

Simultaneous Characterization and Quantification of Immune Cell Subpopulations and PD-L1 Expressing CTCs in Peripheral Blood of Cancer Patients

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Background

- Expression of PD-L1 on tumor, and immune markers in tumor tissue, are associated with improved response to PD-1 and PD-L1 checkpoint inhibitors. However, each alone has limited predictive utility.
- Multimodal characterization of both the tumor and host immune system is an unmet medical need for the improved prediction of response to immunotherapy.
- Metastatic lesions are likely to be under-sampled and require a liquid biopsy, given tumor heterogeneity and evolution and temporal changes in the host immune system.
- We sought to examine expression of PD-L1 on CTCs as well as characterize rare immune cell populations with a non-invasive liquid biopsy. Examining dynamic biomarker changes in longitudinal samples could enable the development of novel diagnostic tools for response prediction and pharmacodynamics studies related to immunotherapy.

Methods

Blood samples were drawn from 3 healthy donors, 5 non-cancerous lung disease patients, and 4 metastatic cancer patients (prior to and on-therapy with checkpoint inhibitor), and sent to Epic Sciences for processing with Epic Sciences' CTC and immune cell assays.

1) SLIDE PREPARATION



2) CELL STAINING



CK, CD45, DAPI, PD-L1, etc.



4) SINGLE CELL DIGITAL PATHOLOGY







Schematic of Epic's CTC platform: CTC enumeration, leukocyte classification, morphology and protein analyses

- 1) Nucleated cells from a 10 mL blood draw are plated onto 10-12 slides (approximately 3 million cells/slide) 2) Slides are stained with 4',6-diamidino-2-phenylindole (DAPI) and a combination of one or more additional markers, including cytokeratin (CK),
- CD45, PD-L1, CD3, CD4, CD8, CD14, Ki-67, PD-1, Lag-3, and Tim-3 3) Slides are scanned using a rapid fluorescent scanning method, which images each nucleated cell 4) All cells are analyzed with a multi-parametric digital pathology algorithm for morphological features and protein expression



CTC Phenotypic Analysis:

- PD-L1 protein • Calreticulin
- Tumor Cell
- Heterogeneity

Immune Cell

- Characterization: • Activated T Cells
- Regulatory T Cells
- Exhausted T Cells MDSCs
- NK cells

Schematic of Epic Single Cell Tools for Immune Monitoring 1) A single blood draw is used to make slides for CTC and immune cell analysis. Extra slides are archived for staining at a later date 2) Slides can be used for CTC and immune cell characterization as well as downstream genomic analysis

3) SCANNING



s	Cytoplasmic Features	Cell Features
	Cytoplasmic Area	AR Expression
	Cyto Convex Area	CK Expression
	Cyto Major Axis	N/C Ratio
5	Cyto Minor Axis	
,	Cyto Circularity	
	Cyto Solidity	
	Cyto Entropy	
	Cyto Speckling	
	Cyto Presence	

CTC Genomic

Analysis:

 Genomic Instability Mutational Burden

PD-L1(+) Peripheral Circulating Cells in Lung Cancer



¹Boffa D. J. Cellular Expression of PD-L1 in the Peripheral Blood of Lung Cancer Patients is Associated with Worse Survival. Cancer Epidemiology, Biomarkers & Prevention. 2017; 26(7):1139-

Tools to Characterize Rare Immune Cell Populations



Representative images of immune cell staining in control cell lines and healthy donor or patient leukocytes. Cells were plated and stained with DAPI and one or more immune cell markers including CD4, CD8, Ki-67, CD3, PD-L1, CD45, PD-1, Tim-3, Lag-3, and CD14.



Cell Ratios (CD4:CD8) 1:1

CD4 CD8 DAPI	04 08. ∆PI	CD4 CD8 DAPI	
	3:1	1:1	
% of CD4 Cells	70%	49%	
% of CD8 Cells	23%	44%	

3:1

Detecting large scale immune cell changes with the CD4, CD8 Assay. Contrived samples were prepared with immunomagnetically purified CD4 and CD8 cells plated in ratios of approximately 3:1, 1:1, and 1:3. Table shows measured cell counts as a percentage of total cells (approximately 1 million total cells per slide).

Representative scatter plots demonstrating CD4, CD8, and Ki-67 threshold determination in a healthy donor sample. Healthy donor leukocytes were plated and stained with CD4, CD8, Ki-67, and DAPI. Mean fluorescent intensity (MFI) was plotted for approximately . million cells to generate histograms and determine positivity thresholds for each marker, and to quantify CD4⁺ and CD8 ⁺ cell counts. Thresholds were applied to scatter plots for guantification of double positive values for CD4+Ki67+ and CD8+Ki-67+.





Detecting small scale immune cell changes: auantitation and linearity of the CD8 assay. CD8+ cells were spiked into healthy donor leukocytes at target ratios of approximately 1 - 9%. After staining and quantification of CD8⁺ cells, percentages of detected spiked CD8⁺ cells for three technical replicates (Percent CD8⁺ Measured) were plotted against target percentages (Percent CD8+ Target). Bars show mean and standard error of the mean.





healthy donor replicates were analyzed as pairs (D).

determined for patients and healthy donor samples (C). Percent change between baseline and on-therapy was calculated for patient samples and

Conclusions

• Multimodal characterization of both the tumor and host immune system is an unmet medical need for the improved prediction of response to immunotherapy.

• Detection of PD-L1 in circulating cells from NSCLC liquid biopsy samples was associated with worse overall survival, highlighting the potential utility of minimally invasive liquid biopsies for tumor profiling.

• Epic Sciences has developed tools to simultaneously detect, characterize, and quantify CTCs and rare immune cell subpopulations from a single blood draw.

Examining dynamic biomarker changes with minimally invasive liquid biopsies in longitudinal samples could enable the development of novel diagnostic tools for response prediction and pharmacodynamics studies related to immunotherapy.