).

## Conclusion

1 µM calcein AM



/ell containing a single colo

- An optimized fluorescent workflow to implement monoclonality assurance was demonstrated. Calcein-AM fluorescence viability dye was used and stained cells were imaged with a fluorescence-capable CloneSelect Imager system, showing similar viability compared to label-free conditions.
- 1µM of calcein-AM was ideal for single cell detection on the CloneSelect Imager while not inducing any cytotoxicity. A
  similar assay optimization step would be needed for other dyes, cell types, imaging systems, and other experimental
  conditions.
- The fluorescence approach for confirming monoclonality is significantly beneficial in reducing the time required to manually identify single cells. The analysis can be immediately performed on day 0, providing information on clonality earlier in the workflow.



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