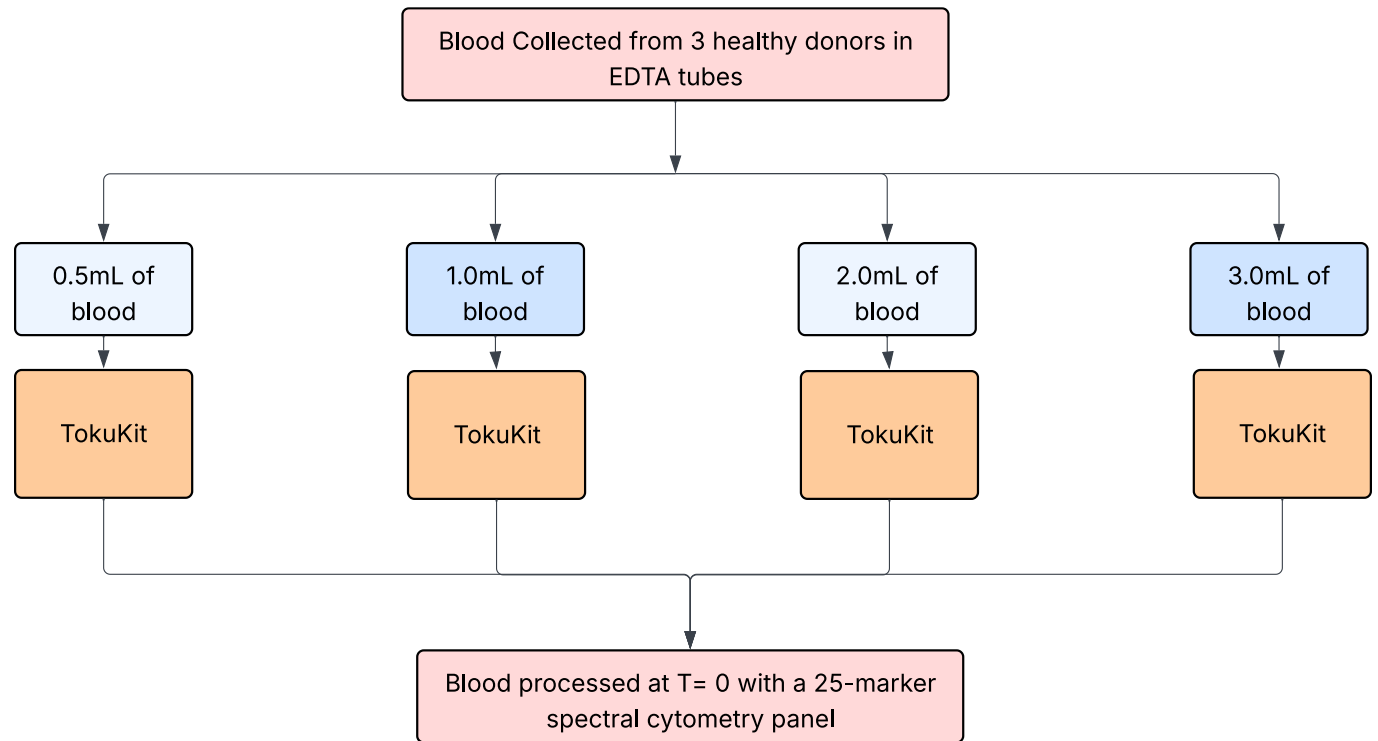


# **Does Blood Volume Affect Immune Cell Preservation with TokuKit?**

This study shows that regardless of blood inputs from from 0.5 mL to 3 mL, TokuKit preserved immune cell populations within  $\pm 2\%$  of the standard 2 mL blood volume.

# Experimental Design



# Experimental Design

1

## Sample Collection

Whole blood was collected from **3 healthy donors** into EDTA tubes and processed with **TokuKit** at **4 volumes: 0.5, 1, 2, and 3 mL**.

2

## Timeline

Samples were processed at  $T = 0$  and analyzed immediately after.

3

## Red Blood Cell Lysis

Since TokuKit contains its own **lysis buffer**, no additional ACK or BD lysis step was required.

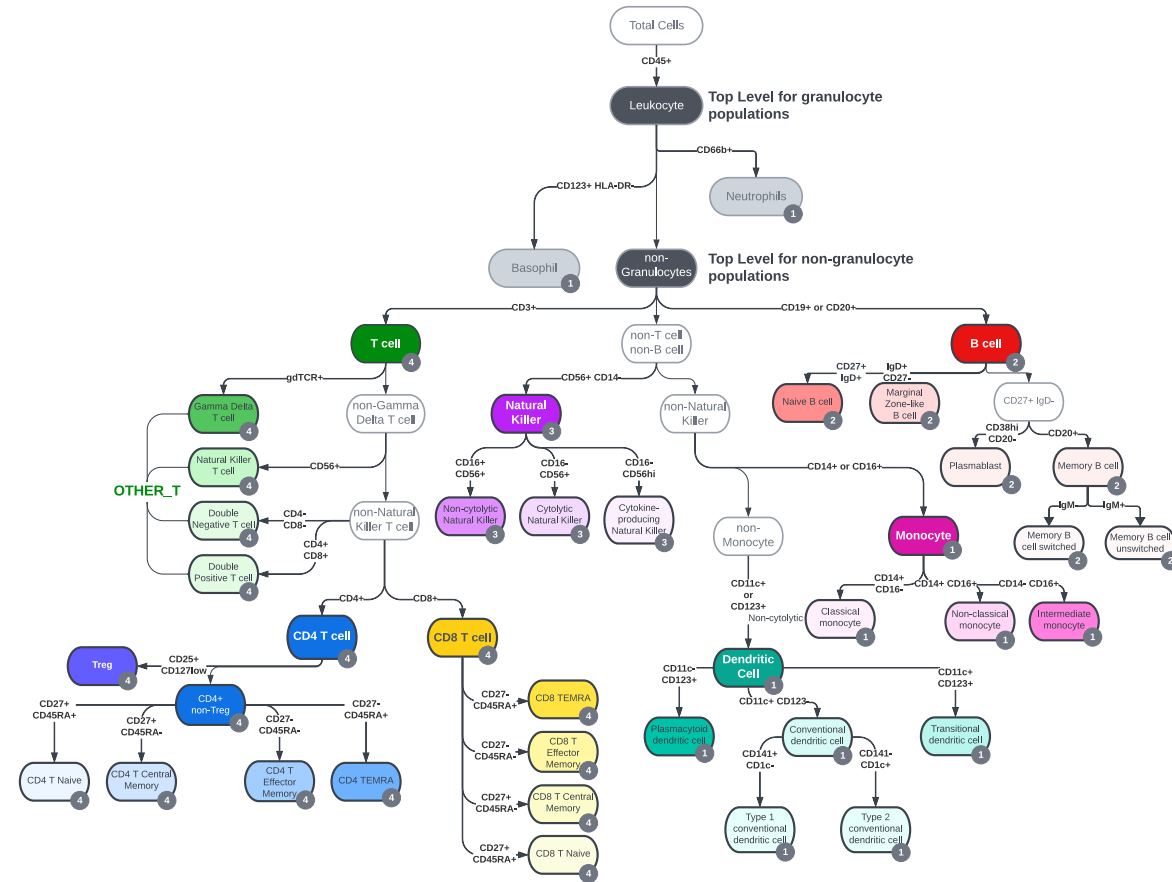
4

## Instrument

Cytometry analysis was performed on a **Cytek® Aurora**.

# Panel Design

A 25-marker spectral flow cytometry panel was applied to assess the immune cell populations visualized in the gating strategy displayed.



# Lower blood volumes did not impact cell population frequencies

## 95% of immune populations

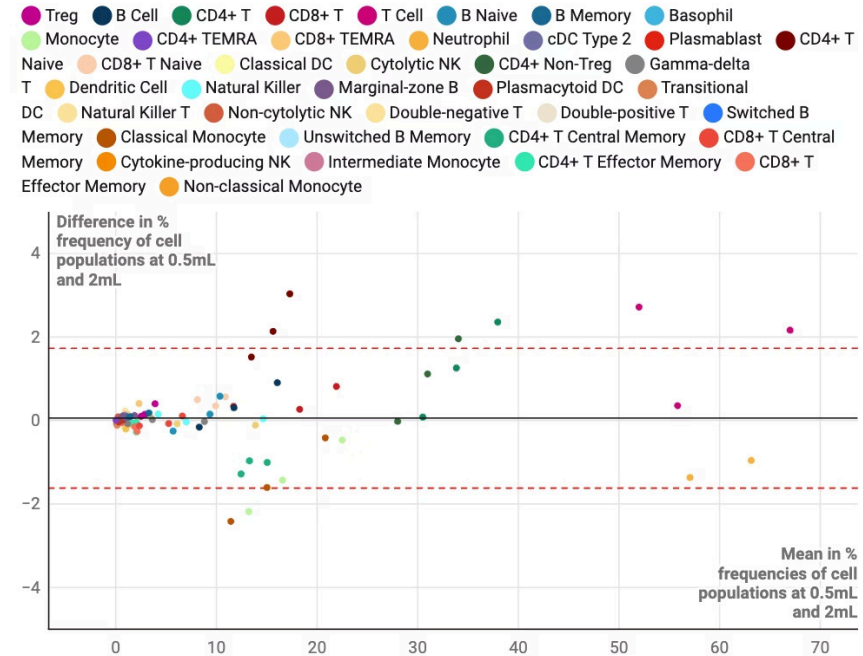
had stable frequencies across volumes ( $\geq 2$  of 3 donors within  $\pm 2\%$ )

## <0.1 mean difference

virtually no systematic shift between volumes (0.5 mL or 2 mL)

## 4 conditions

4 blood volume conditions were tested

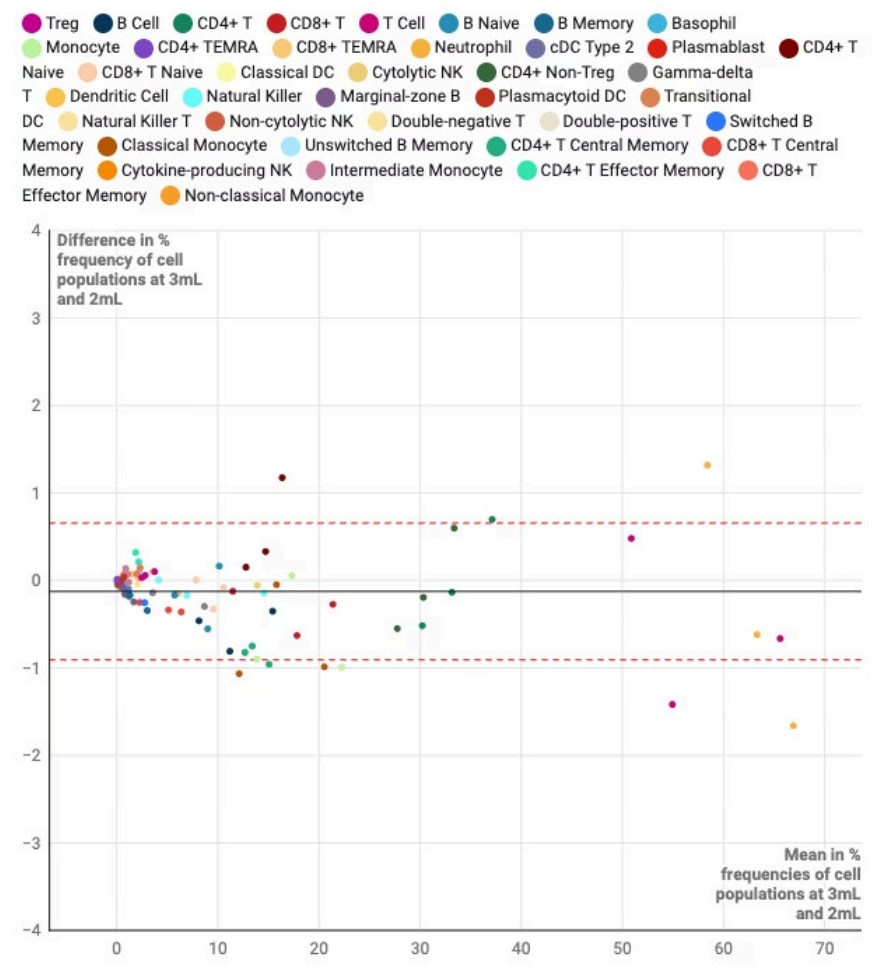


# Higher blood volumes did not impact cell population frequencies

**95% of immune populations**  
had stable frequencies across volumes ( $\geq 2$  of 3 donors within  $\pm 1\%$ )

**<0.2 mean difference**  
virtually no systematic shift between volumes (0.5 mL or 2 mL)

**4 conditions**  
4 blood volumes conditions were tested



# Why Sample Preservation Matters



## **Degradation distorts patient immune profiles**

When preservation methods degrade or introduce artifacts, immune profiles shift, creating results that reflect preservation instability rather than a patient's true biology.



## **Failed samples waste time and budget**

Degraded samples often fail QC, requiring costly redraws, repeat shipments, and delays in data delivery.



## **Inconsistencies undermine PD insights**

Differences in how long samples sit before processing introduce noise, making it harder to detect real pharmacodynamic signals, mechanism of action, or biomarkers of response.

# Want to learn more?

Ask us about the results of our TokuKit blood volume study, and what studies with TokuKit are coming next:

[info@teiko.bio](mailto:info@teiko.bio).

