

# Simultaneous quantification of activated immune cells and PD-L1 expressing circulating tumor cells (CTCs) in peripheral blood of cancer patients receiving checkpoint inhibitor therapy

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**Epic Sciences** 

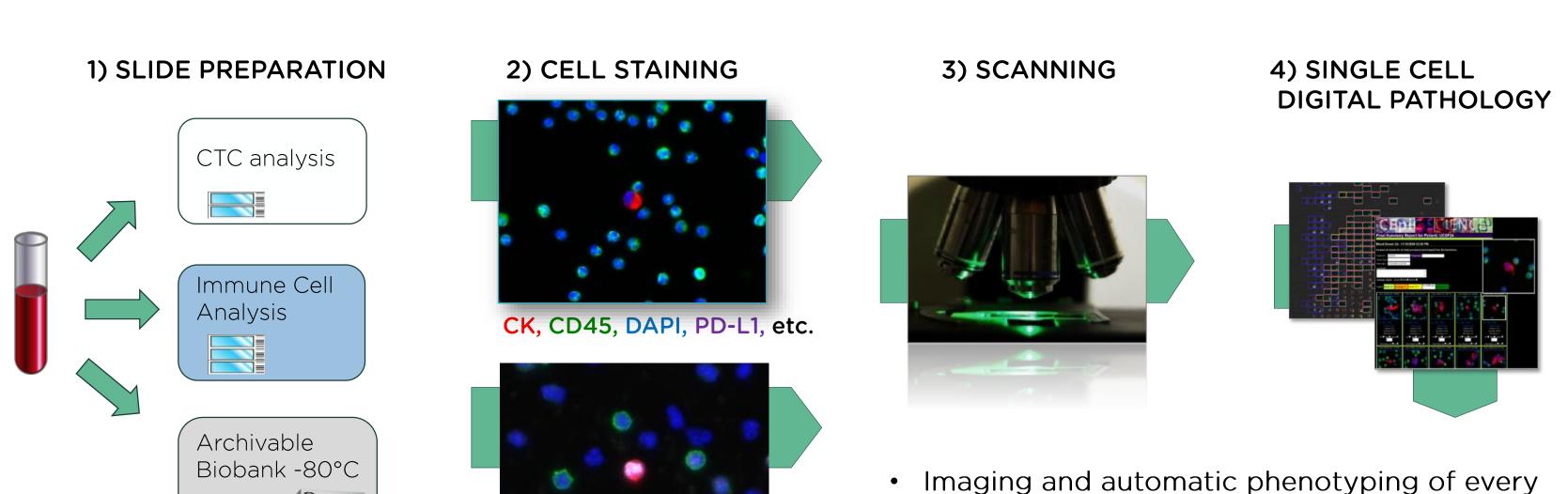
### Background

- Expression of PD-L1 on tumor and tumor infiltrating lymphocytes has been associated with improved response to PD-1 and PD-L1 immune checkpoint inhibitors (ICI), however clinical utility is limited and tumor biopsies are not always available.
- Analysis of the peripheral immune cell subpopulations could aid in the prediction of response to immunotherapy, however current technology robust enough for a clinical setting utilizing flow cytometry suffers from low sensitivity and a lack of standardization.
- Epic's Functional Cell Profiling (FCP) platform uses high-throughput microscopy to analyze all nucleated cells within a tube of blood in a robust and reproducible manner.
- Here we sought to characterize activated T-Cell subpopulations using FCP in patients receiving ICI therapy as a proof of concept with the overarching aim of developing better diagnostic tools for immunotherapy response prediction and pharmacodynamics studies.

### Methods

### The Epic Sciences Functional Cell Profiling platform

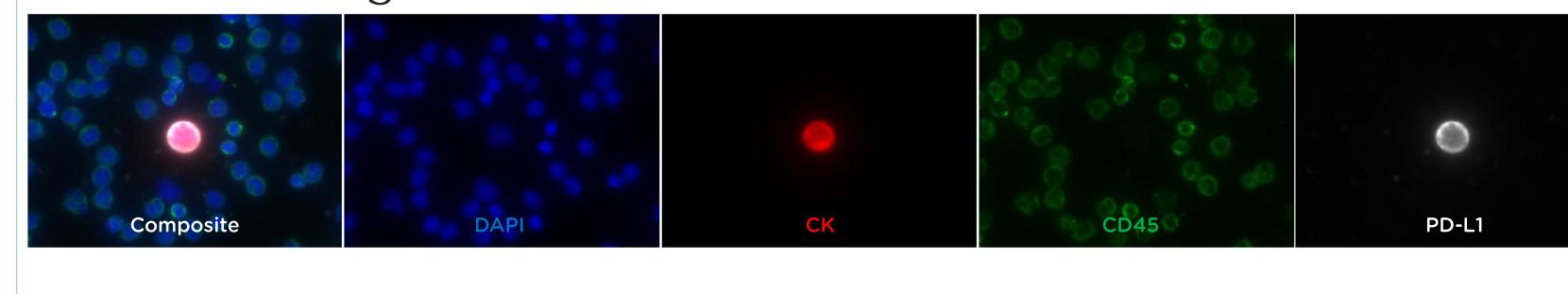
CD8, CD4, DAPI, Ki-67, etc.



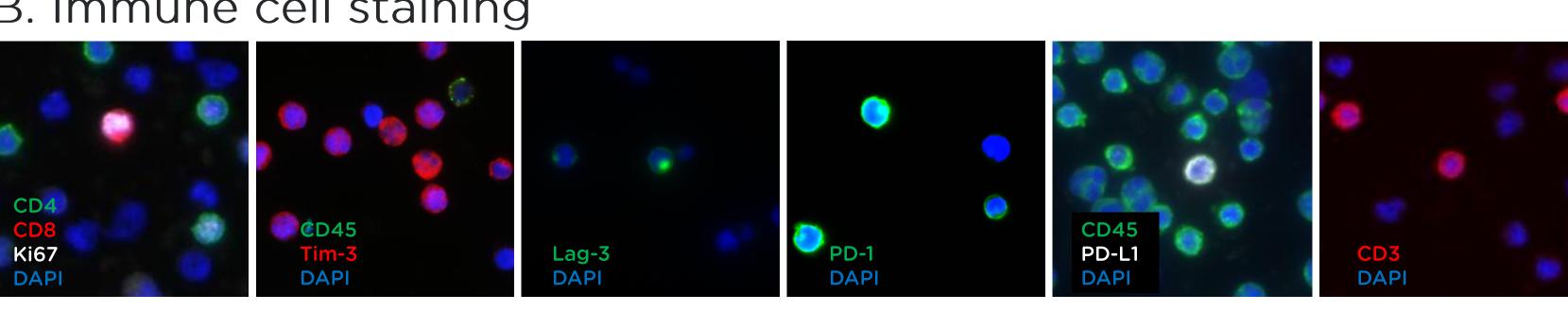
- cell within a tube of blood
- Accurate identification of rare cell populations in a background of  $10^6$  -  $10^7$  other cells

# CTC and Immune Cell Imaging

#### A. CTC staining

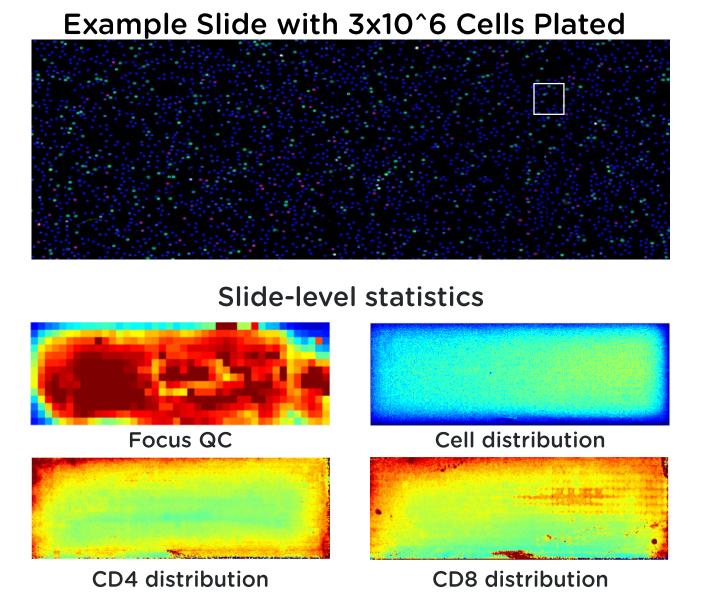


#### B. Immune cell staining

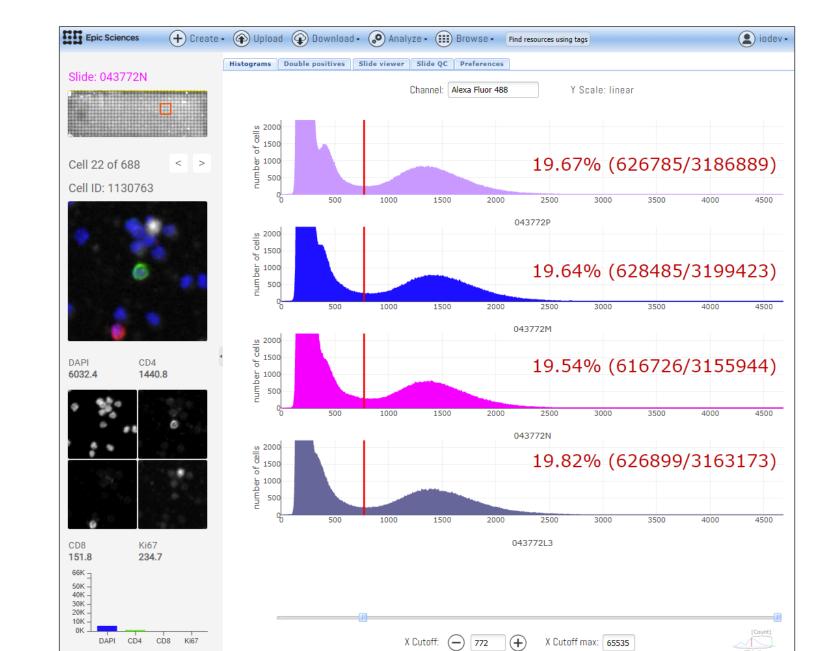


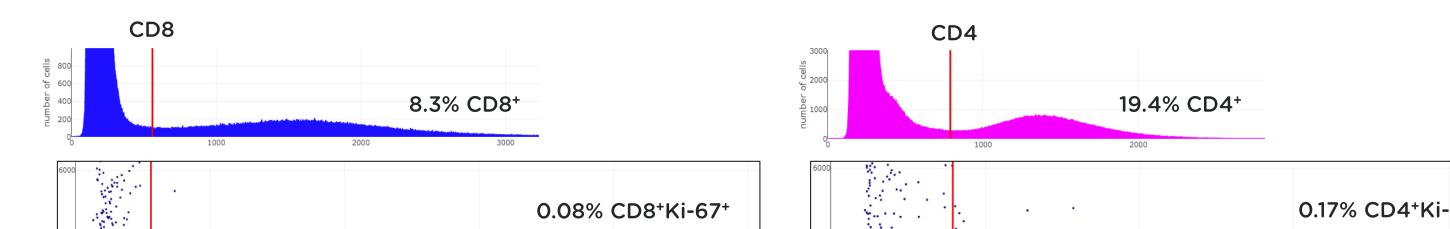
### Cloud-based Epic Discovery Platform

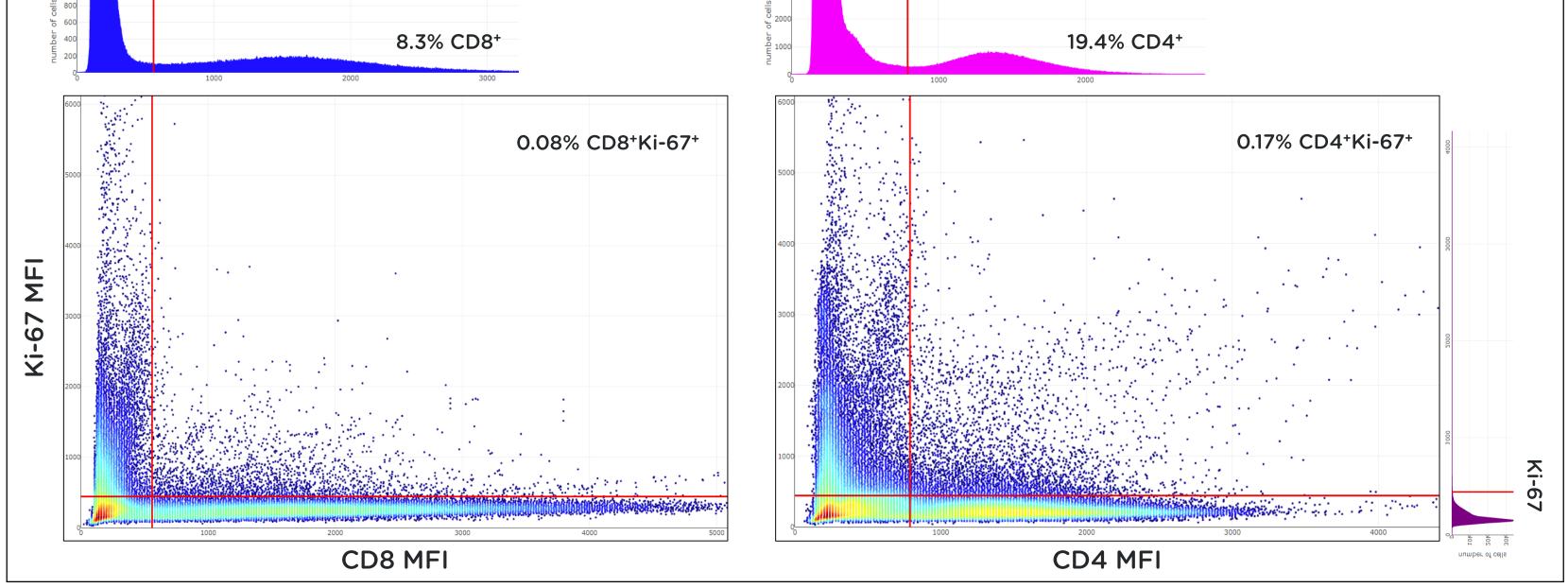
- Epic Discovery Platform provides a new way to quantify leukocyte subpopulations by analyzing all cells and supports image storage, analysis, and provenance management.
- · Cell Explorer links image-based features, cell location, and any relevant metadata for every cell on the slide.



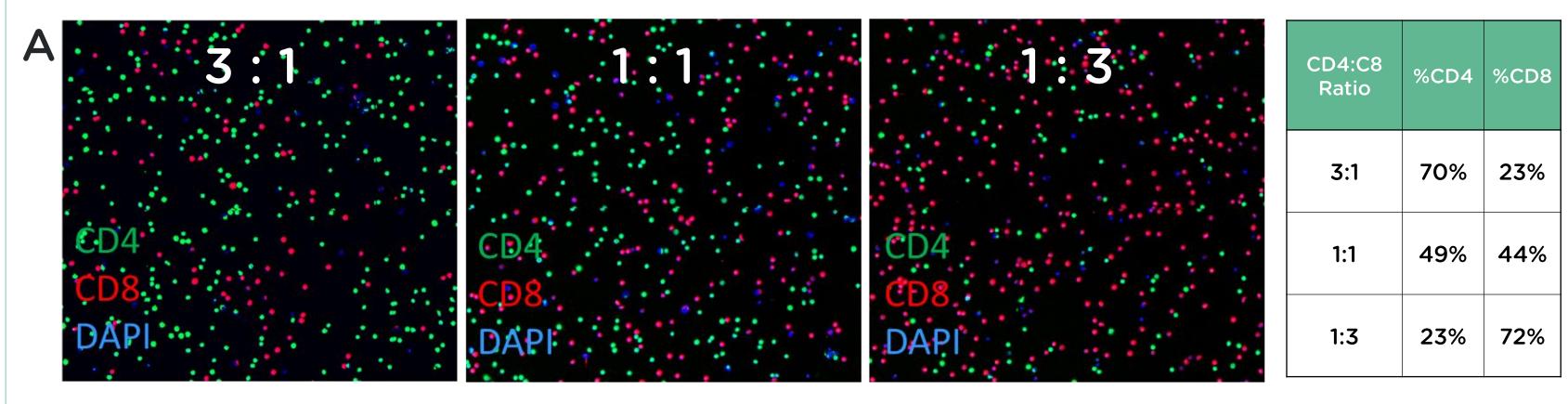
Positivity quantization based on histogram analysis

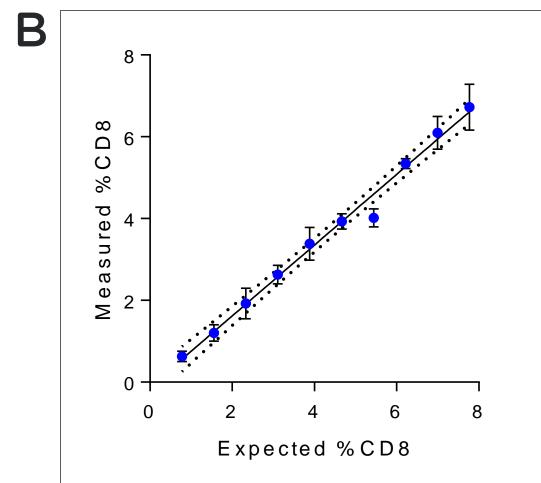






# Large and small scale changes can be detected with the T-Cell Activation assay



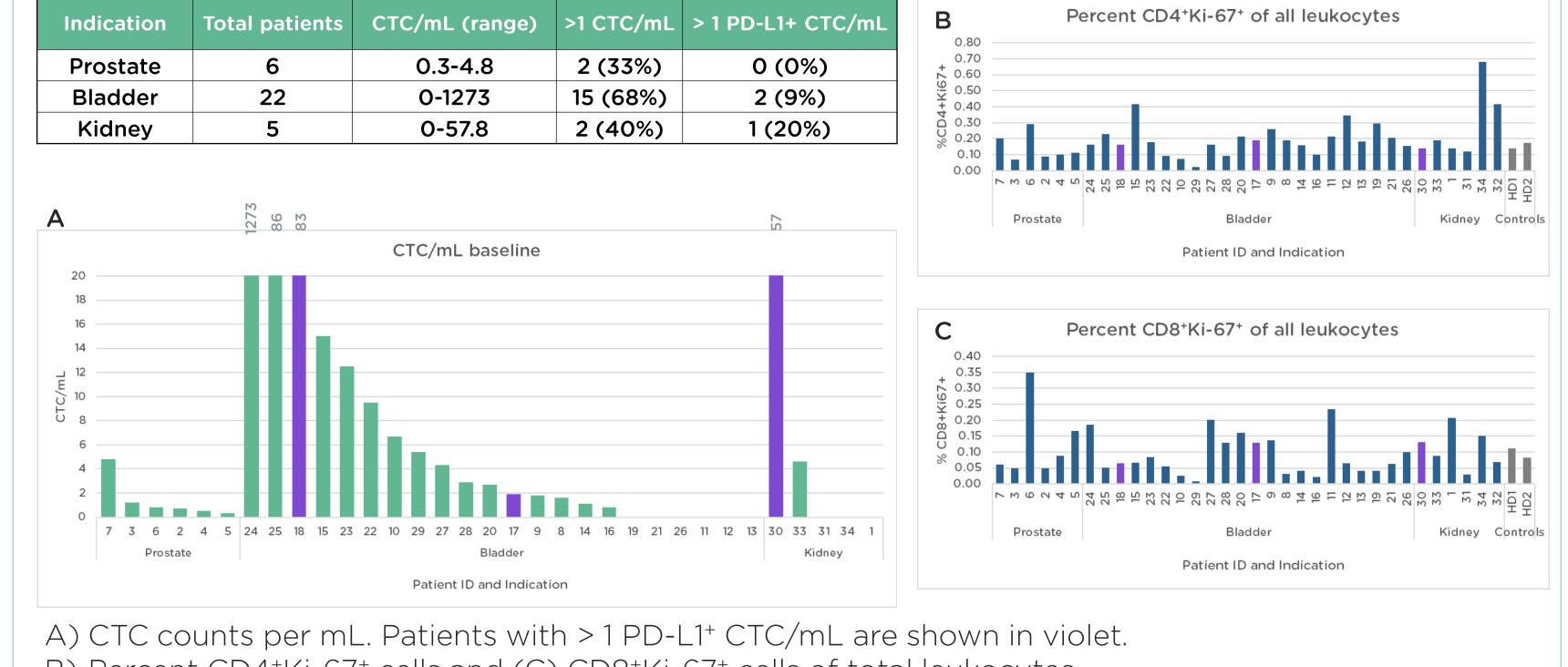


A. Immunomagnetically purified CD4 and CD8 cells plated in ratios of approximately 3:1, 1:1, and 1:3 (left to right). Analyzed ~1 million cells per slide.

B. Immunomagnetically purified CD8 cells spiked into healthy donor blood at target ratios of approximately 1-8% demonstrate linearity of the assay. Mean and SEM of triplicate values are shown.  $R^2 = 0.989$ , slope =

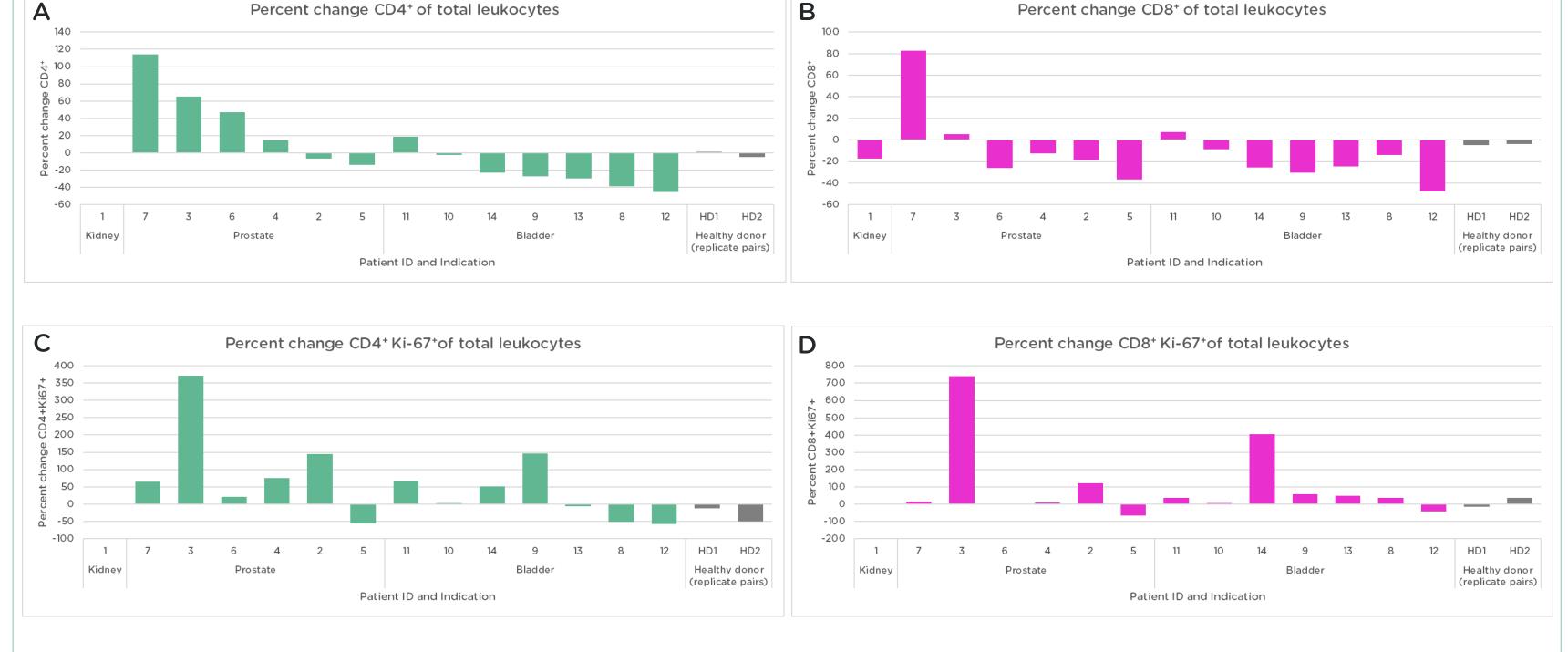
## A single patient sample can be used for characterization of both CTCs and immune cell subpopulations





B) Percent CD4<sup>+</sup>Ki-67<sup>+</sup> cells and (C) CD8<sup>+</sup>Ki-67<sup>+</sup> cells of total leukocytes

# Longitudinal changes were observed upon treatment with checkpoint inhibitor Tx



A and B) Percent change of CD4 and CD8 subpopulations from baseline to on-therapy. C and D) Percent change of CD4<sup>+</sup>Ki-67<sup>+</sup> and CD8<sup>+</sup>Ki-67<sup>+</sup> subpopulations from baseline to on-therapy.

#### Conclusions

- Epic's Functional Cell Profiling (FCP) platform can simultaneously measure immune related biomarkers in CTCs and leukocytes at single cell resolution from a single blood sample.
- Archived samples can be banked for months to years after procurement enabling retrospective testing with multiple assays.
- The platform has extreme sensitivity and reproducibility over conventional methodologies - Cell sub-populations that exist in a background of 10<sup>5</sup>-10<sup>7</sup> other cells are readily identified.
- Changes in immune cell sub-populations in patients receiving ICI therapy were observed across multiple cancer types including bladder, kidney, and prostate.
- Efforts to identify morphological sub-populations within each canonical leukocyte category and comparison of leukocyte and CTC biomarker panels to patient outcomes are ongoing in multiple trials.