

THE MANAFEST ASSAY FOR MONITORING ANTI-TUMOR IMMUNITY

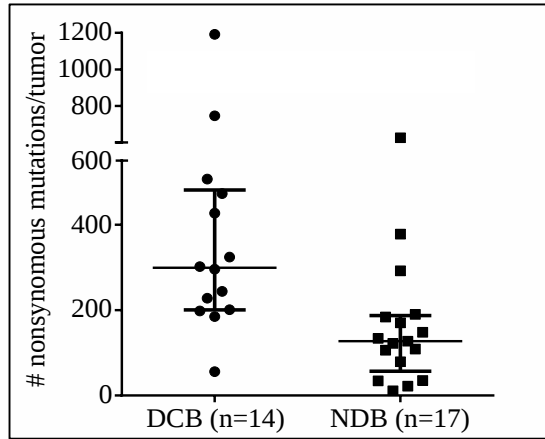
Kellie N. Smith, PhD
SITC Immuno-Oncology Biomarkers:
State of the Art
May 16, 2018
kellie@jhmi.edu



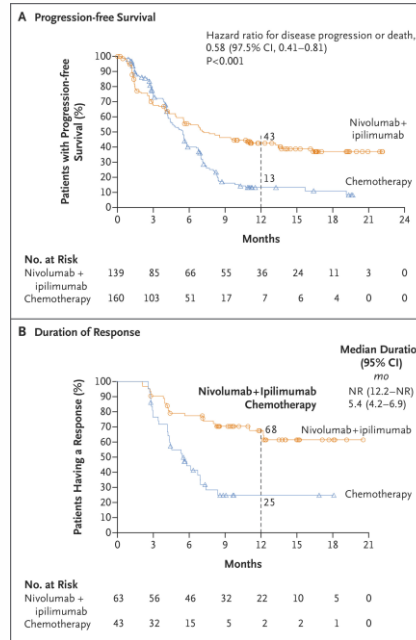
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FOR CANCER IMMUNOTHERAPY

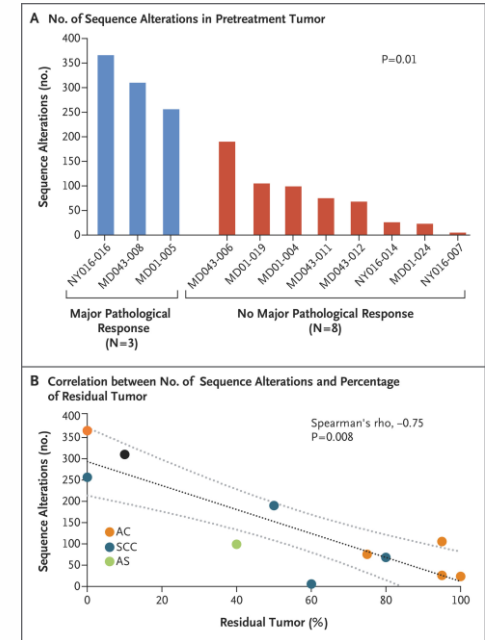
Tumor mutational burden (TMB): a predictive biomarker of response to checkpoint blockade



Rizvi et al, Science, 2015

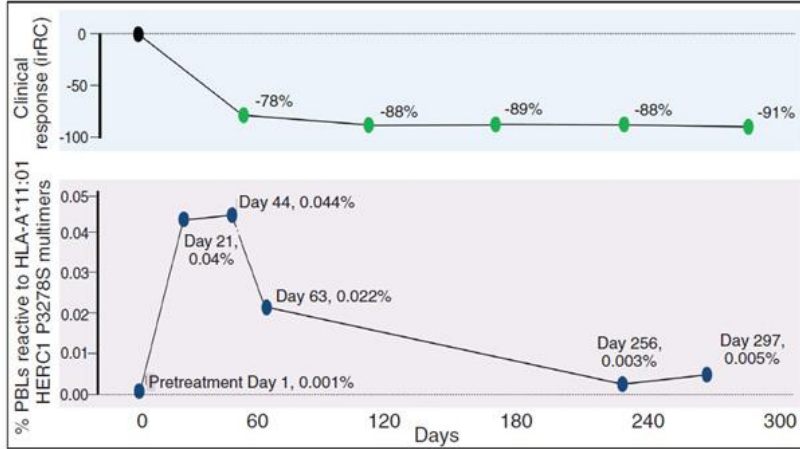


Hellmann et al, NEJM, 2018

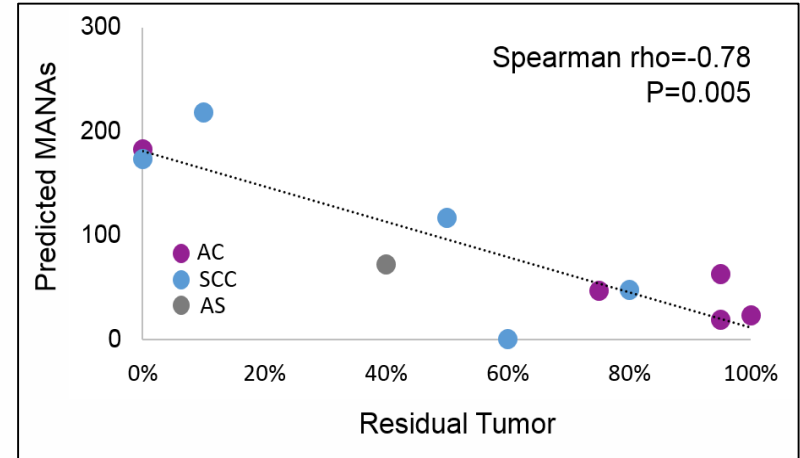


Forde, Chast, and Smith et al, NEJM, 2018

TMB and The Importance of Neoantigens



Rizvi et al, Science, 2015



Forde, Chait, and Smith et al, NEJM, 2018



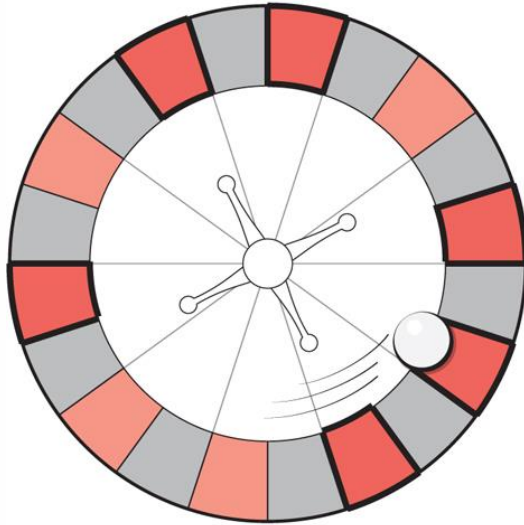
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TMB and The Importance of Neoantigens

Responders

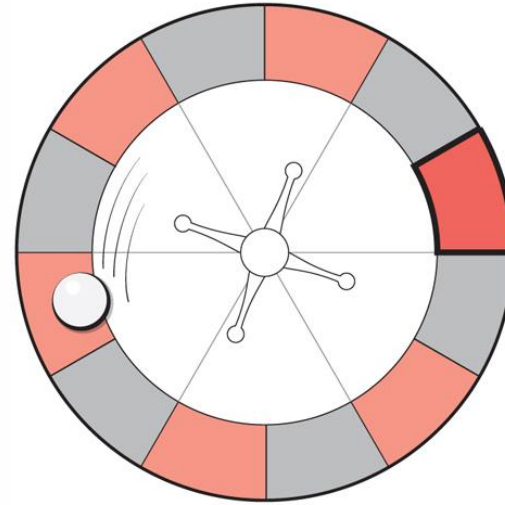
Cancer patients with more mutations have a higher chance of responding to therapy



 Mutations  Ipilimumab-responsive mutation

Nonresponders

Cancer patients with fewer mutations have a lower chance of responding to therapy



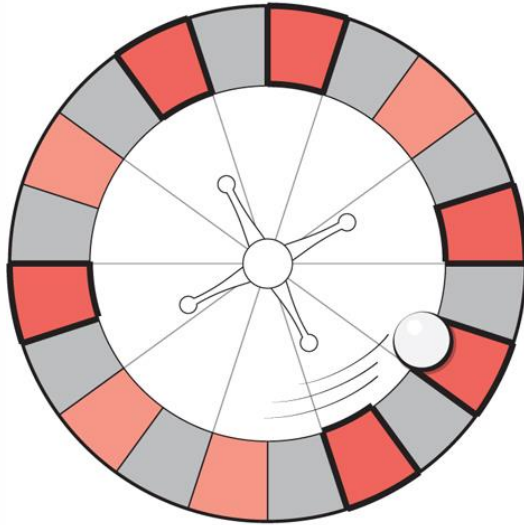
Gubin MM and Schreiber RD, Science, 2015



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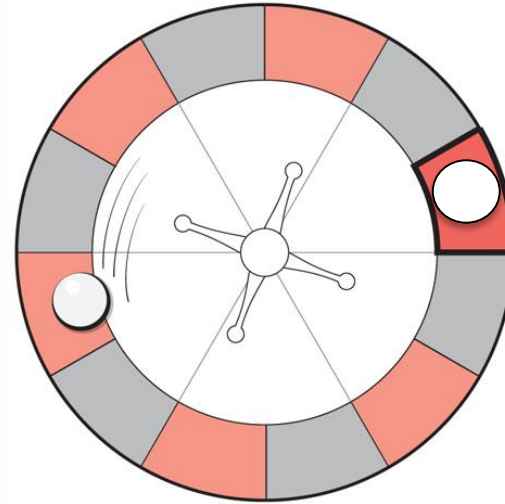
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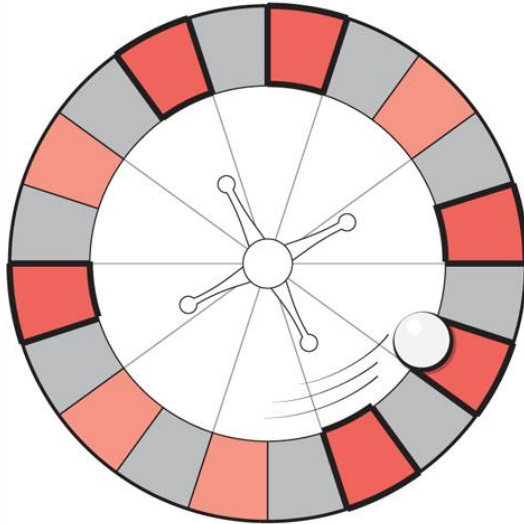
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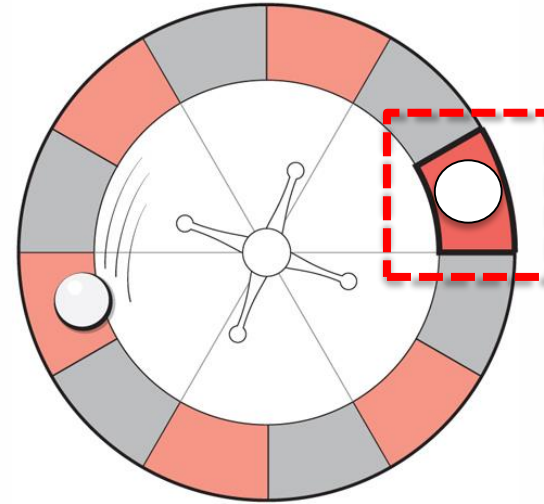
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Identifying
these patients
=
predictive
power

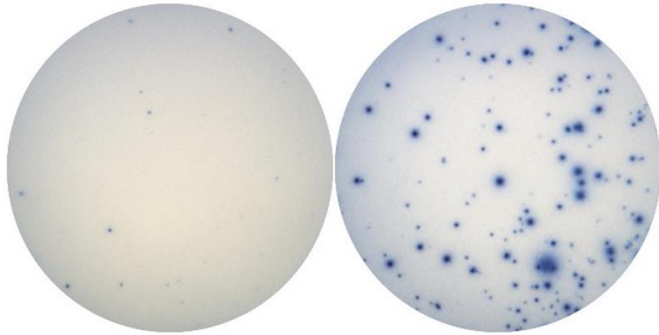
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ELISpot: The gold standard in immunological monitoring



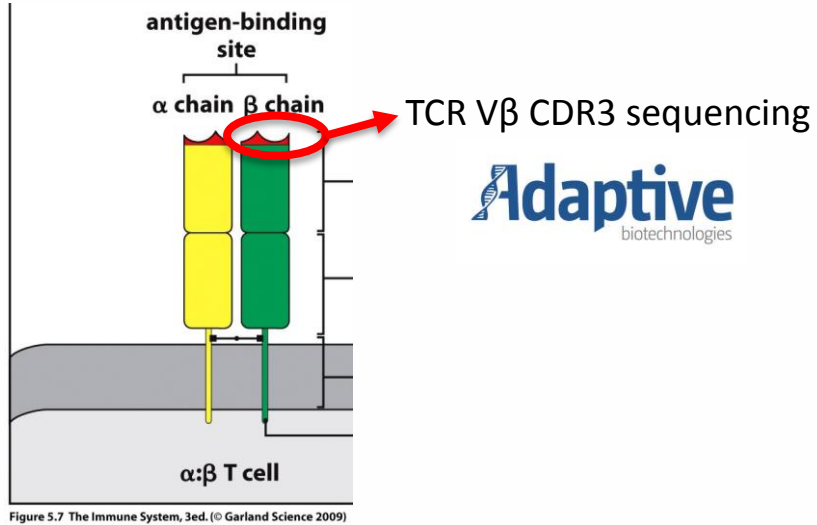
Adapted from ImmunoSpot

Pitfalls:

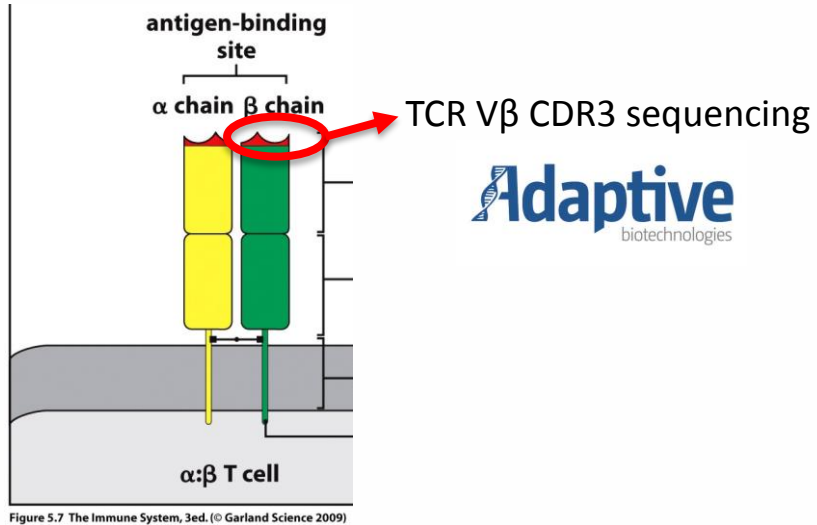
- Can't track response without repeating assay
- Can't link with FFPE specimens
- Can't track geographic distribution
- Background after culture/treated patients
- Cytokine-dependent
- No clonal dynamics



T cell receptor (TCR) sequencing



T cell receptor (TCR) sequencing

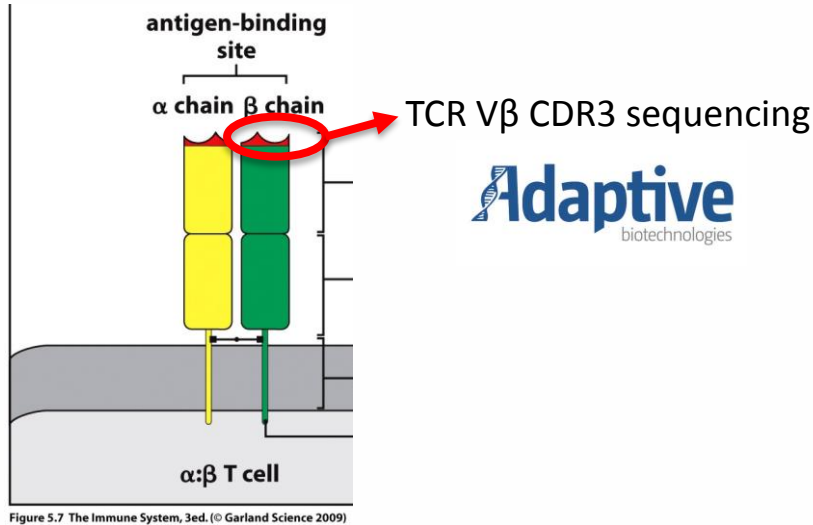


Sample types:

FFPE
Flash frozen
Viably frozen
Cultured cells



T cell receptor (TCR) sequencing



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Viably frozen
Cultured cells



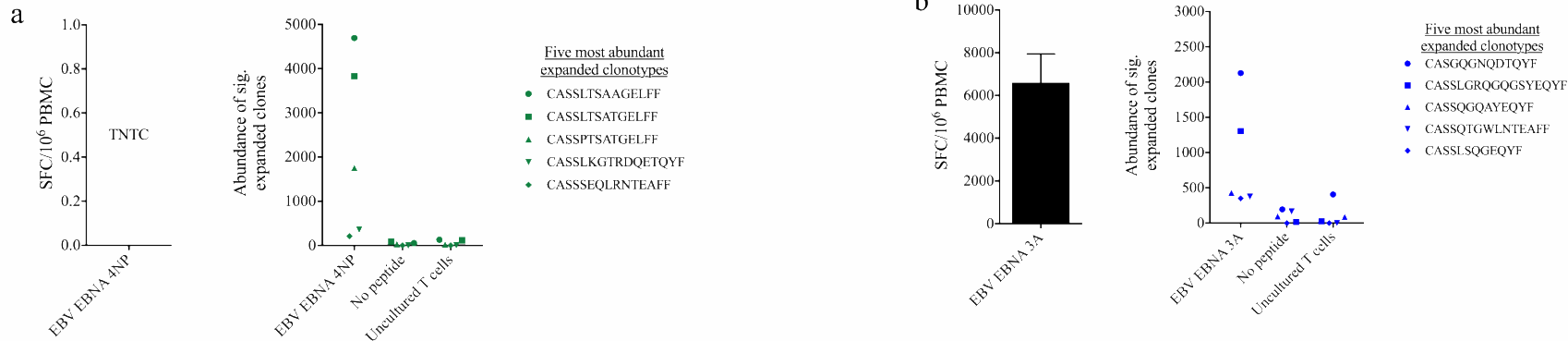
If you can get DNA, you can track it!



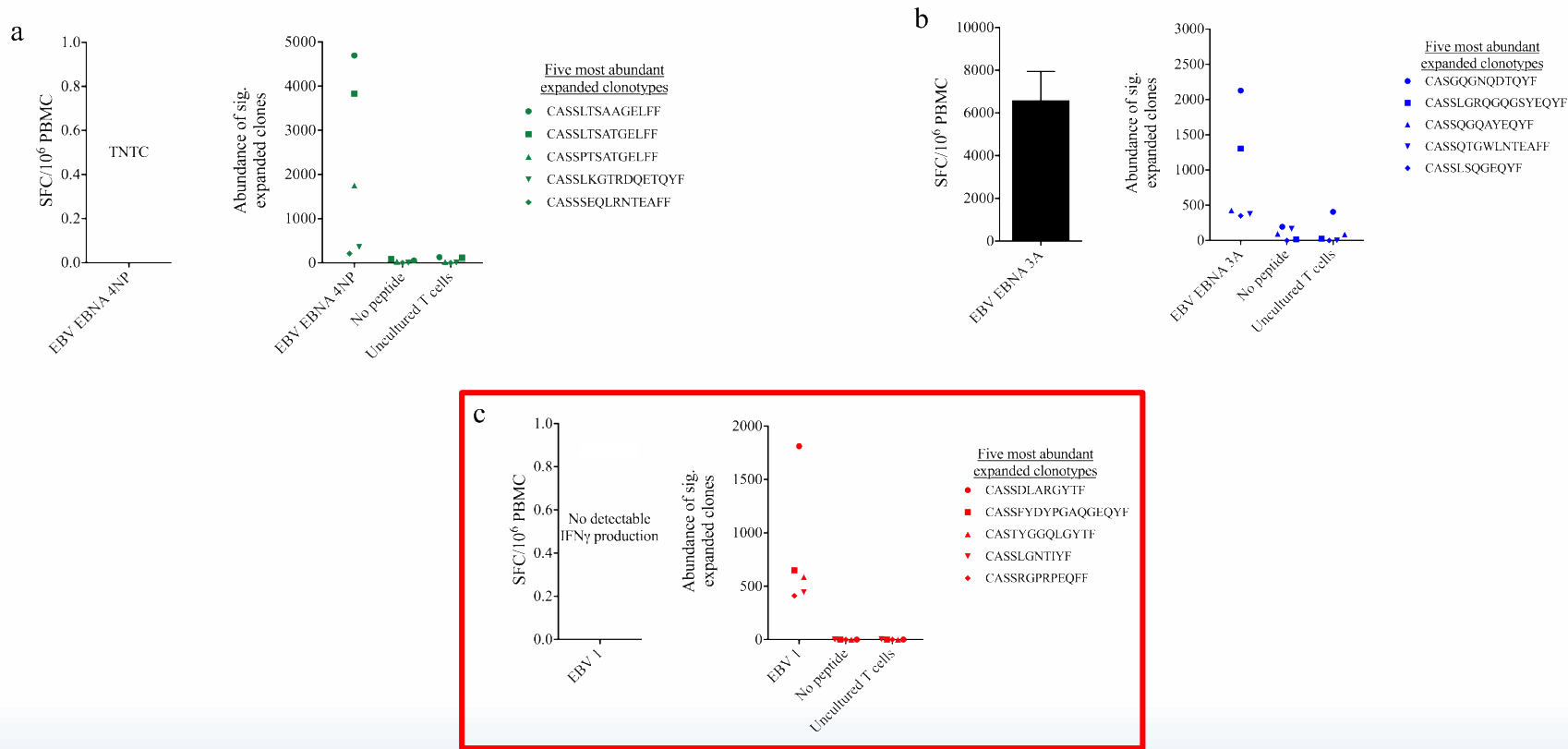
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ELISpot May Underestimate the Anti-Tumor Immune Response



ELISpot May Underestimate the Anti-Tumor Immune Response

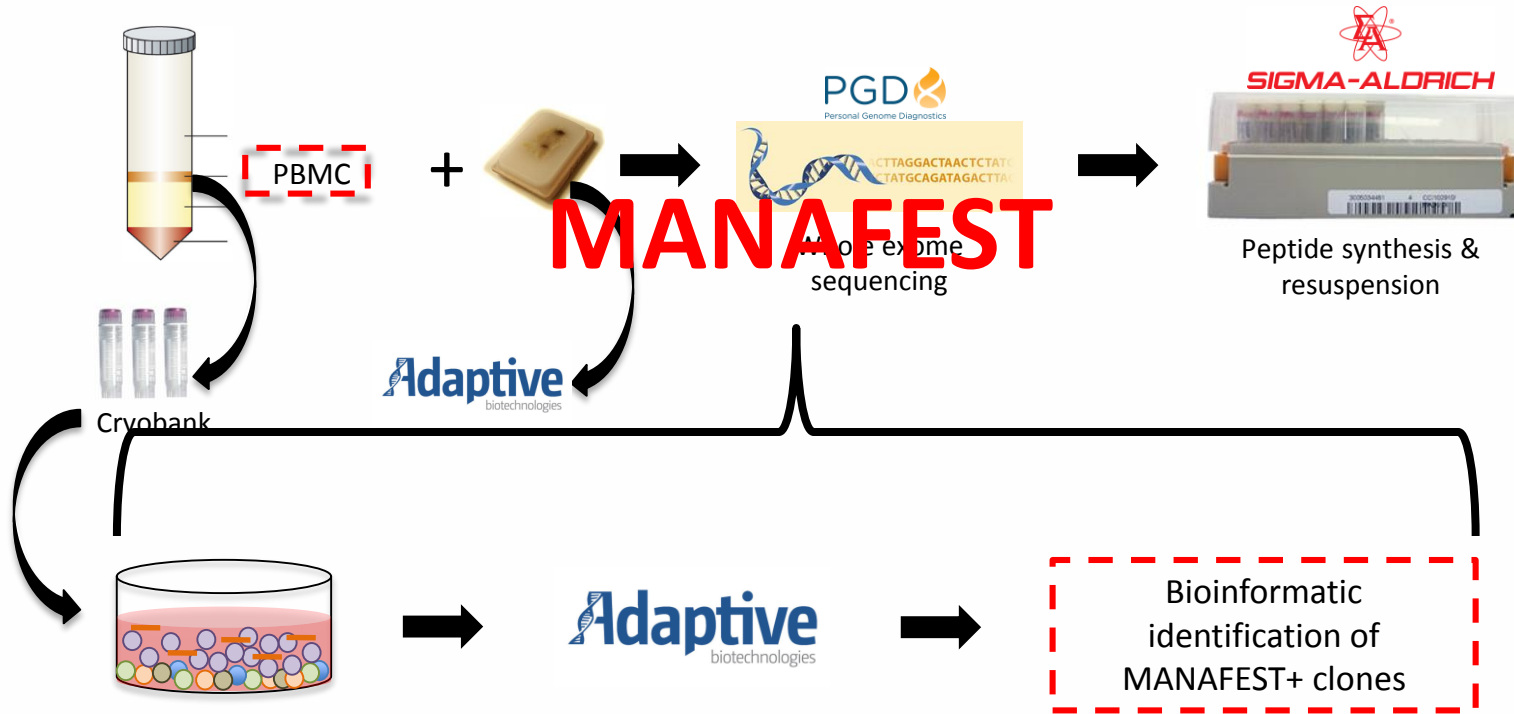


The Functional Expansion of Specific T Cells (FEST) Assays

- Combines experimental T cell culture with in silico identification of clones corresponding to Ag-specific T cells
 - ViraFEST
 - TAAFEST
 - BactiFEST
 - OncoFEST
 - AutoFEST
 - **MANAFEST**



Basic Workflow of the MANAFEST Assay



Data Analysis & Bioinformatic Pipeline

FEST data analysis

User's manual

Upload FEST files (.tsv)

Browse...

No file selected

Save Input Object

OR

Upload an R Object with data (.rda)

Browse...

No file selected

Select a reference sample

None

Select a baseline sample

None

Select samples to exclude

Specify the minimal number of templates (increase in case of large files)

1

☐ Ignore baseline threshold

Specify clone confidence

0.99

Estimated number of CD8+ T cells per well

100000

☐ Use nucleotide level data

Run Analysis

Specify FDR threshold

0.05

Specify odds ratio threshold

5

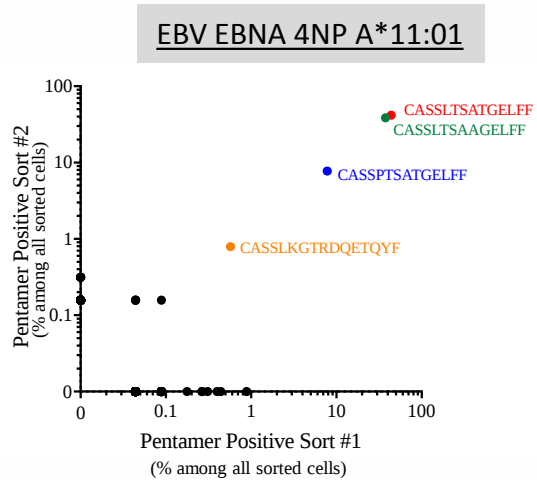
Download Results

Create heatmaps

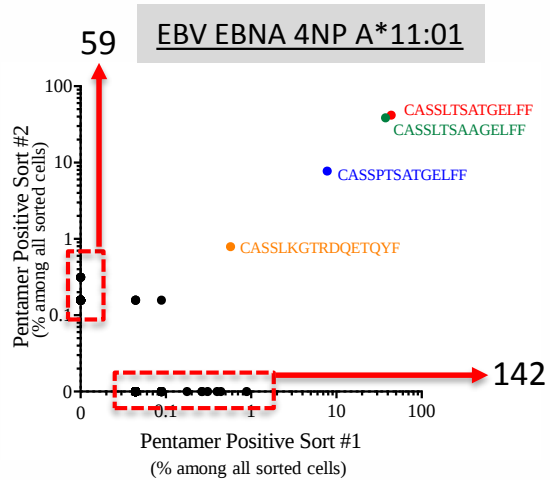
- What is considered “MANAFEST+”?
 - Significant expansion compared to “no peptide”
 - Significant expansion compared to every other peptide
 - Odds ratio >5
 - No significant expansion in response to any other peptide
 - ~46 peptides for optimum specificity
- Match w/ clones found in tumor
- Evaluate peripheral dynamics of MANAFEST+ clones



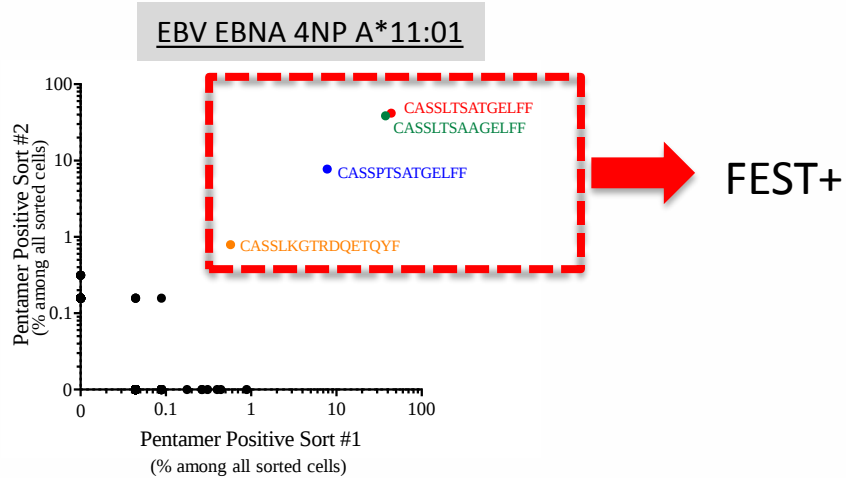
Sensitivity & Specificity



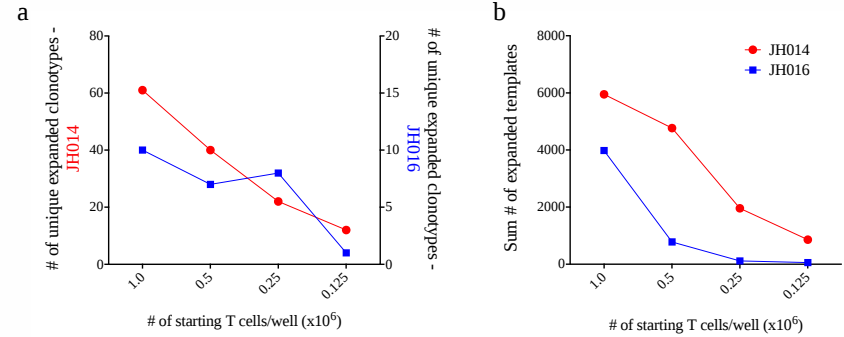
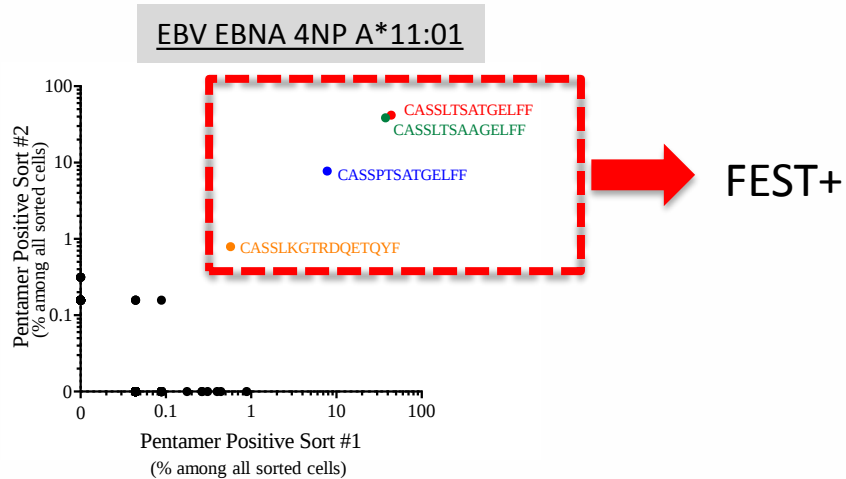
Sensitivity & Specificity



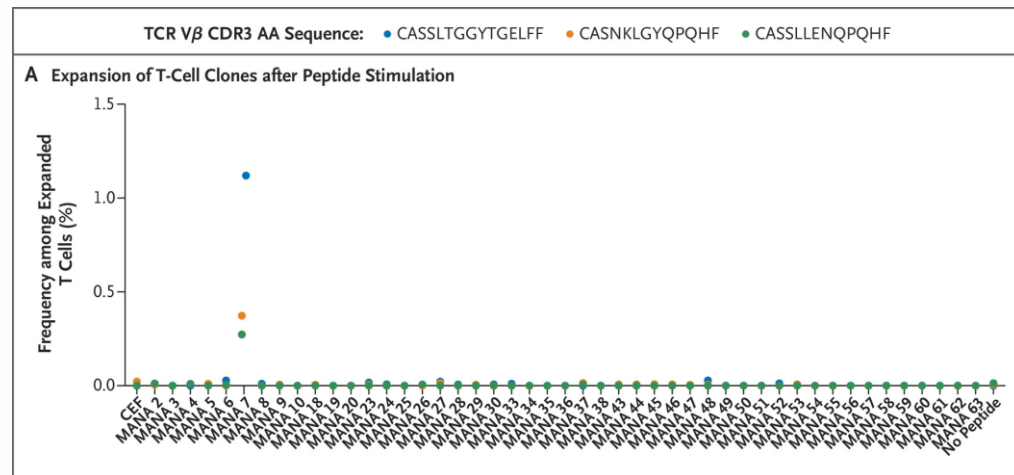
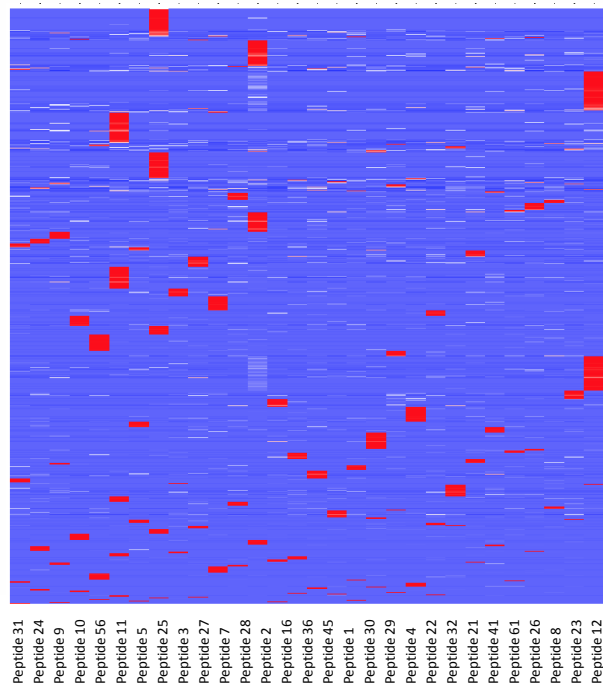
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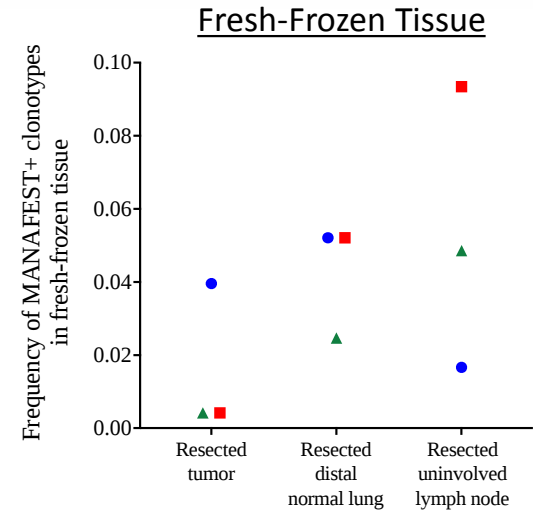
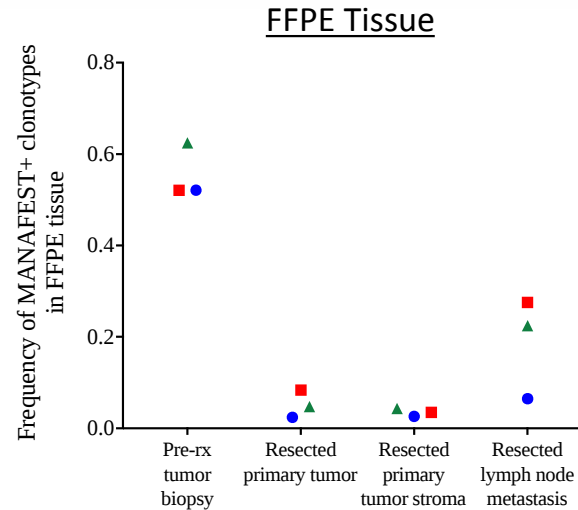
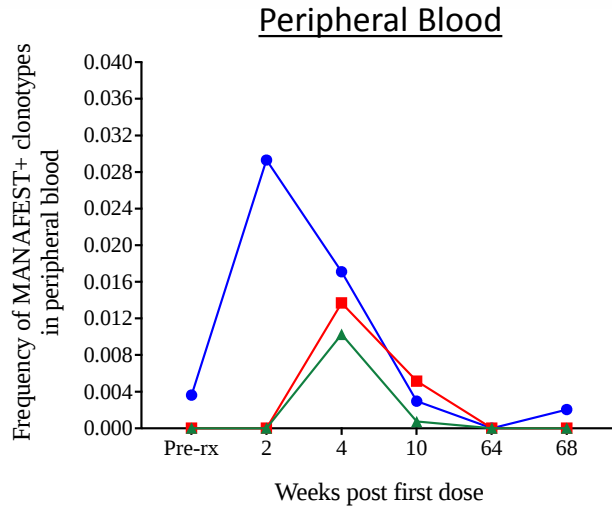
Example MANAFEST Data Visualization



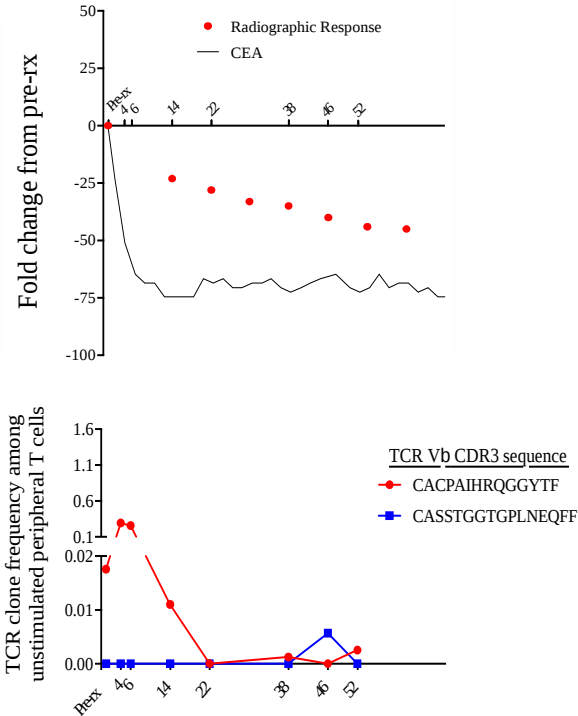
Forde, Chaft, and Smith et al, NEJM, 2018



Possibilities: tracking antigen-specific T cells in every biological compartment & specimen type



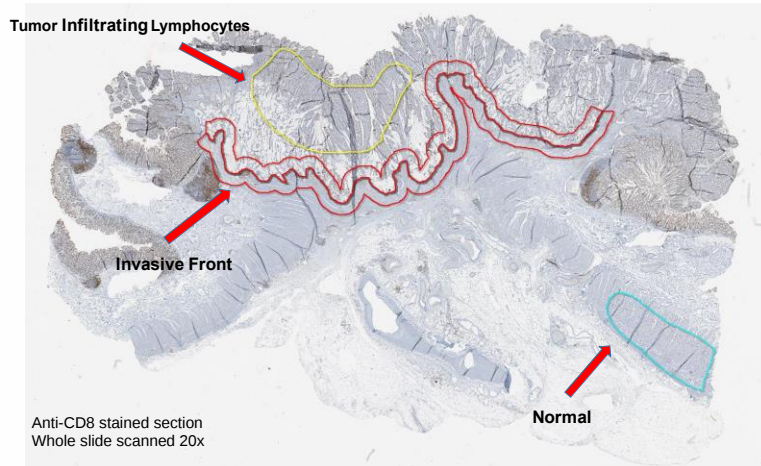
Possibilities: associating antigen-specific T cell dynamics with clinical parameters



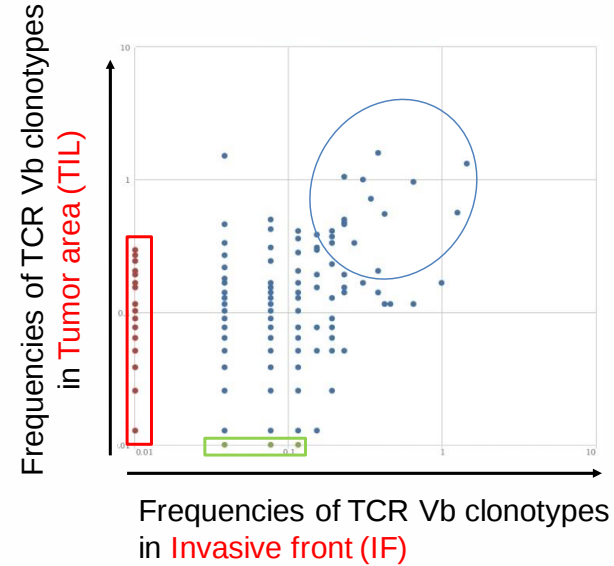
Adapted from Le DT, Science, 2017



Possibilities: tracking geographic distribution of antigen-specific T cell repertoire



Nicolas Llosa, MD
Franck Housseau, PhD



Quantitative outputs: a wealth of data

- # of positive peptides
- Frequency of each positive clone
- Combined frequency of positive clones
- Expansion compared to baseline
- Common motifs
- Peripheral and intratumoral dynamics



How does MANAFEST stack up?

	ELISpot/FLUOROSpot	MULTIMERS	F.E.S.T.
Assay	Enzymatic/fluorescence	Flow cytometry	NGS
T cell activation	In vitro	Ex vivo	In vitro
Signals	Colorimetric/fluorescence	Fluorescence	TCR sequences
Biospecimens	Fresh PBMC Antibodies Peptides	Fresh PBMC Antibodies Peptides HLA monomers	Fresh PBMC
Instrumentation	ELISPOT reader	Flow cytometer	None
Quantification	Frequency/intensity	Frequency	# clonotypes Frequency Fold change
Sensitivity	1/10,000-20,000	1/5,000-10,000	>1/100,000
Specificity	++ Background	++ Background Spectrum overlaps	+++
Resolution	>10-50 cells/well	>250 events/gate	>10 reads/clonotype
Throughput	Low	High	High
Functional readout	Cytokine / Granzyme	Multiparameter	N/A
Time course	1 ELISPOT/time point	1 staining/time point	1 test total
Archival specimens	No	No	FFPE/snap frozen
Geographic analysis	No	No	Yes, using FFPE

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Future Directions

- Opportunities for machine learning approaches
- Phenotyping neoantigen-specific clonotypes
- Association with clinical parameters in a variety of disease and treatment settings
- Assay development and optimization: cross-institutional validation with MSKCC



Acknowledgements

Cancer Immunology

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PATIENTS AND THEIR FAMILIES!



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