



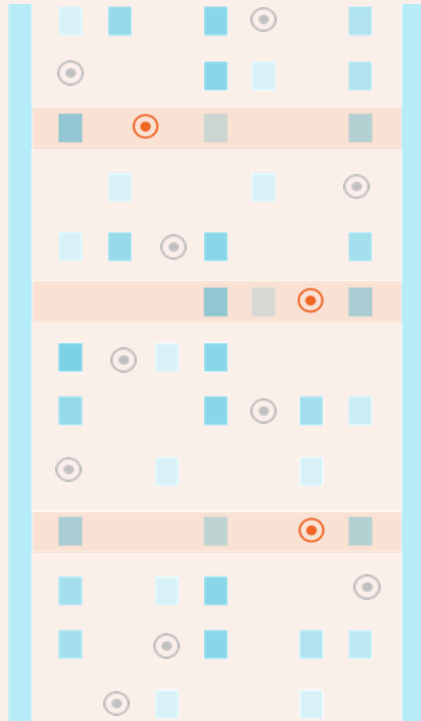
Precision Profiling
...predictive single-cell response.

November 9th, 2017

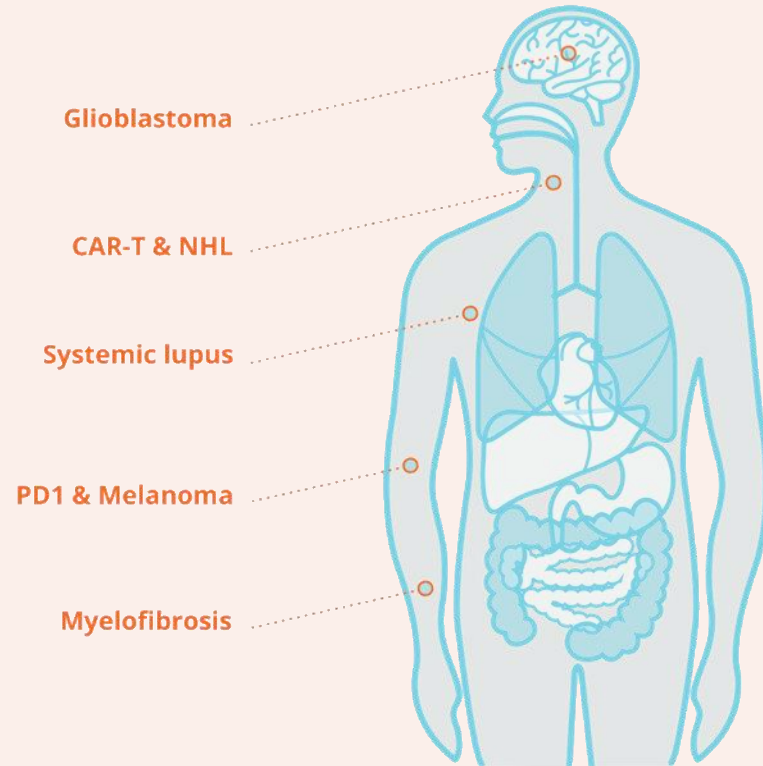
This presentation is not available for CME/CE credit

Target the right therapies, as early as possible to highest urgency cancer patients

Predictive Single Cell Response

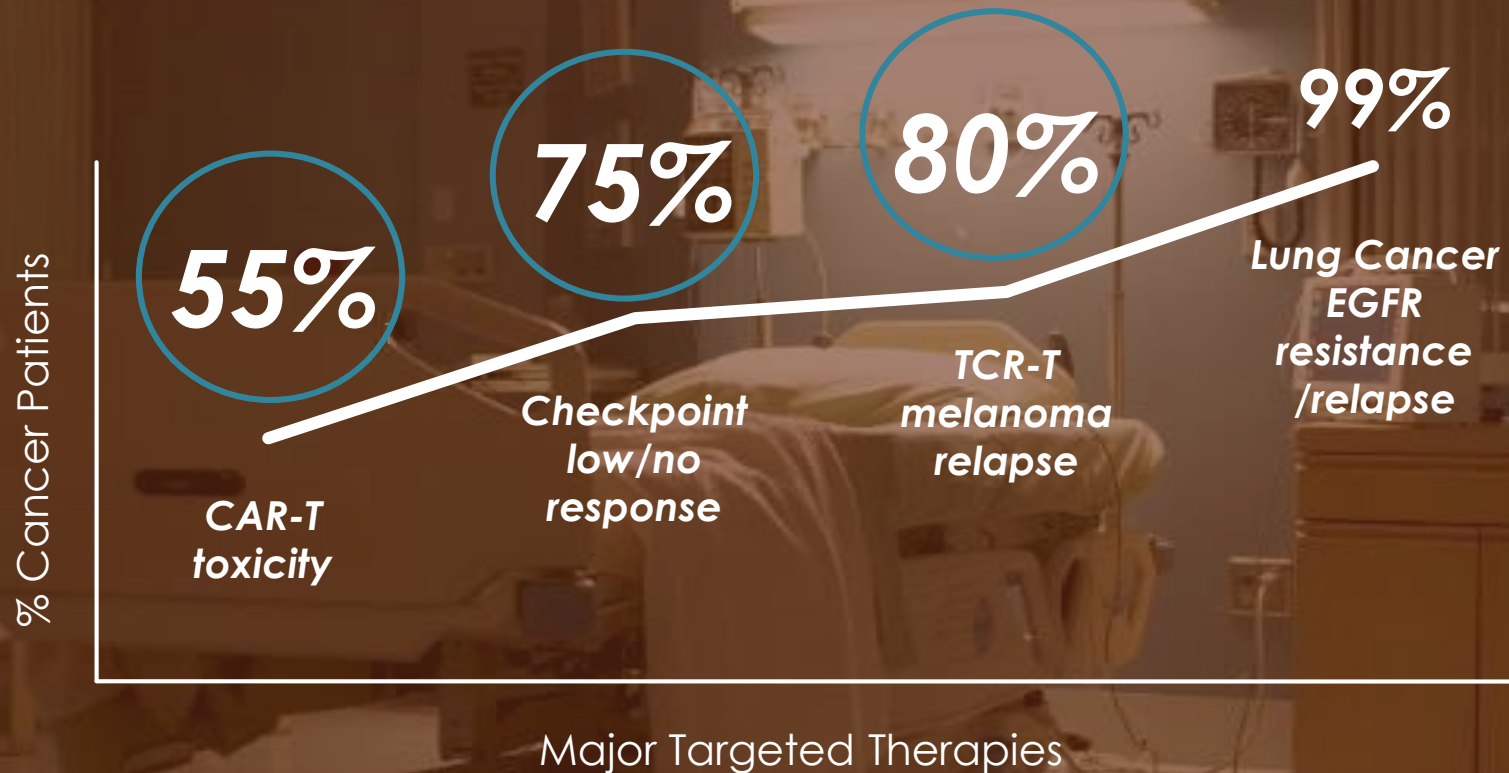


Precision Profiling



Three broader challenges in cancer immunotherapy

IsoPlexis aims to lower three major statistics in immuno-oncology



IsoPlexis founding scientists & mission

Lead immuno-oncology clinical and engineering areas,
urgent applications of predictive T-cell response



Antoni Ribas, MD, PhD
UCLA



Arnie Levine, PhD
Princeton



Jim Heath, PhD
Caltech



Rong Fan, PhD
Yale



Ross Levine, MD
MSK



David Ho, MD
ADARC

Functional capacity of antigen-specific T cells

Polyfunctionality correlates with quality and durability in patient responses

Nature Reviews Immunology **8**, 247–258 (1 April 2008) | doi:10.1038/nri2274

T-cell quality in memory and protection:
implications for vaccine design

Robert A. Seder , Patricia A. Darrah & Mario Roederer

T cells mediate effector functions through a variety of mechanisms. Recently, multiparameter flow cytometry has allowed a simultaneous assessment of the phenotype and multiple effector functions of single T cells; the delineation of T cells into distinct functional populations defines the quality of the response. New evidence suggests that the quality of T-cell responses is crucial for determining the disease outcome to various infections. This Review highlights the importance of using multiparameter flow cytometry to better understand the functional capacity of effector and memory T-cell responses, thereby enabling the development of preventative and therapeutic vaccine strategies for infections.

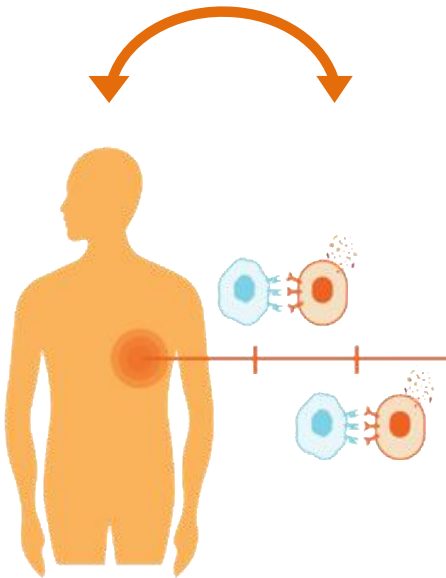


Proposes that **polyfunctionality**, the ability for a T cell to co-produce multiple cytokines, is likely a better correlate to the quality of T cells in memory and protection.

IsoPlexis uniquely addresses challenges in I-O

IsoCode assay links T-cell function to patient outcome

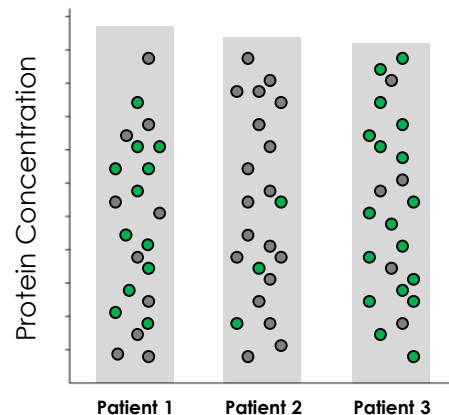
Link



IsoCode's more sensitive detection can lead to

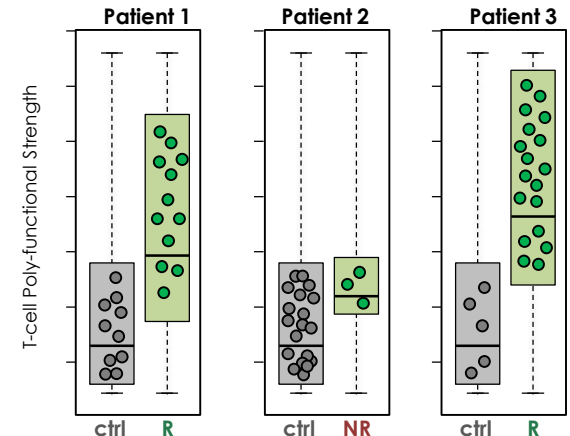
- Development feedback and excellence
- Improvement in patient management:
Predictive biomarkers

Bulk blood or T-cell proteins



T-cell function looks similar, despite differences in outcome

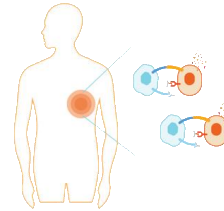
IsoPlexis single T-cell function



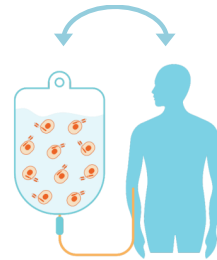
Detect patient T-cell functional differences, correlated to outcome

IsoPlexis recent data: T-cell function links to outcome

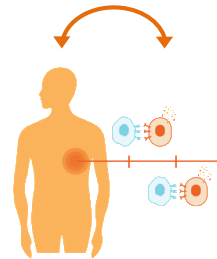
Predictive single-cell response applied to I-O



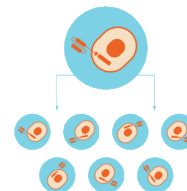
TIL function predicts response and non-response in PD1/CTLA4 treatment



Pre-infusion product profiles predict CAR-T Patient Response



TCR-T product tracks / predicts response and relapse



CAR-T Product Characterization for Manufacturing

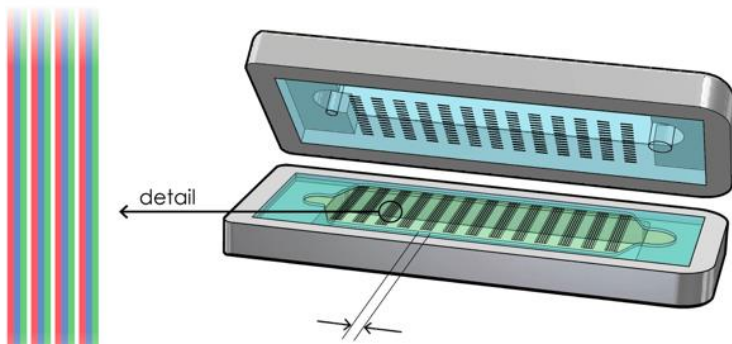
IsoCode Platform Technical Overview & Validation

IsoPlexis: **I**solated single-cells, Multi**P**lexed data

Precision T-cell Profiling enabled by the IsoCode Chip

Sensitivity captured

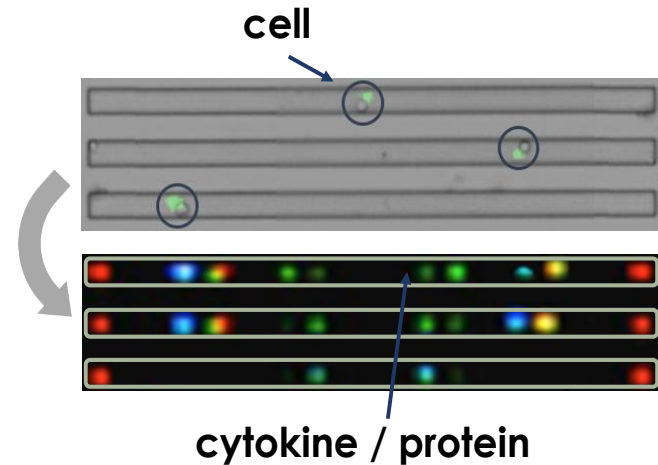
IsoCode Single-cell Chip



Single T-cell & high throughput

Depth of data required

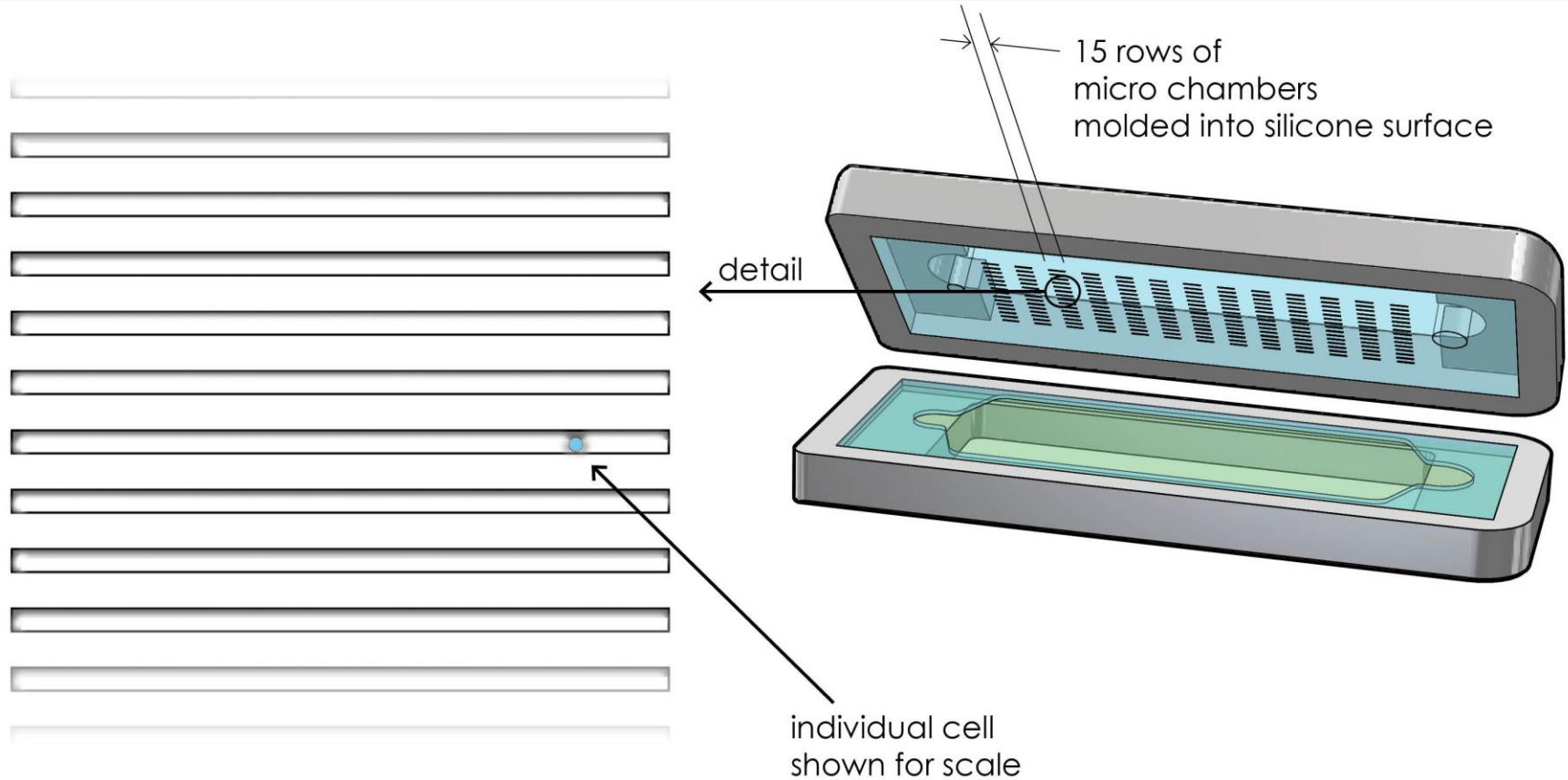
ELISA 40+plex secreted proteins, per cell



Required T-cell functions

IsoCode Chip: Cells

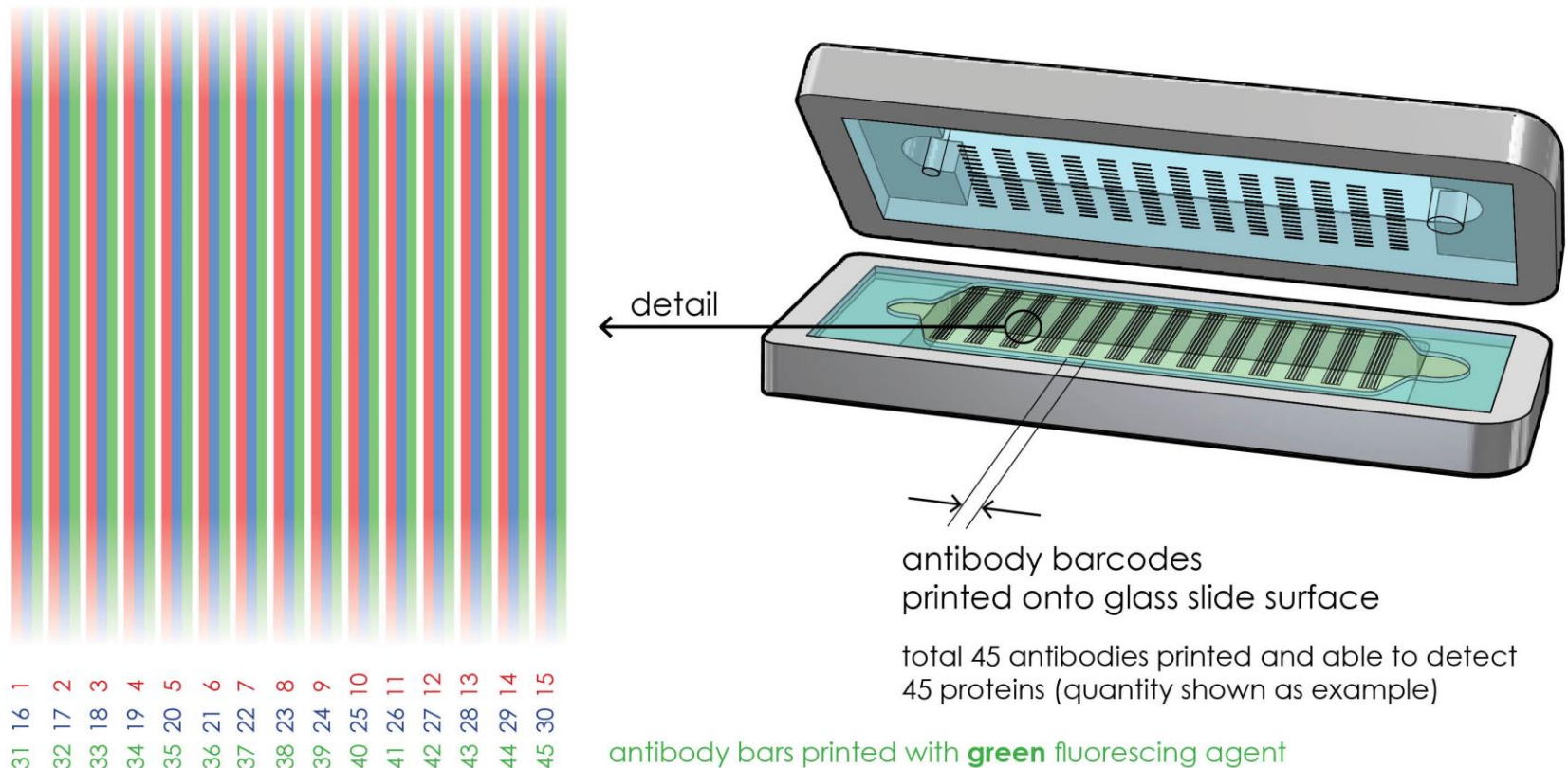
Micro-chamber cell capture



micro-chambers are molded into silicone top surface

IsoCode Chip: Array

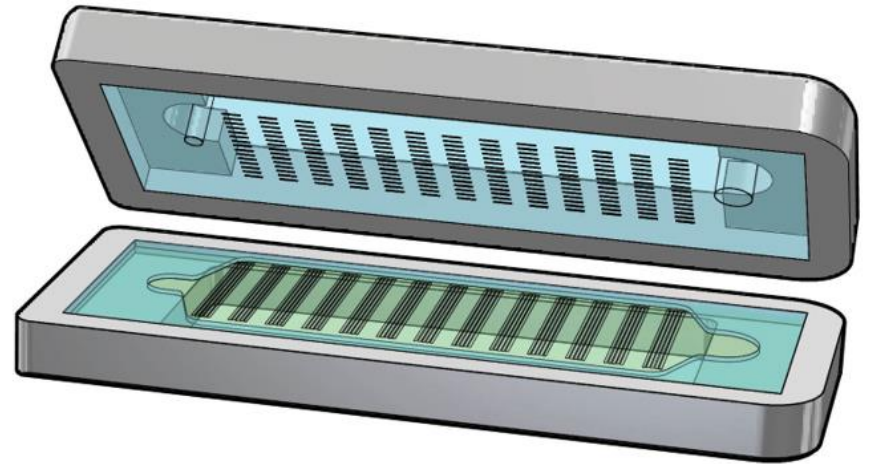
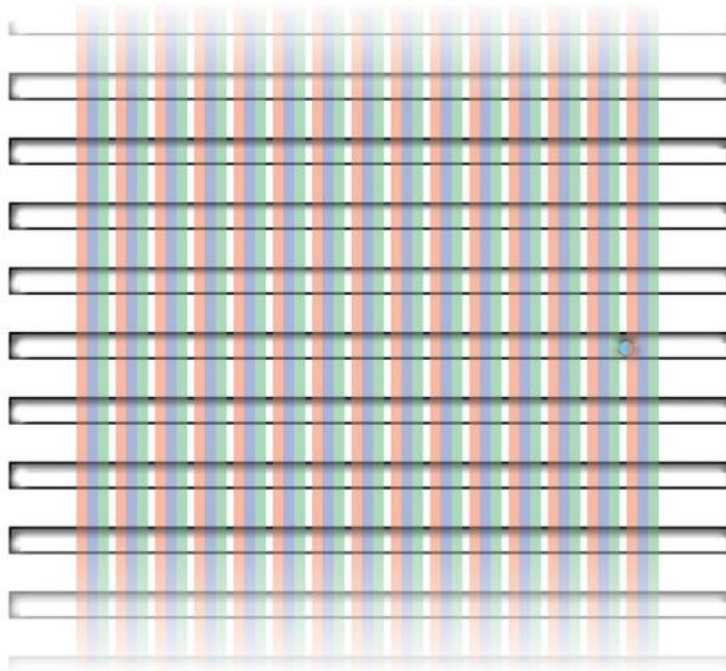
High-density antibody barcode array



antibody barcodes are printed onto glass bottom surface

IsoCode Chip: Cell Enclosure

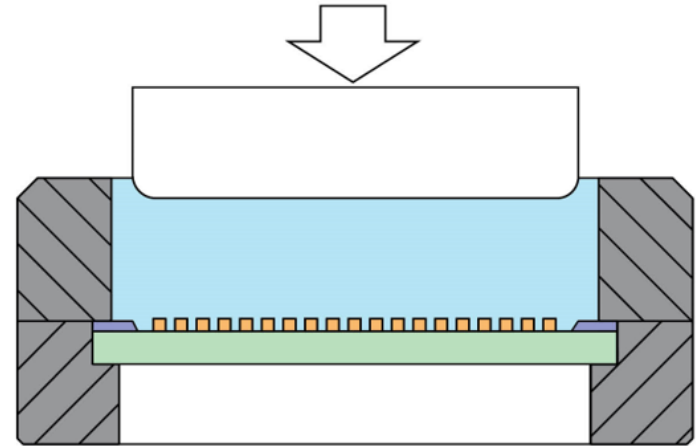
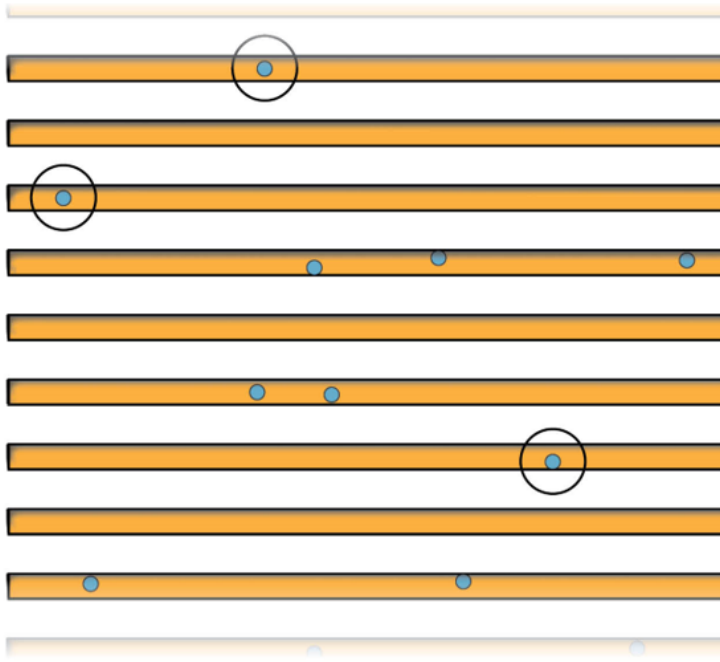
Combining micro-chambers with antibody barcode



micro-chambers are combined with the antibody barcode for highly multiplexed detection

IsoCode Chip: Cell Identification – Imaging Step 1

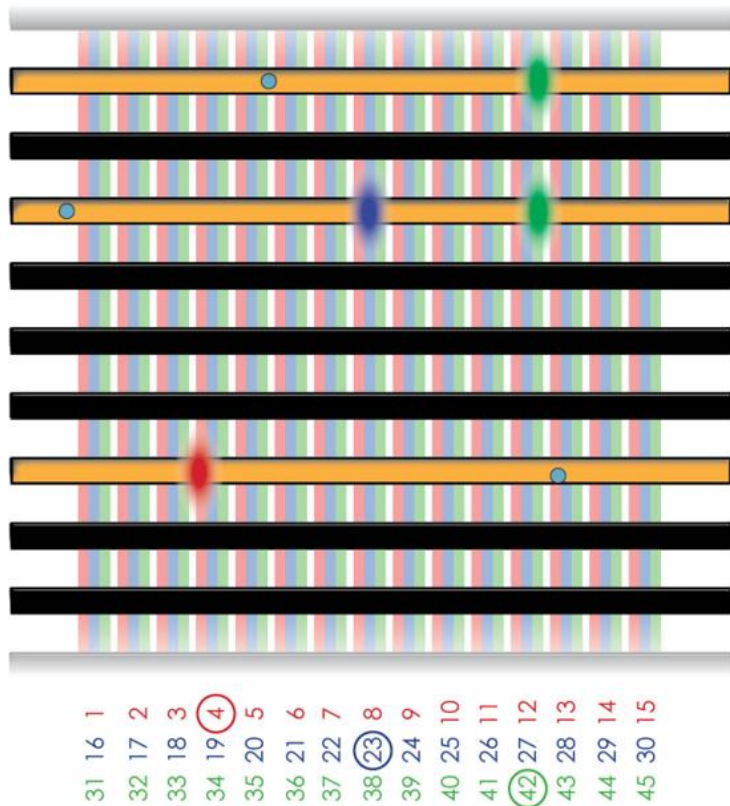
Automatic cell detection and counting: 3 surface markers per cell



Scan identifies micro-chambers containing single cells

IsoCode Chip: Protein Quantitation – Imaging Step 2

Single cell protein profile: 32+ secreted proteins per cell



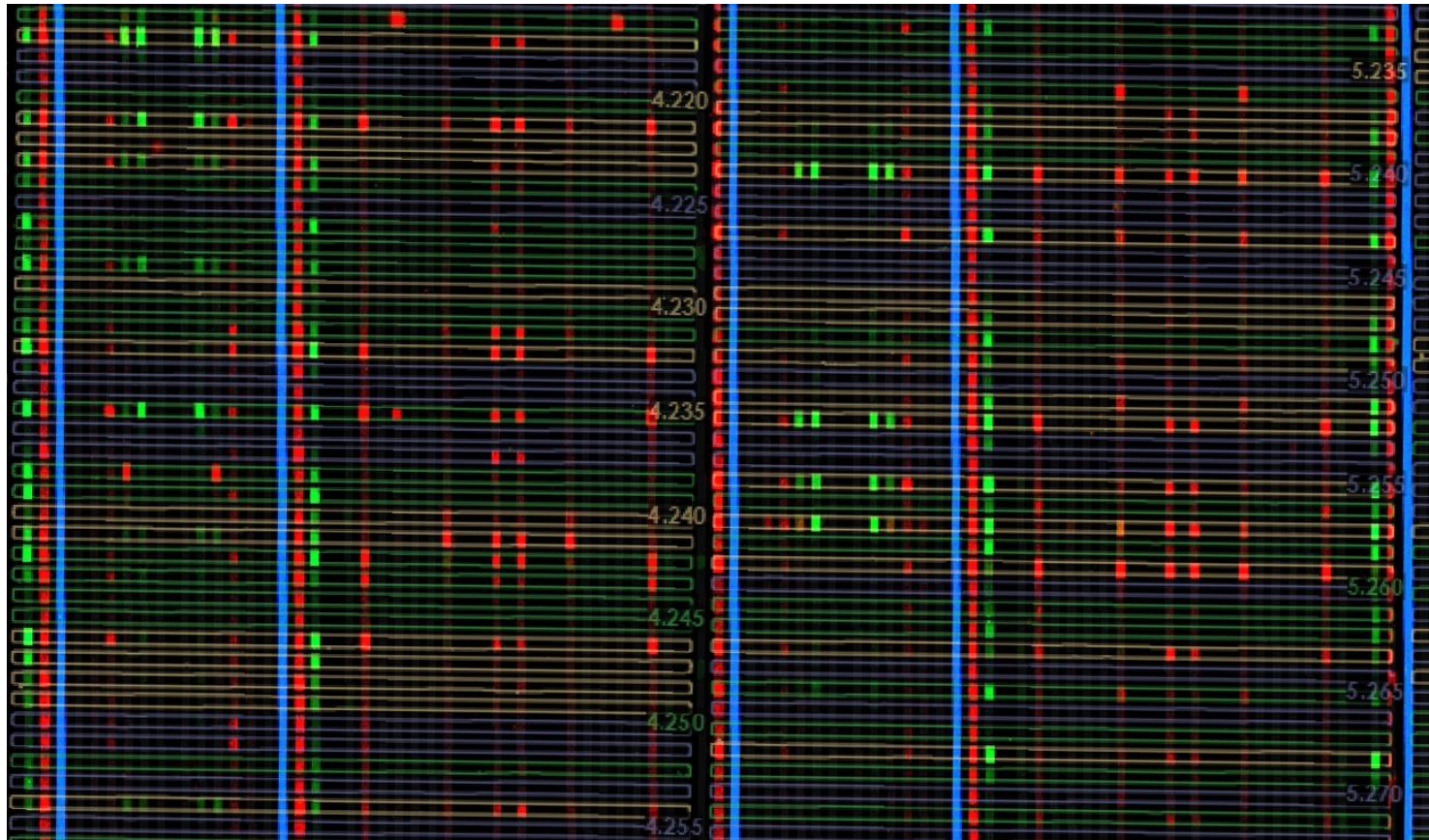
- IsoCode Chips are incubated 12 - 24 hours, causing each captured cell to secrete proteins throughout each individual micro-chamber
- each secreted protein is captured onto a corresponding antibody bar
- reagent is then introduced into the flow path, causing the captured proteins to fluoresce

scan disregards micro-chambers containing multiple or no cells

Incubated cells secrete proteins which bond to antibody bars for identification

IsoCode Chip: Actual Protein Readout, Per Chamber

Representative IsoCode Chip data



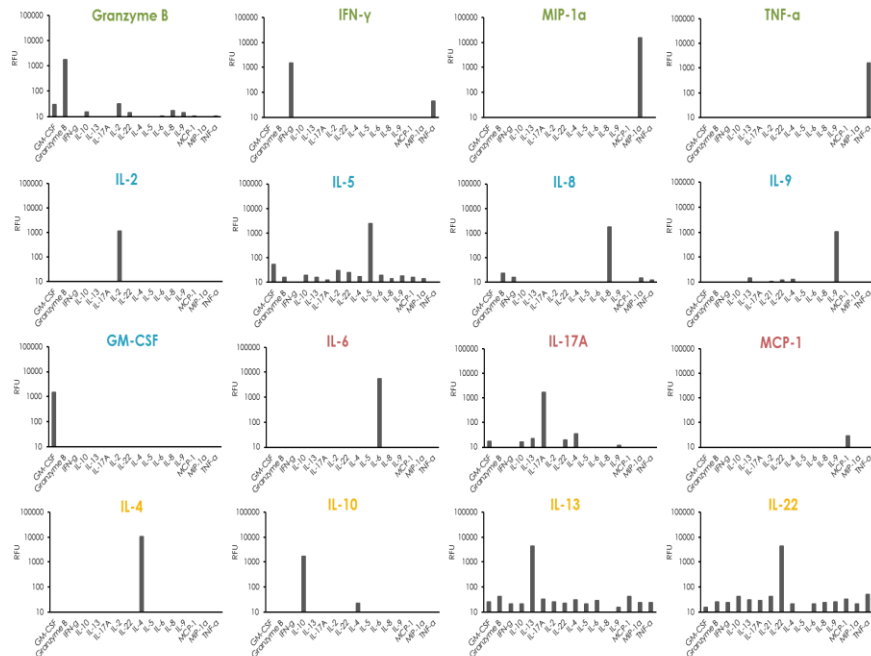
scaled detail of actual IsoCode Chip with fluorescing proteins

Validation: QC and calibration of antibody panels

Antibody panels are rigorously validated for both sensitivity and specificity

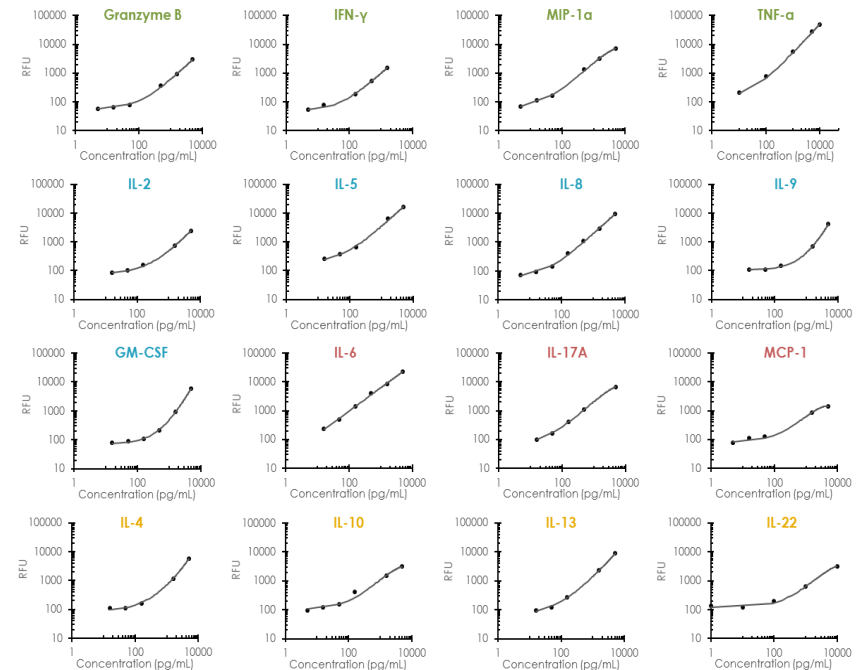
A) rigorous specificity of antibody capture

no cross reactivity of antibodies; SNR of 10



B) sensitivity and linearity of antibody pairs

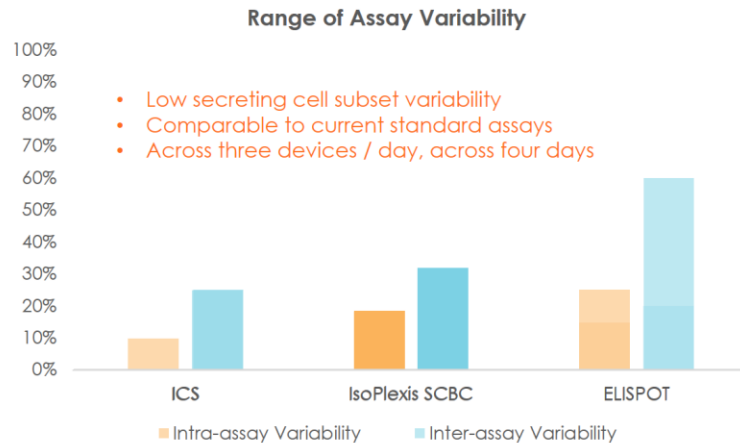
dynamic range of ~5 to 5000 pg / ml



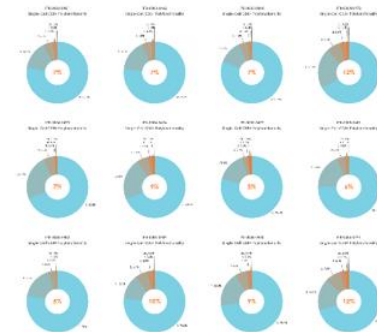
Validation: Low T-cell subset CV across metrics

Low variability leads to consistent data

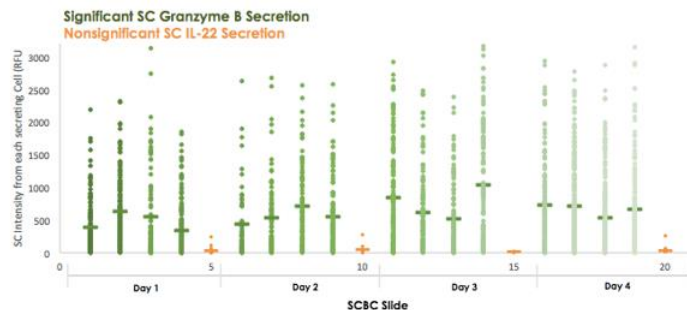
A) low secreting cell subset variability: 18.5% - 32%



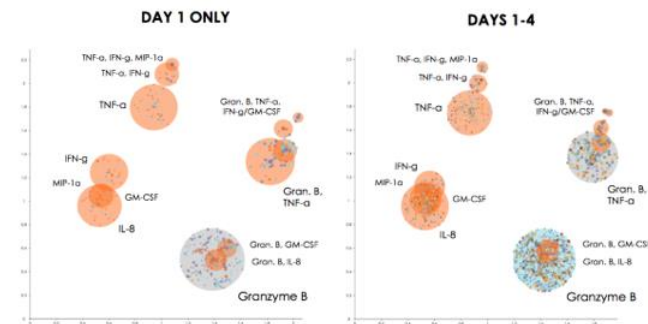
B) poly-functionality cell subset CV of 28.8% across samples & days



C) cell subset signal CV of 28.5% across 12 tests of same samples



D) PAT PCA subsets conserved as well across 12 tests of same samples

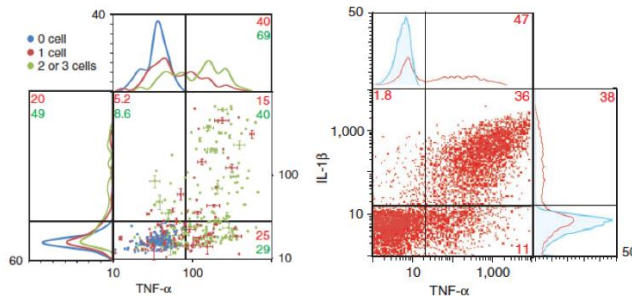


confidential

Validation: IsoCode compared to other immunoassays

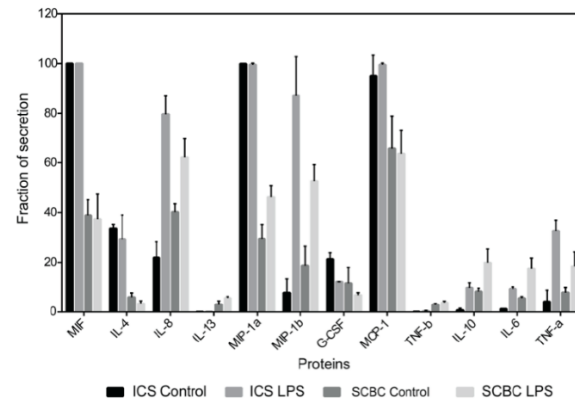
IsoCode has strong correlation with traditional low-plex single cell assays

SCBC vs Flow ICS



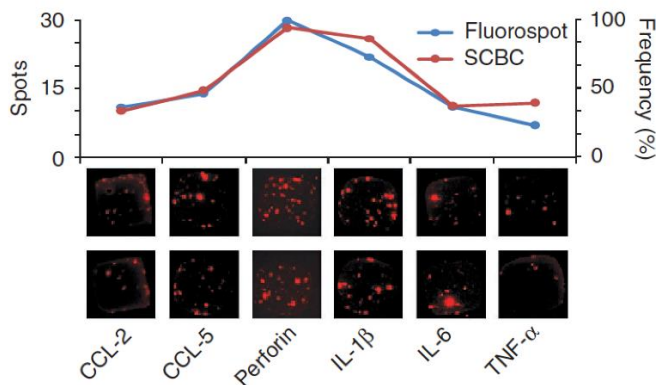
Ma, Fan, et al., *Nature Medicine*, 17, 738-743 (2011).

SCBC vs Flow ICS



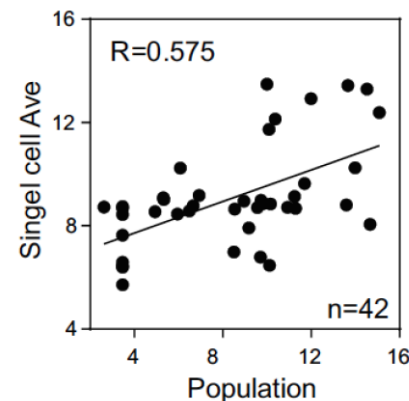
Lu, Xue, et al., *Proc. Natl. Acad. Sci. U.S.A.*, 112(7), 607-615 (2015).

SCBC vs FLUOROSpot



Ma, Fan, et al., *Nature Medicine*, 17, 738-743 (2011).

SCBC vs Population

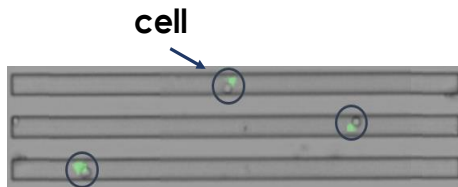


Lu, Xue, et al., *Proc. Natl. Acad. Sci. U.S.A.*, 112(7), 607-615 (2015).

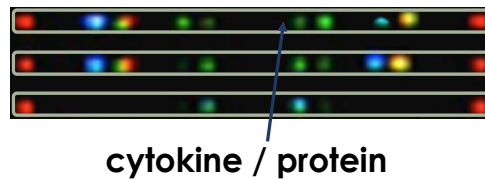
Informatics: Ease of understanding, implementation

The IsoPlexis workflow enables validated analysis in larger trials

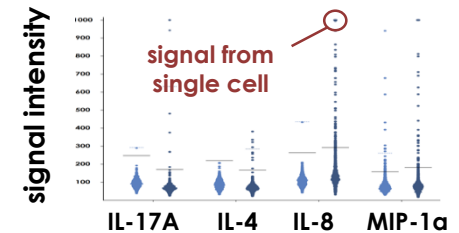
DETECT CELLS



DETECT PROTEINS

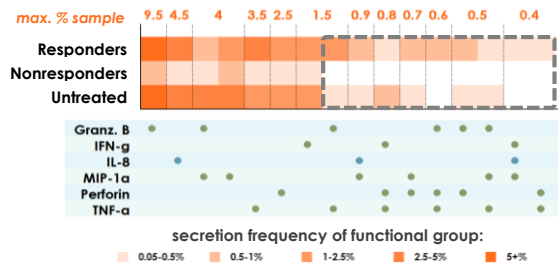


QUANTIFY READOUT



PROFILE DONORS

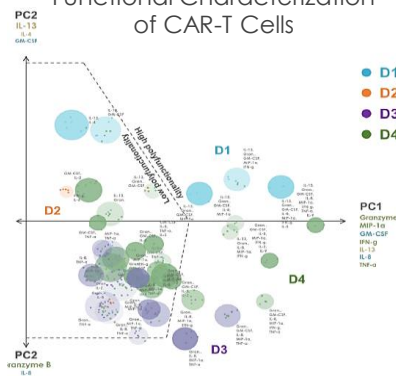
Functional Heat Map of Anti-PD-1 Treated and Untreated Patient Groups



Presented at ASCO 2017

IDENTIFY DIFFERENCES

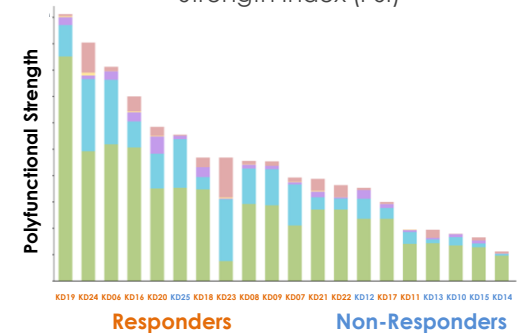
Functional Characterization of CAR-T Cells



Presented at BioMAN Workshop 2016

CORRELATE TO OUTCOME

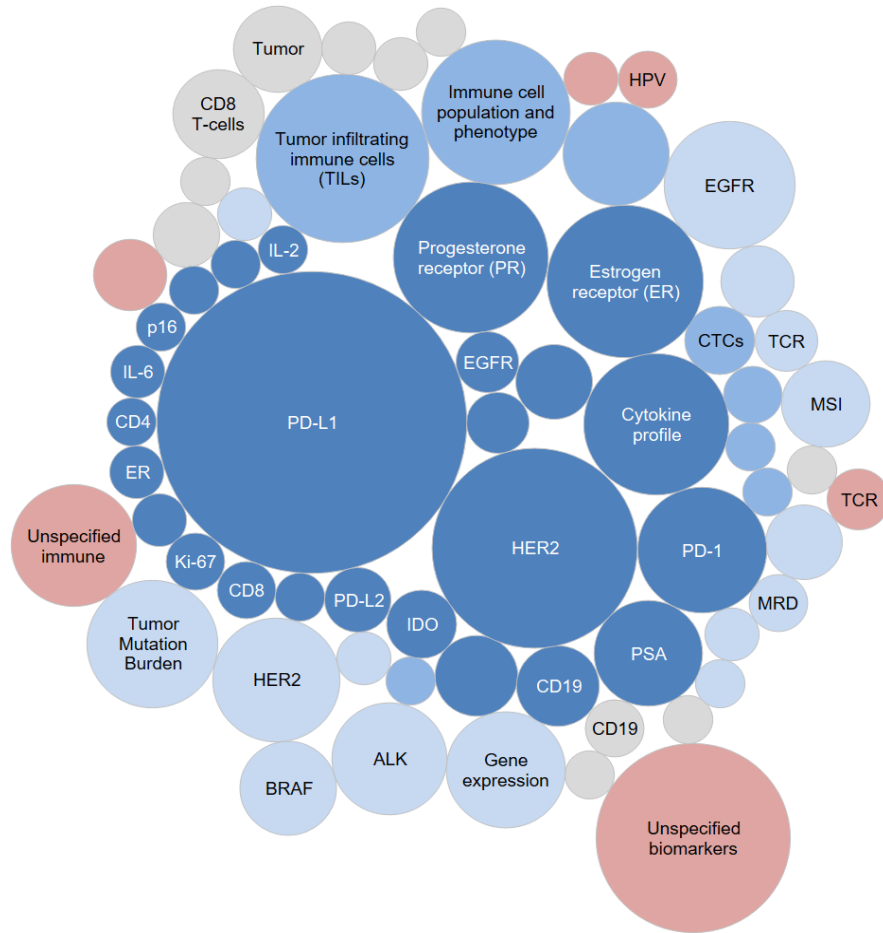
Single-Cell CAR-T Polyfunctional Strength Index (PSI)



Presented at AACR 2017

Immuno-Oncology Biomarkers: Landscape Summary

DeciBio Report



Immuno-Oncology Biomarkers mentioned in at least 5 different clinical trials.

The top “novel” immuno-oncology biomarkers identified in this analysis are:

- Tumor-infiltrating immune cells (mentioned 70 times in 61 different trials)
- Immune cell populations / phenotypes (mentioned 60 times in 45 different trials)
- **Cytokine profiles** (mentioned 52 times in 43 different trials)
- Tumor mutation burden / genomic mutation profiling (mentioned 45 times in 35 different trials)
- Gene expression signatures (mentioned 33 times in 29 different trials)

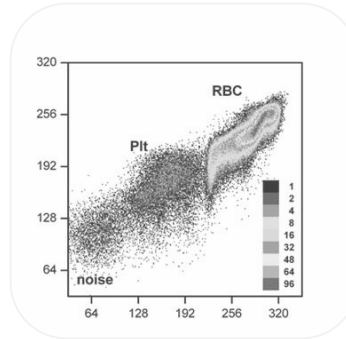
Technology used for T-cell characterization

IsoPlexis links T-cell functional data to mode of action

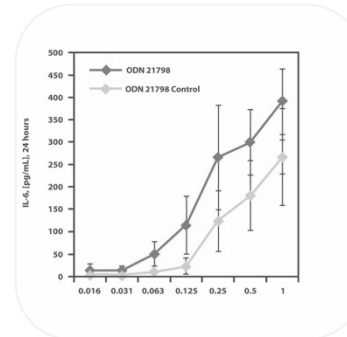
Patient Gene Profile



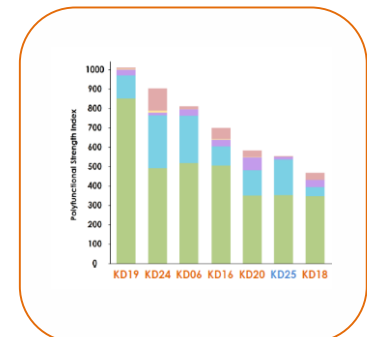
Cell Type Frequency



Blood Levels



T cell Functional Capacity



System	Gene expression	Flow and mass cytometry	ELISA and Luminex	IsoCode
Correlation to MOA in T cells	x	x	x	✓
Single Cell	✓	✓	x	✓
Multiplexed secreted protein data	x	x	✓	✓

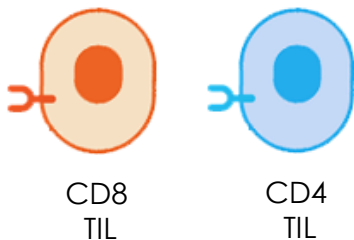
I-O Case Studies

Tumor dissociation and stimulation to ensure specific readouts

Sample Enrichment

Sample: Fresh melanoma samples are immediately processed upon arrival to single-cell suspensions.

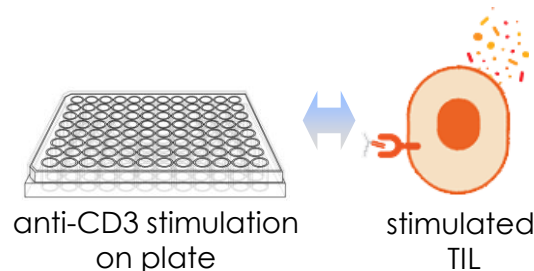
CD8+ and CD4 TILs are enriched by anti-CD8 and anti-CD4 microbeads respectively.



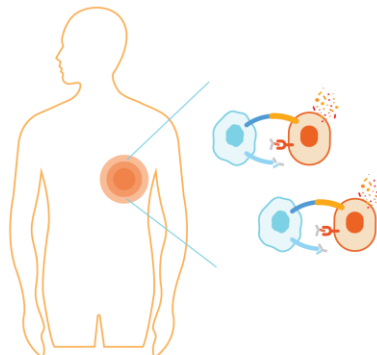
Number of cells dependent on donor, tumor, and viability

Antigen Stimulation

Stimulated: anti-CD3 stimulation at 37°C, 5% CO₂ for 20 hours.

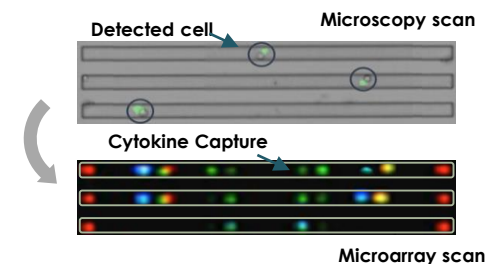
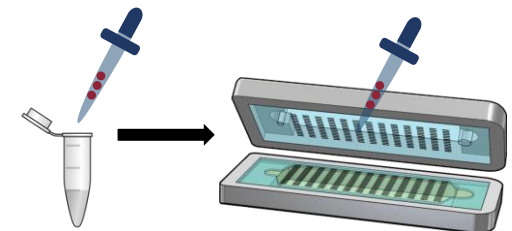


Treated vs. Untreated Patients



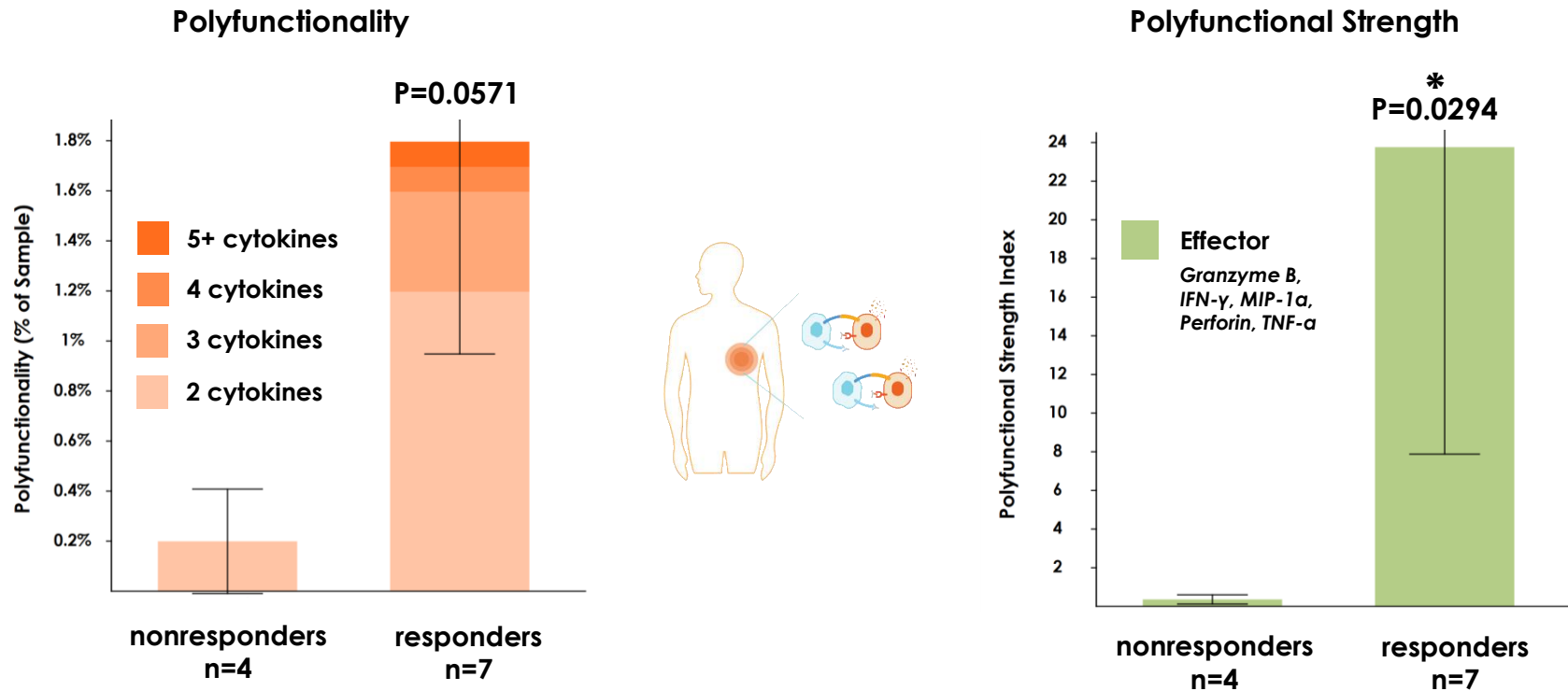
TILs Loading & Analysis

Loading: Stimulated TILs are collected, then pipetted from single-cell suspension and loaded onto IsoPlexis' IsoCode system



Case 1: checkpoint biomarkers

Profiling T Cells for Improved Patient Response Markers to anti PD-1 and/or anti CTLA4 therapy

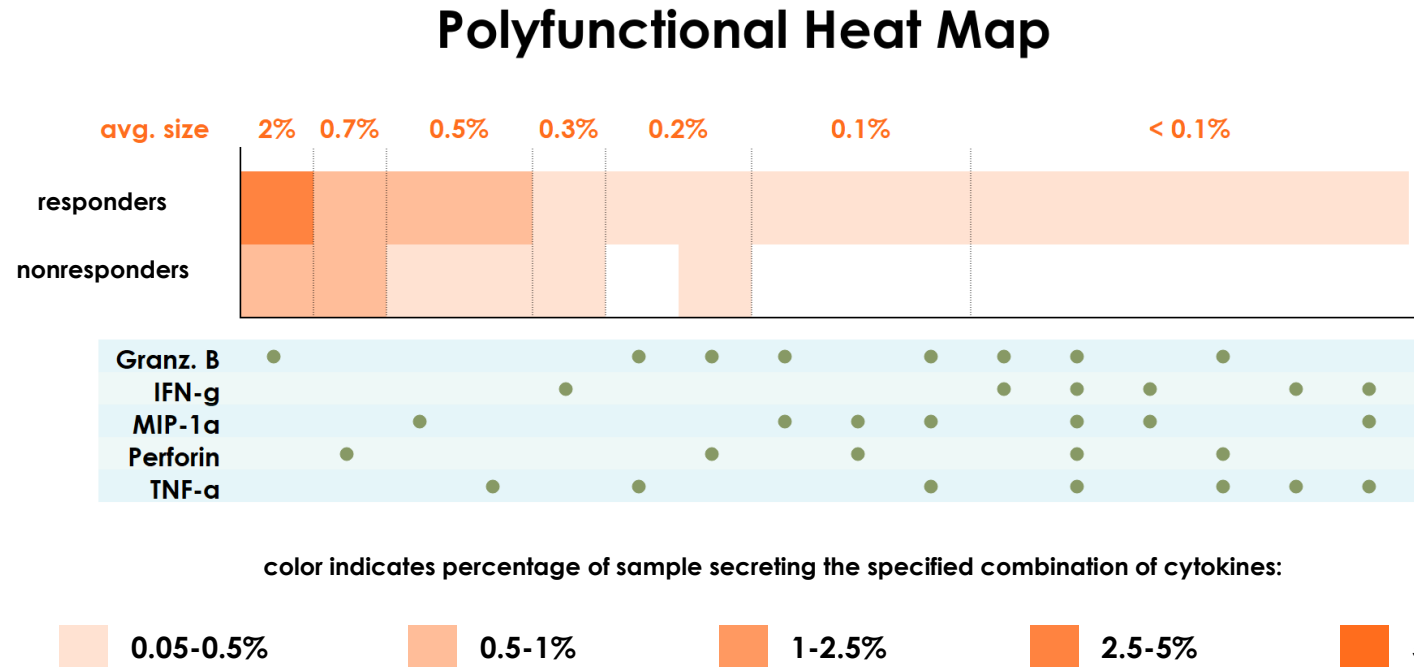


Objective: Clarity of mechanism & markers of patient selection

* P<0.05, Mann-whitney U test)
• All detail on type of response is available in appendix, and that response includes mixed responders and resistant disease in otherwise responders

Case 1: checkpoint biomarkers

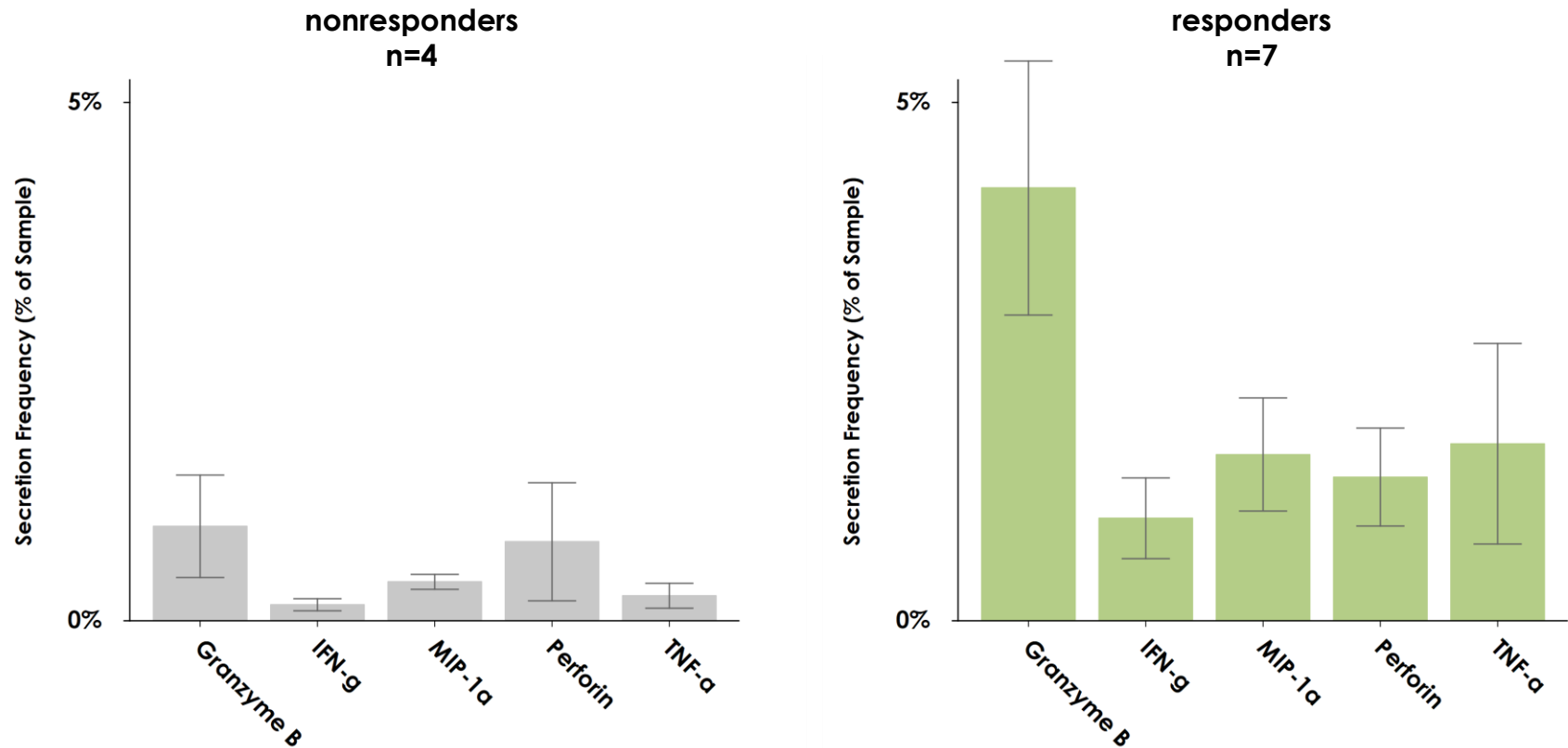
Emergence of unique polyfunctional cell subsets



Identify unique polyfunctional cell subsets in patients who responded to anti PD-1 and/or anti CTLA4 therapy

Case 1: checkpoint biomarkers

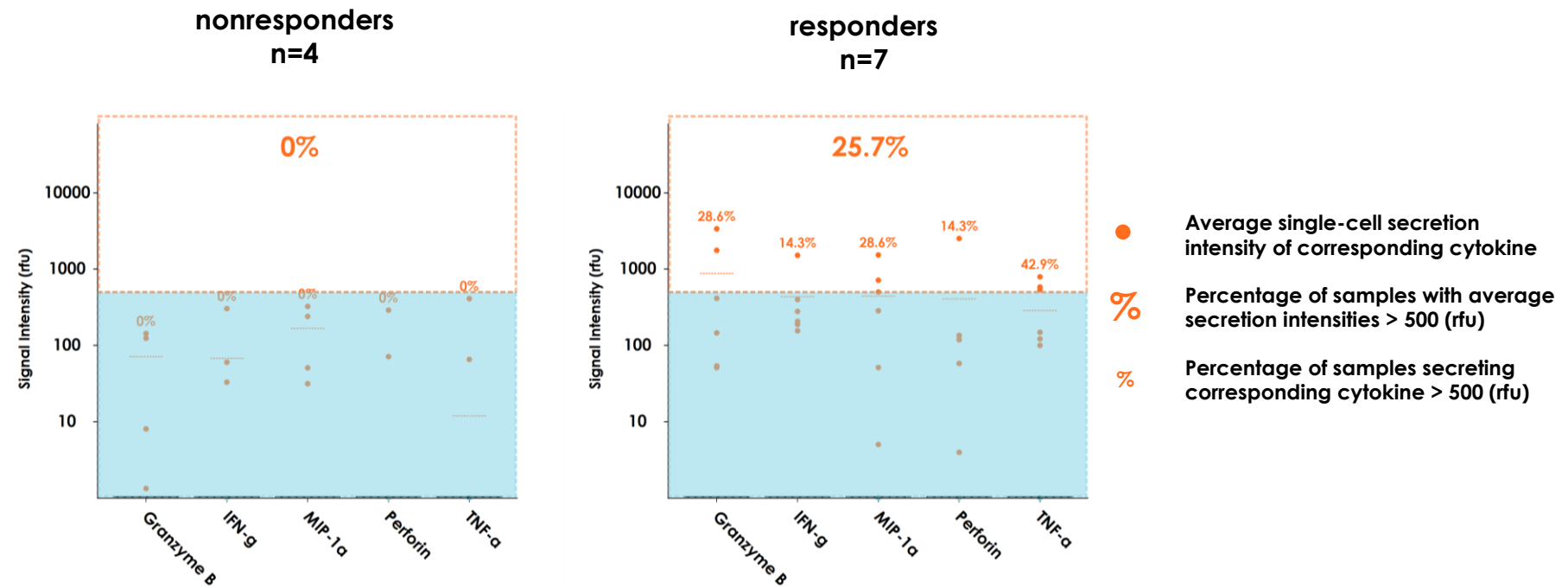
Functional differences in responding patients



Increased antitumor protein secretions in patient responding to anti PD-1 and/or anti CTLA4 therapy

Case 1: checkpoint biomarkers

Enhanced secretion intensity in responding patients

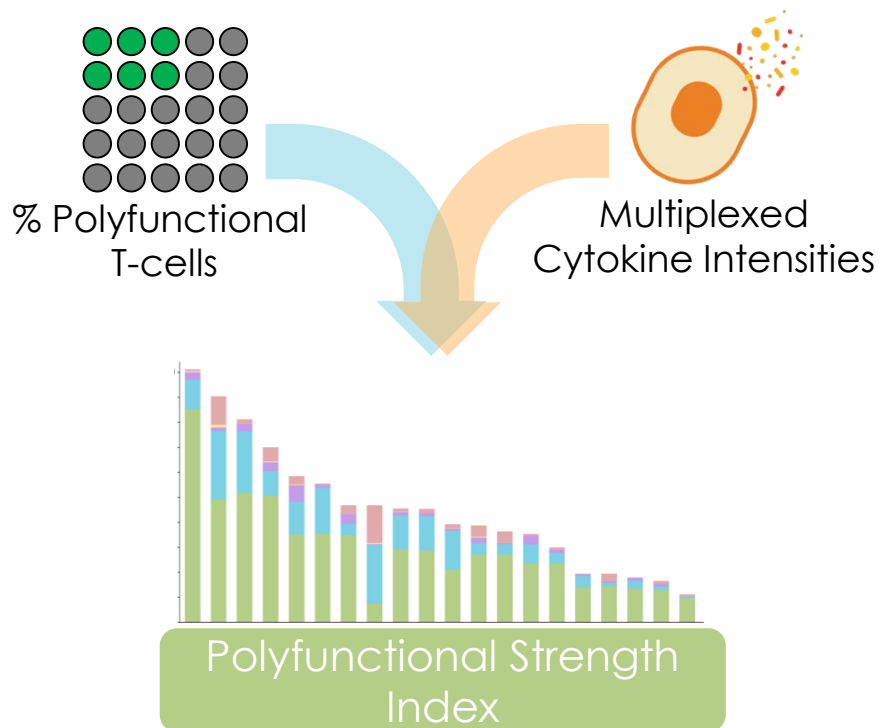


Enhanced average secretion intensities of proteins associated with antitumor immunity in patients who responded to anti PD-1 and/or anti CTLA4 therapy

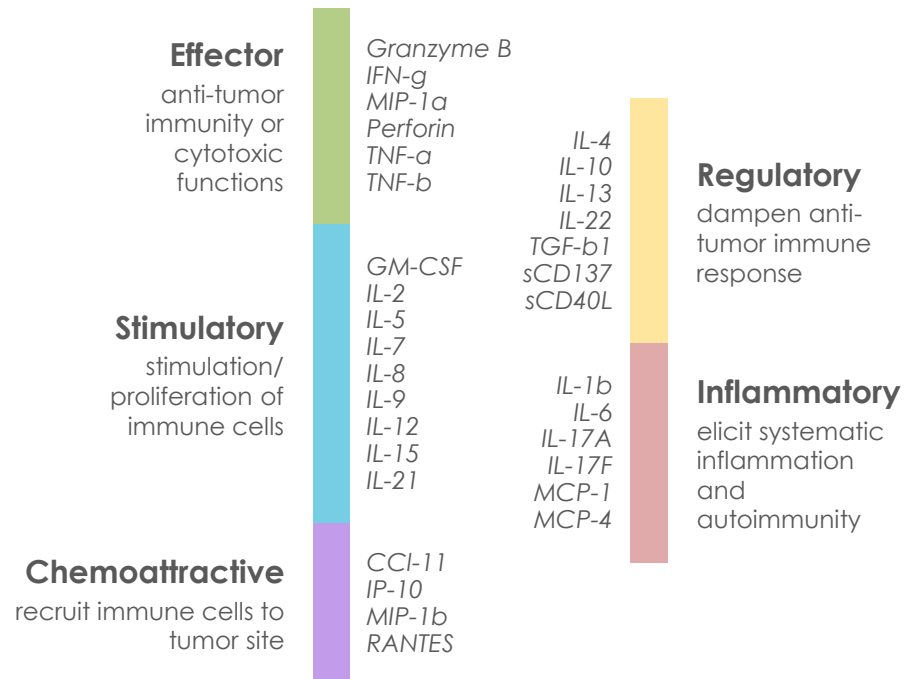
Case 2: CD19 CAR-T Polyfunctionality for Kite Product

Polyfunctional Strength Index (PSI)¹ metric links to patient outcomes

Product Metric: Requires IsoCode



IsoCode Single-Cell Panel



1. IsoPlexis IsoCode Technology and Ma et al 2013

CCL, chemokine ligand; GM-CSF, granulocyte macrophage colony-stimulating factor; IFN, interferon; IL., interleukin; IP, interferon-gamma-inducible protein; MCP, monocyte chemoattractant protein; MIP, macrophage inflammatory protein; PSI, polyfunctional strength index; RANTES, regulated on activation, normal T cell expressed and secreted; TGF, transforming growth factor; TNF, tumor necrosis factor.

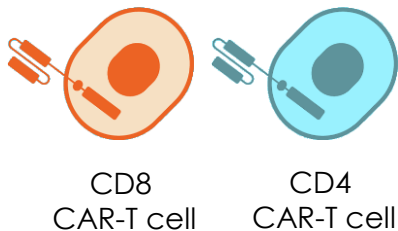
Case 2: IsoCode CAR-T Workflow Overview

Stimulation & loading to ensure antigen specific readouts

Sample Enrichment

Sample: Cryopreserved samples are thawed upon arrival to RT.

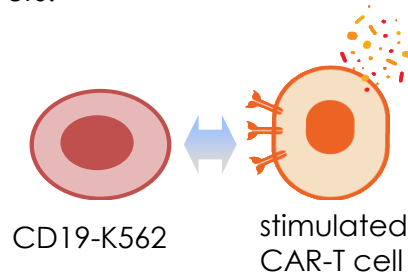
CD8+ and CD4 CAR-T cells are enriched by anti-CD8 or anti-CD4 microbeads respectively.



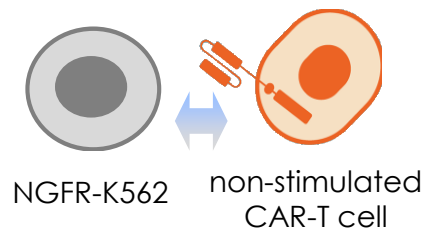
2 million cells requested per donor, viability

Antigen Stimulation

Stimulated: CD19-specific response at 37°C, 5% CO2 for 20 hours.

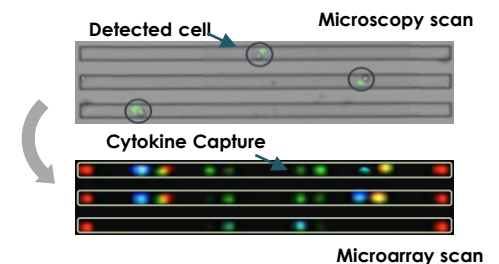
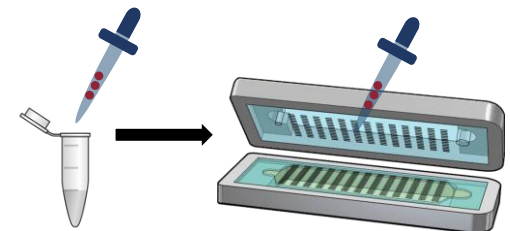


Control: Non CAR specific (allogeneic) response at 37°C, 5% CO2 for 20 hours.



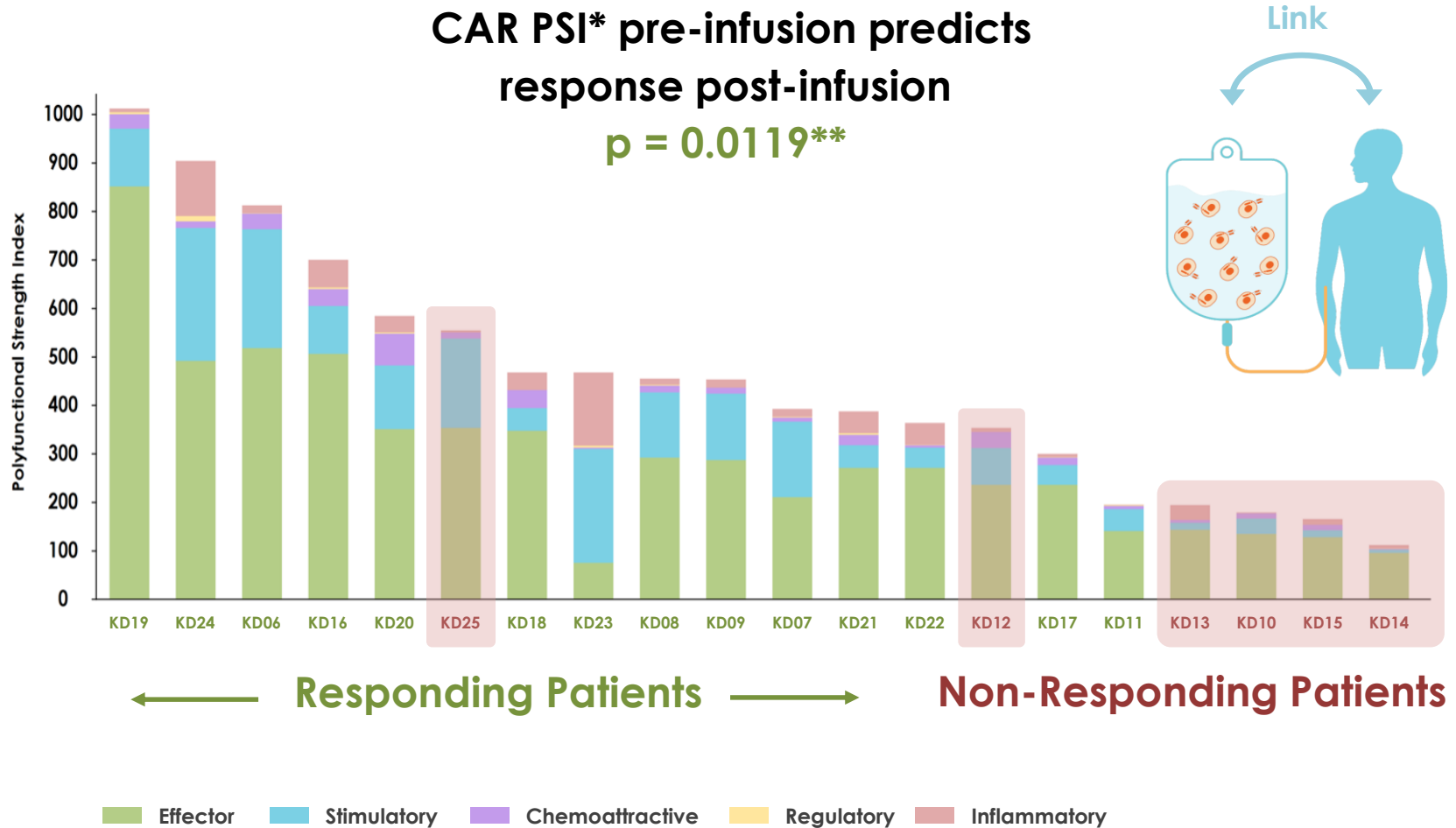
Depletion & CAR-T Loading

Depletion: Cancer target cells are depleted, then stimulated. CAR-T cells are pipetted from single-cell suspension and loaded onto IsoPlexis' IsoCode system.



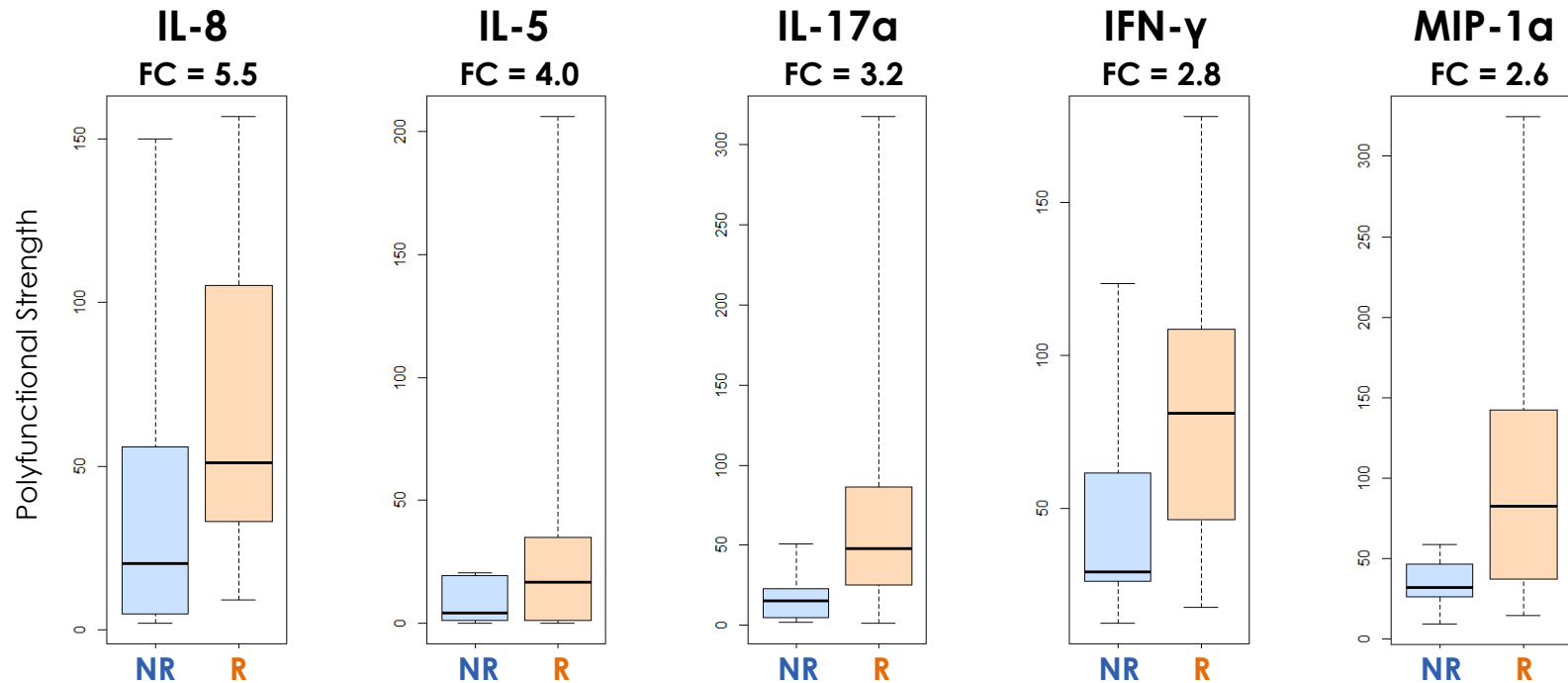
Case 2: ranking CAR-T donors by PSI

Patient pre-infusion product profiles enable early intervention



Case 2: drivers of CAR-T product potency

Further, from the readouts, non-redundant cytokines contributed to these PSI differences (CD4 ex.)



additionally, pre-infusion product is an independent variable vs. CAR-T expansion

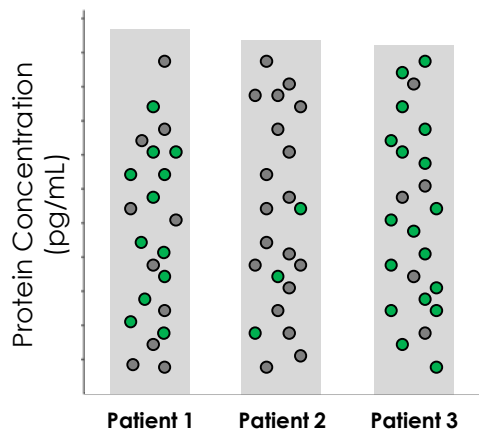
Deeper biomarkers for response / non-response

Applying improved sensitivity to patient stratification

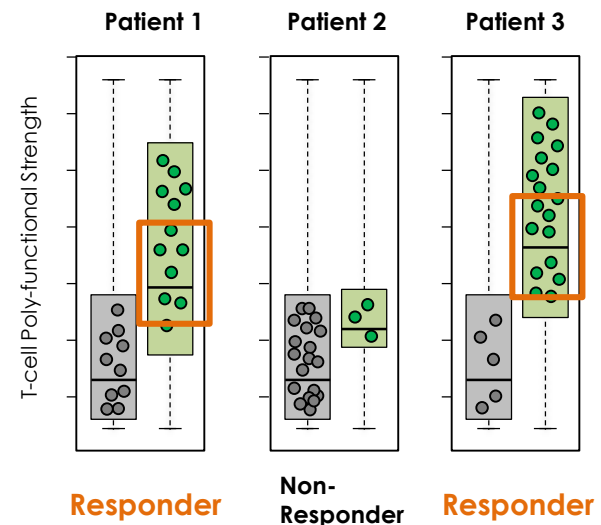
Isoplexis technology provides
data driven by & related to patient outcome

Current Methodologies

(flow cytometry / bulk assay)



Single T-cell Patient Response



- T-cell biomarkers are masked in current analysis and methodologies.
- Functional differences in cellular sub-populations are unable to be differentiated.

- Improve / optimize development and administration to achieve T-cell poly-functional potency metrics, linked to outcome
- Improve multi-dimensional assessment of patient response to enable stratification

Patient biomarker methods & differentiation

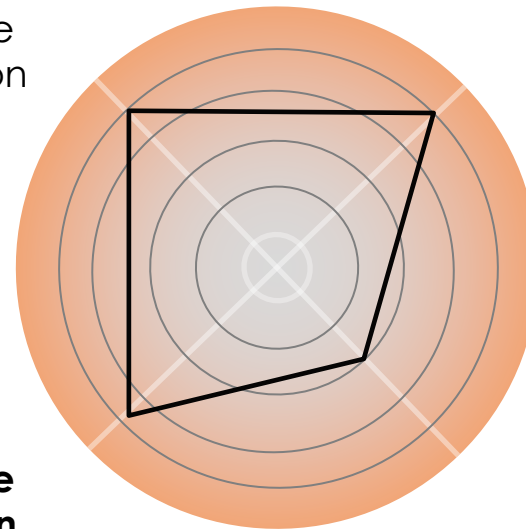
Predictive solutions add value to characterizing T-cell function

Immunotherapy Assessment
Genetic profiling
Flow and Mass Cytometry
Histology
Serum profiling
IsoCode T cell Functionality

Multiple T-Cell & Tumor Readouts for assessment necessary: ImmunoGram¹

Immune Infiltration

Tumor Foreignness



Immune Function

Checkpoint absence

IsoCode offers critical T cell function signature linked to outcome to complement tumor signature solutions

Thank You & Appendix

