

Novel biomarkers: pitfalls, limitations, emerging options.

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BIOMARKERS

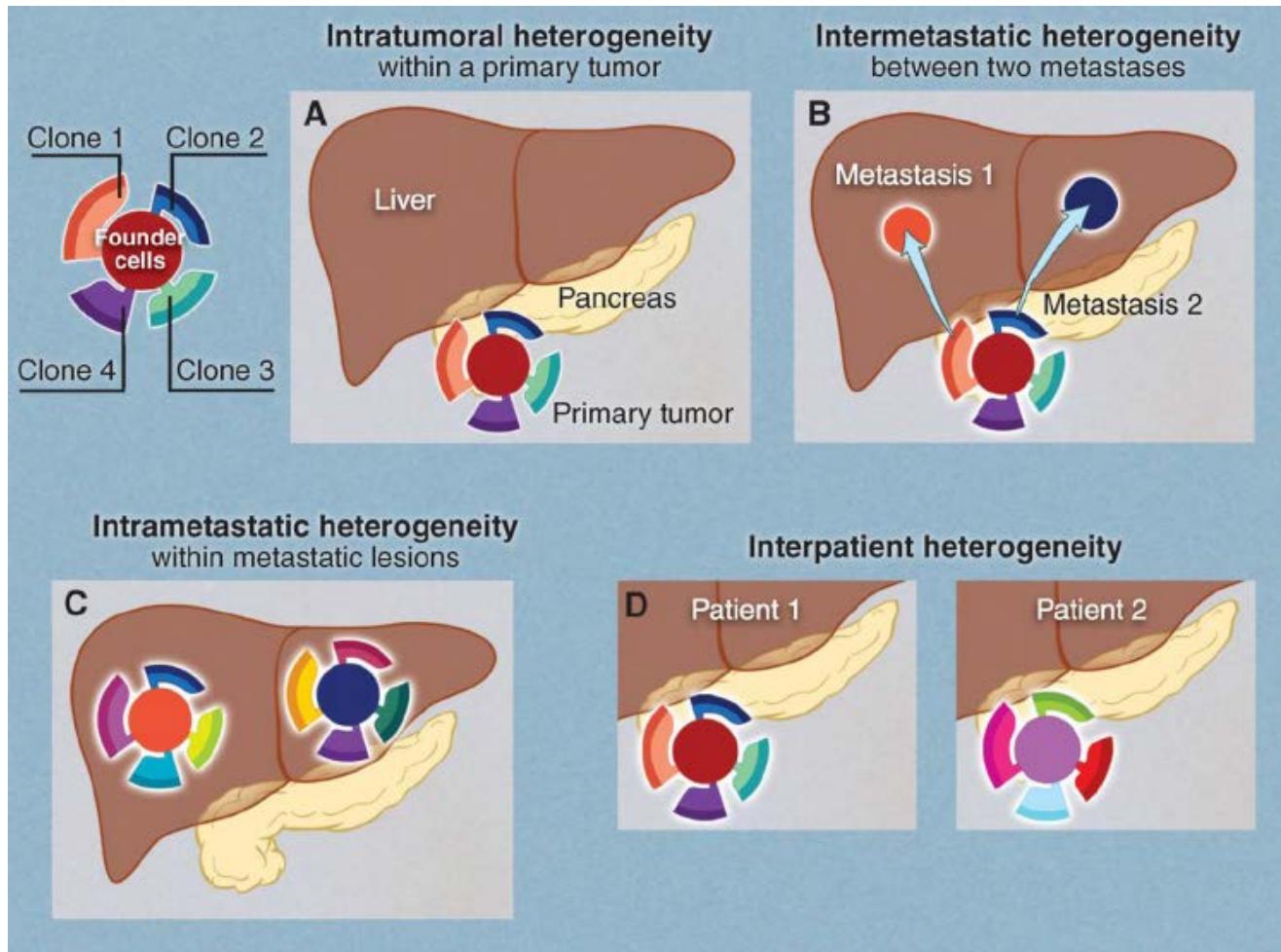
- Tumor:
 - Histol
 - Molec
 - Single IHC/ISH
 - Multiplex
 - DNA mutations
 - RNASeq/NanoString
- Blood:
 - PBMC
 - Sera
 - Phenotyping
 - Functional
 - Ab Response (pre)
 - Mass Spec
- “Systemic”
 - Imaging
 - Microbiome



“Hurdles”

- Tumor Heterogeneity (antigenic)
- Checkpoint blockade and Costimulatory Abs only effective against immunogenic tumor (preclinical)

Cancer Heterogeneity Mandates Broad Immunity



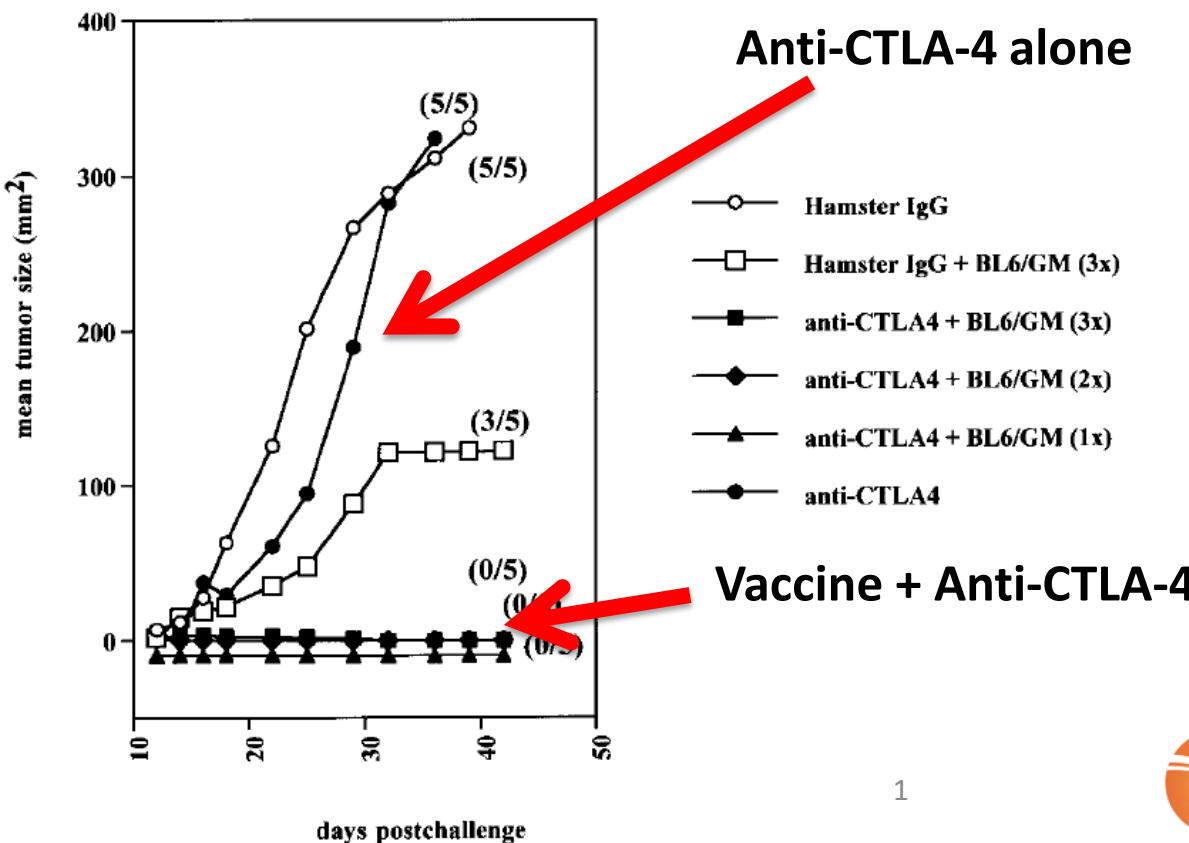
Hypothesis:

Effective treatment of metastatic cancer will require an immune response to many antigens

Many Cancers not seen by immune system

Combination Immunotherapy of B16 Melanoma Using Anti-Cytotoxic T Lymphocyte-associated Antigen 4 (CTLA-4) and Granulocyte/Macrophage Colony-Stimulating Factor (GM-CSF)-producing Vaccines Induces Rejection of Subcutaneous and Metastatic Tumors Accompanied by Autoimmune Depigmentation

By Andrea van Elsas, Arthur A. Hurwitz, and James P. Allison



J. Exp. Med 190 (3) : 355-366, 1999

Hypothesis:

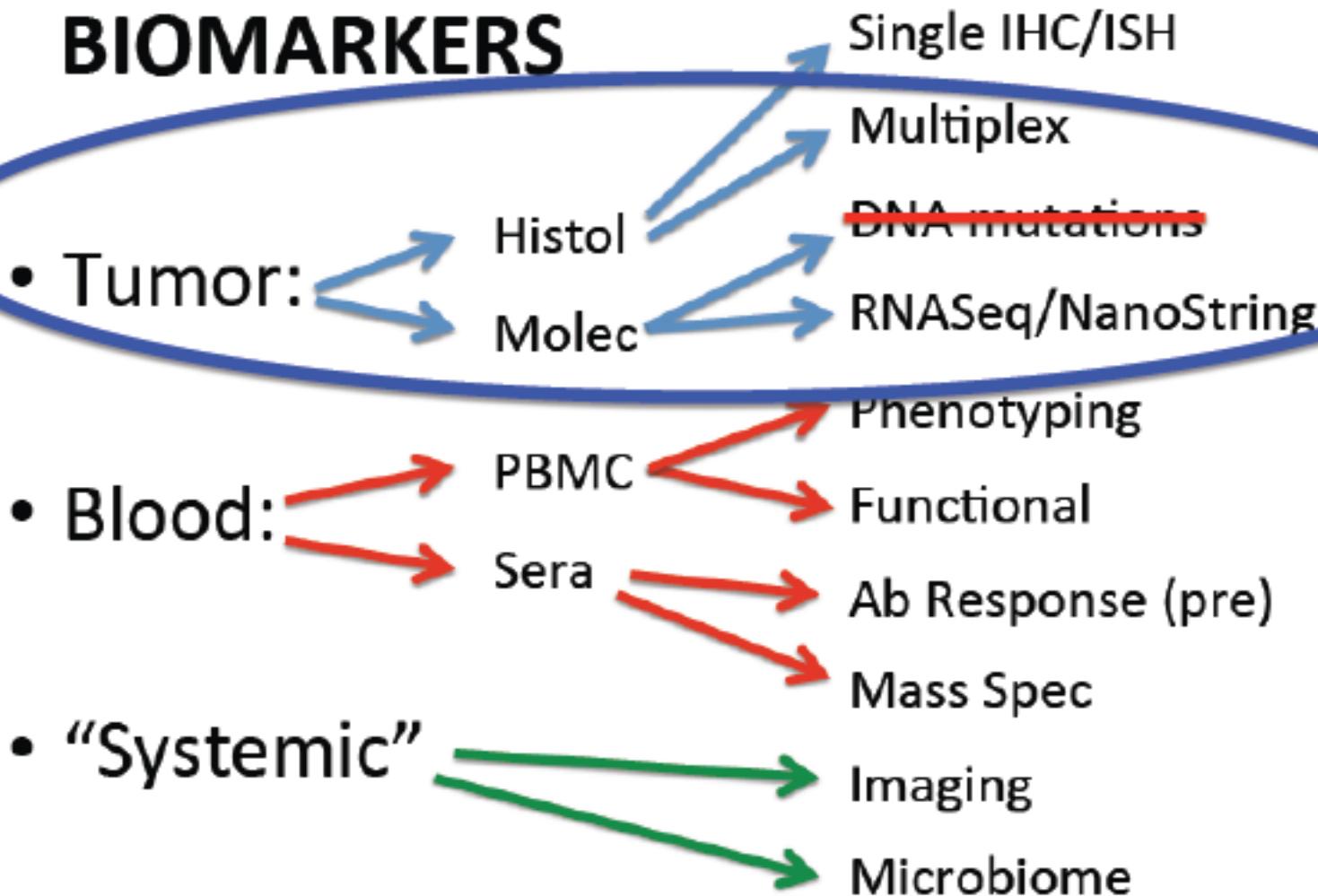
Tumors that are less immunogenic need something to prime anti-cancer immunity.

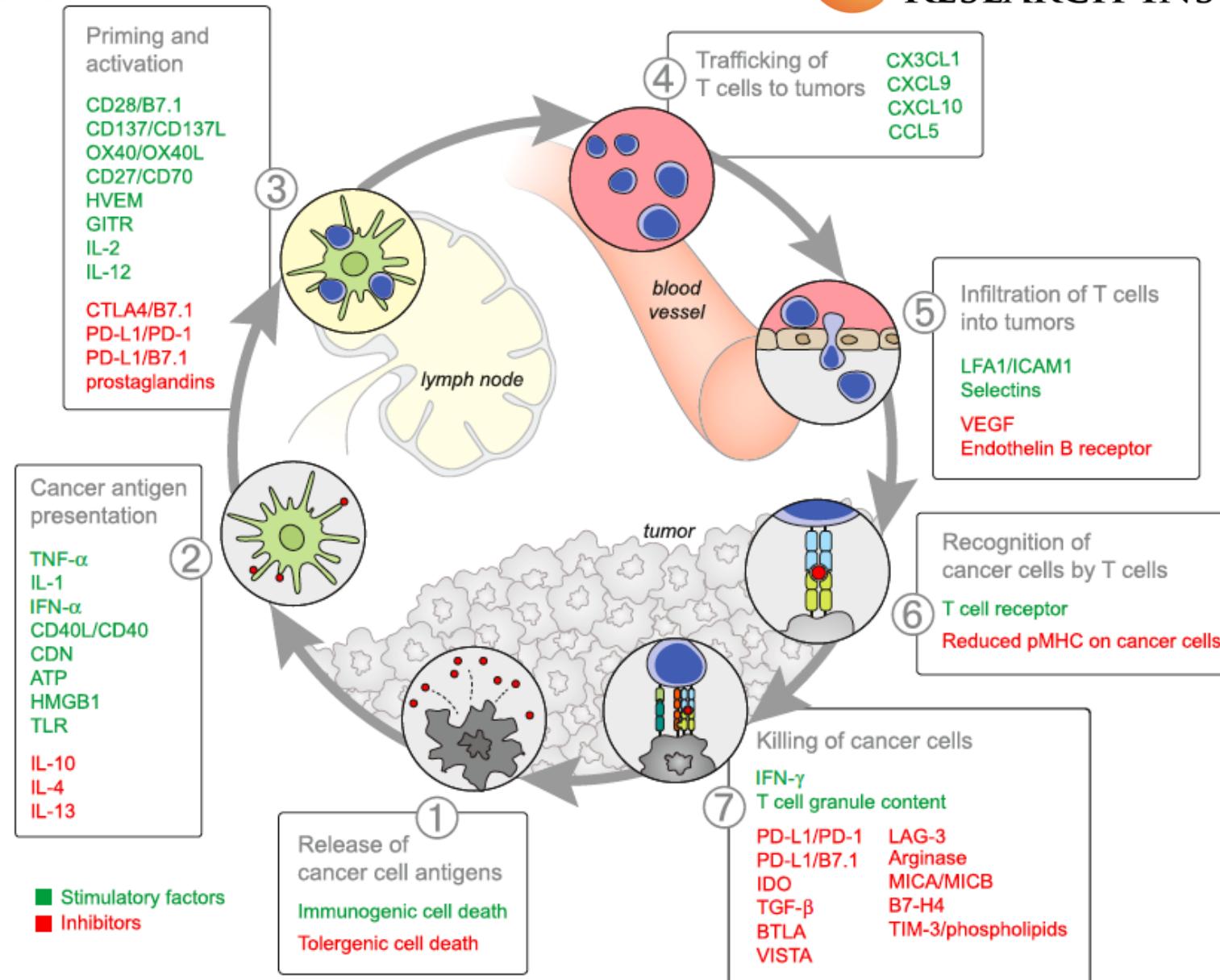
- **Vaccines**
 - Chemo/Rad
 - Antibodies
 - BiTEs / DARTs
 - Oncolytic viruses

Cancer Immunogenicity and heterogeneity important

- Underscores limitations of sampling
- Gives direction for “Future” – More effective treatments with potential to CURE.

BIOMARKERS







Zipei Feng

red = PD-L1
yellow = CD8
green = Foxp3
purple = CD20
aqua = melanoma cocktail
pink = CD163
blue = DAPI

Biomarkers Not Predictive of tumor-specific TIL cells in the tumor



NOT PREDICTIVE

Number of CD3

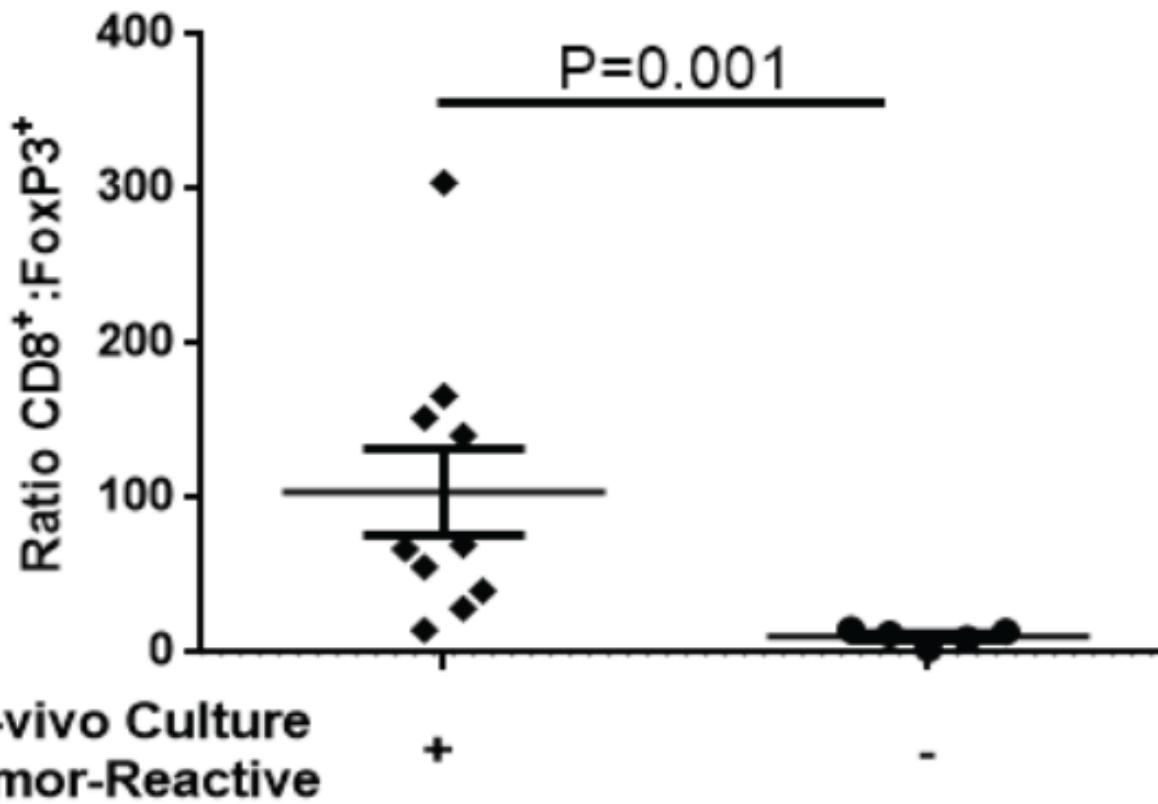
Number of CD8

Number of FoxP3

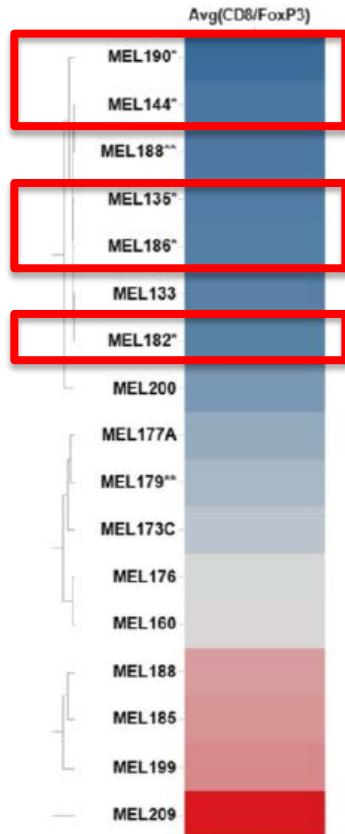
Feng et al. *Journal for ImmunoTherapy of Cancer* (2015) 3:4
DOI 10.1186/s40425-015-0091-z



The CD8:FoxP3 ratio is predictive

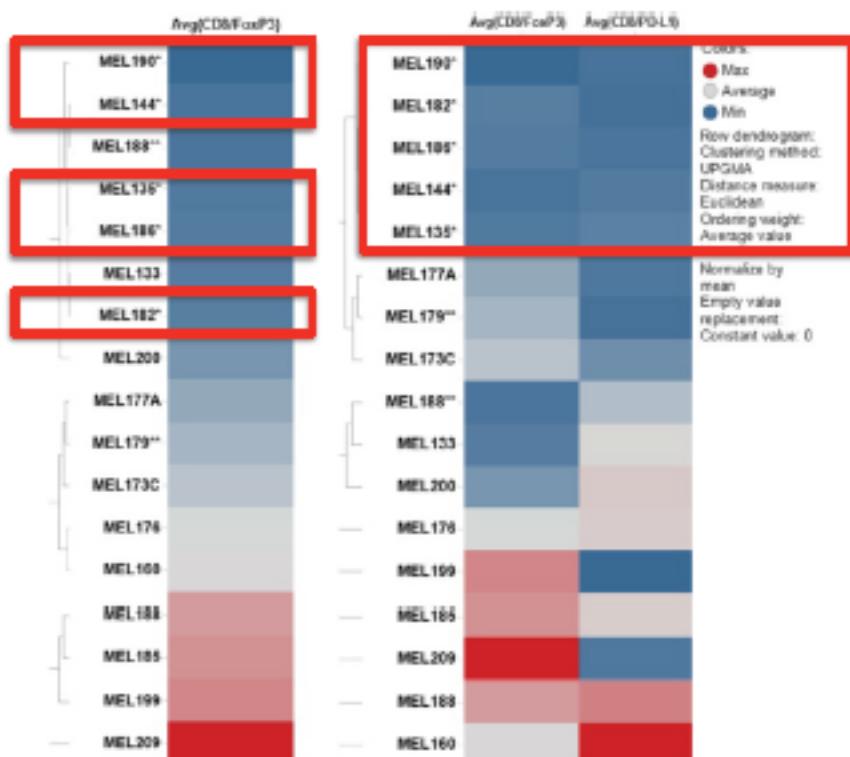


Combining CD8:FoxP3 Increases Power to Predict TIL



Red box Identifies tumors that fail to grow TIL

Combining Both CD8:FoxP3 and CD8:PD-L1 Increases Predictive Power - Melanoma



Red box Identifies tumors that fail to grow TIL

**Collaboration with Cliff Hoyt (PE)*

**Applying to NSCLC and HNSCC*



How to think about relationships?



How to think about relationships?

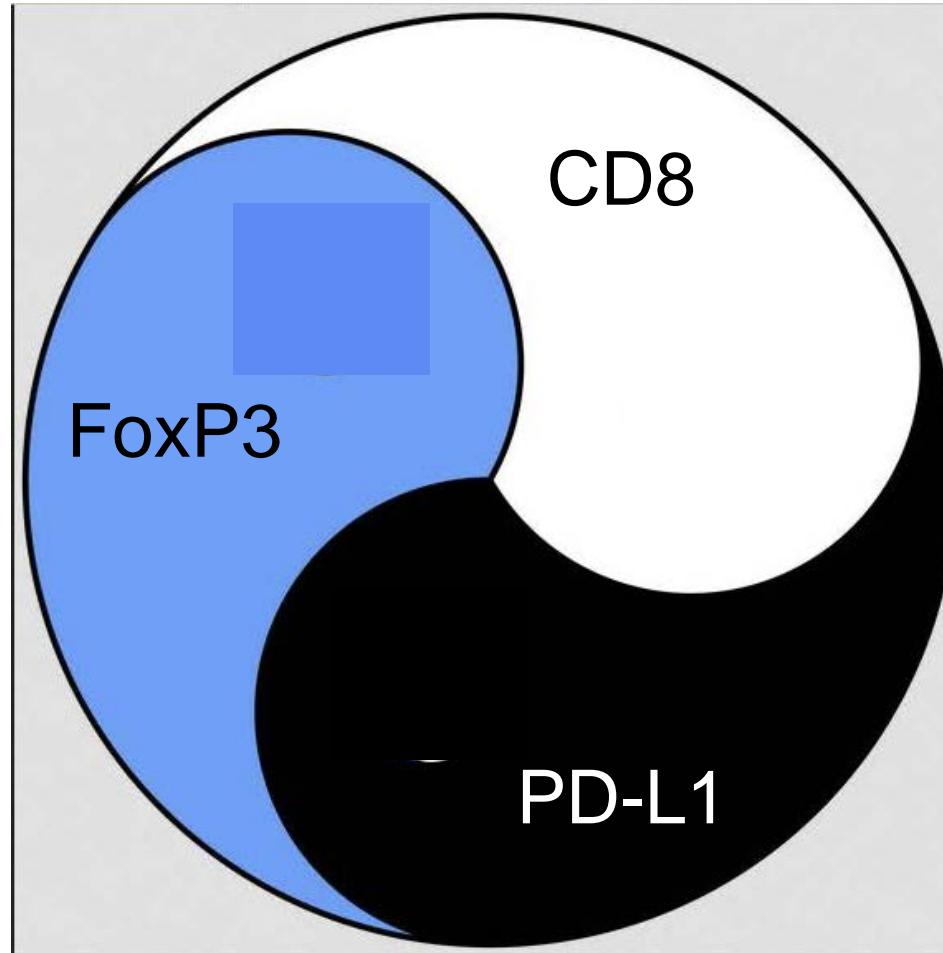


Effectors

Suppressors

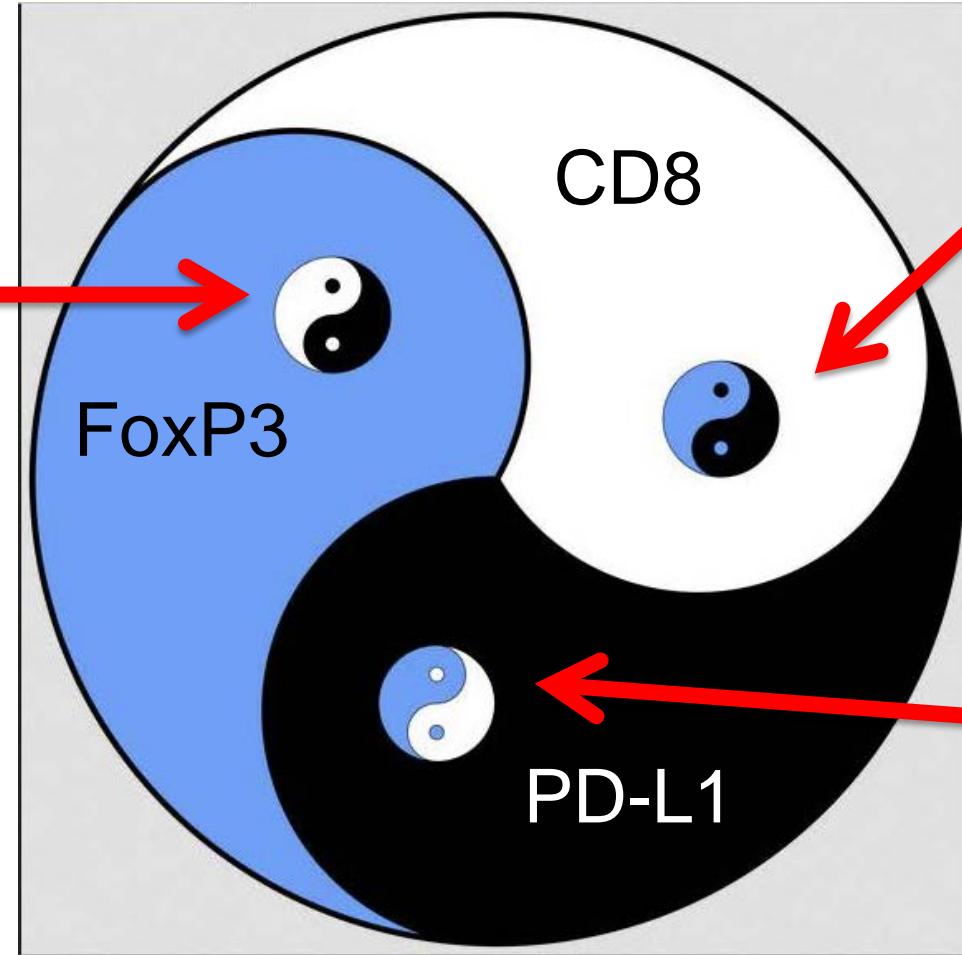


Relationships are complex



Relationships are complex

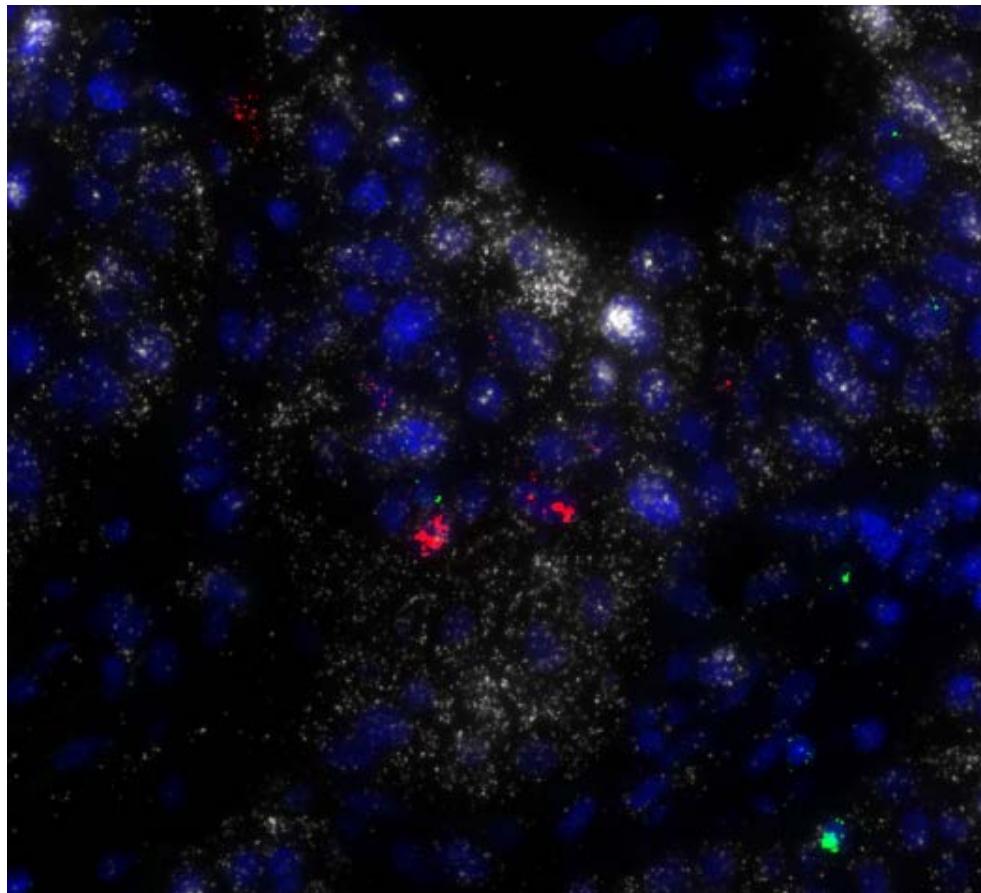
Ratio to CD4s /
Tu – induced
vs
Natural
Where are they
located?
Perimeter /
Center



TCR
Specificity /
Maturation
status
CM, EM, Eff
IFN, TNF, ...

Where
expressed?
What other
inhibitors?

Use ISH to Identify Functional Properties – Collaboration with ACD Bio



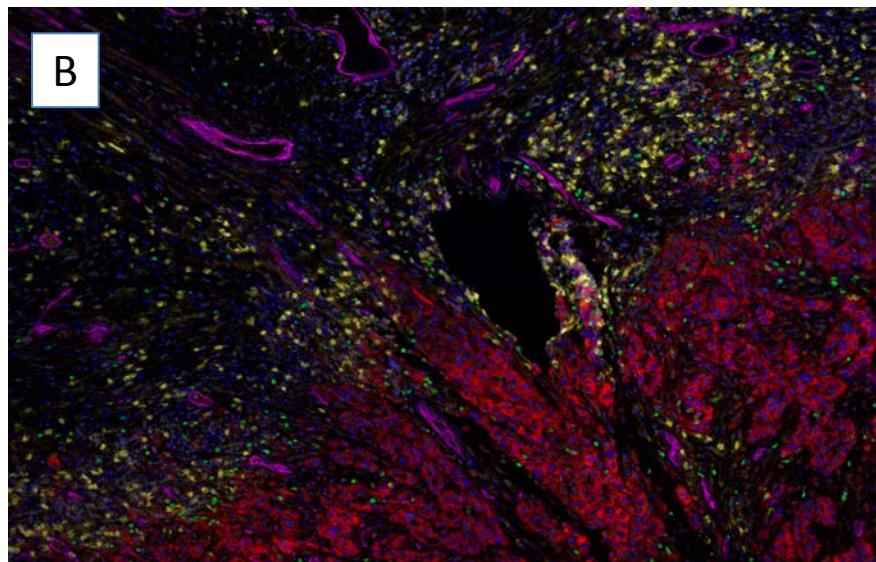
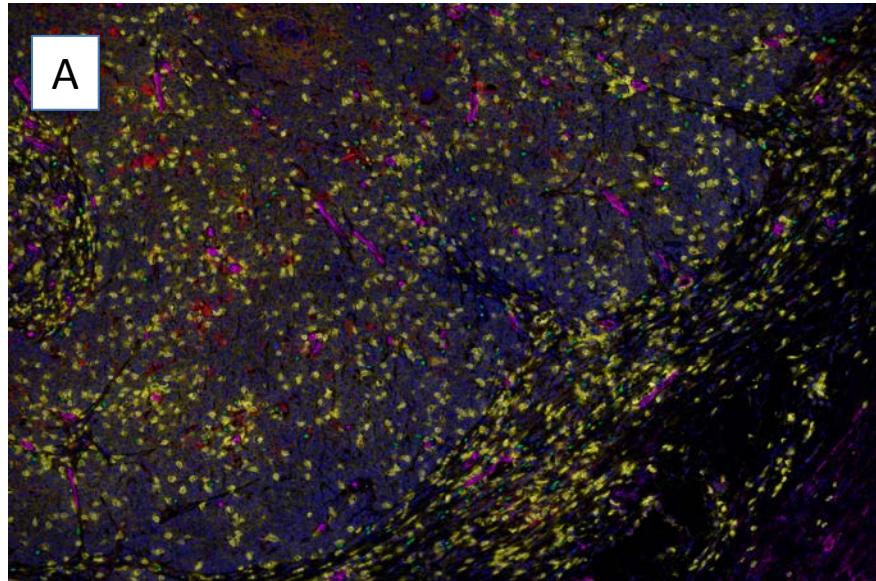
White – TGF β

Red – TNF α

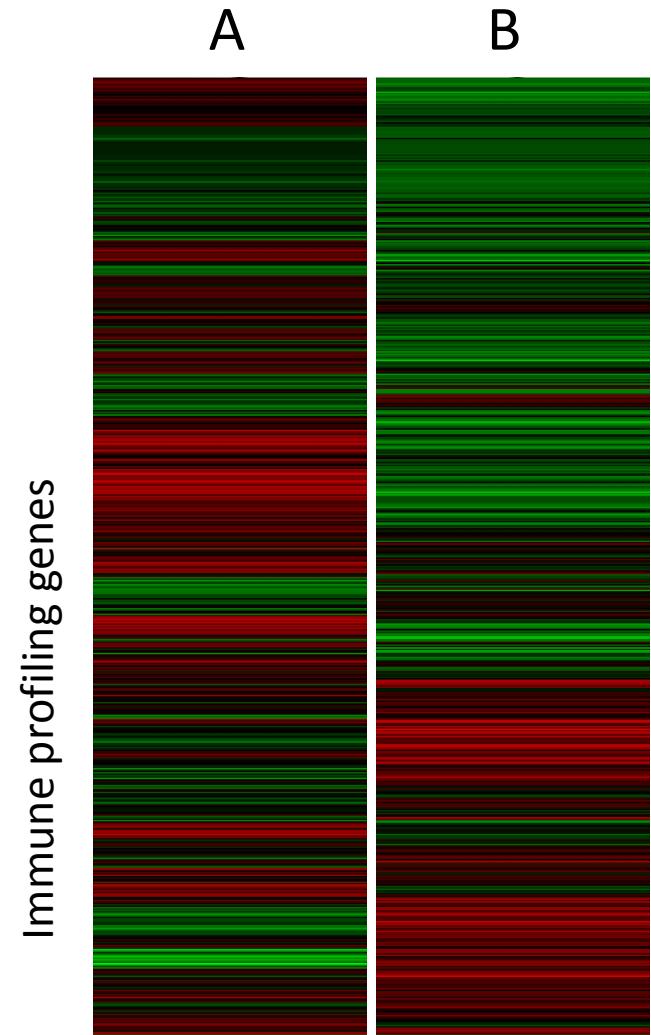
Green – IFN γ

**Collaboration with
Emily Park &
Xiao-Jun Ma
(ACD Bio)*

NanoString Gene Expression Immune Profiling Analysis Performed on Two OHNSCC



PD-L1
FoxP3
CD-3
DAPI





***Possible Today:
Use Multispectral***

- ***Assess tumor biopsies***
 - *T cell infiltrates*

Could Stratify patients

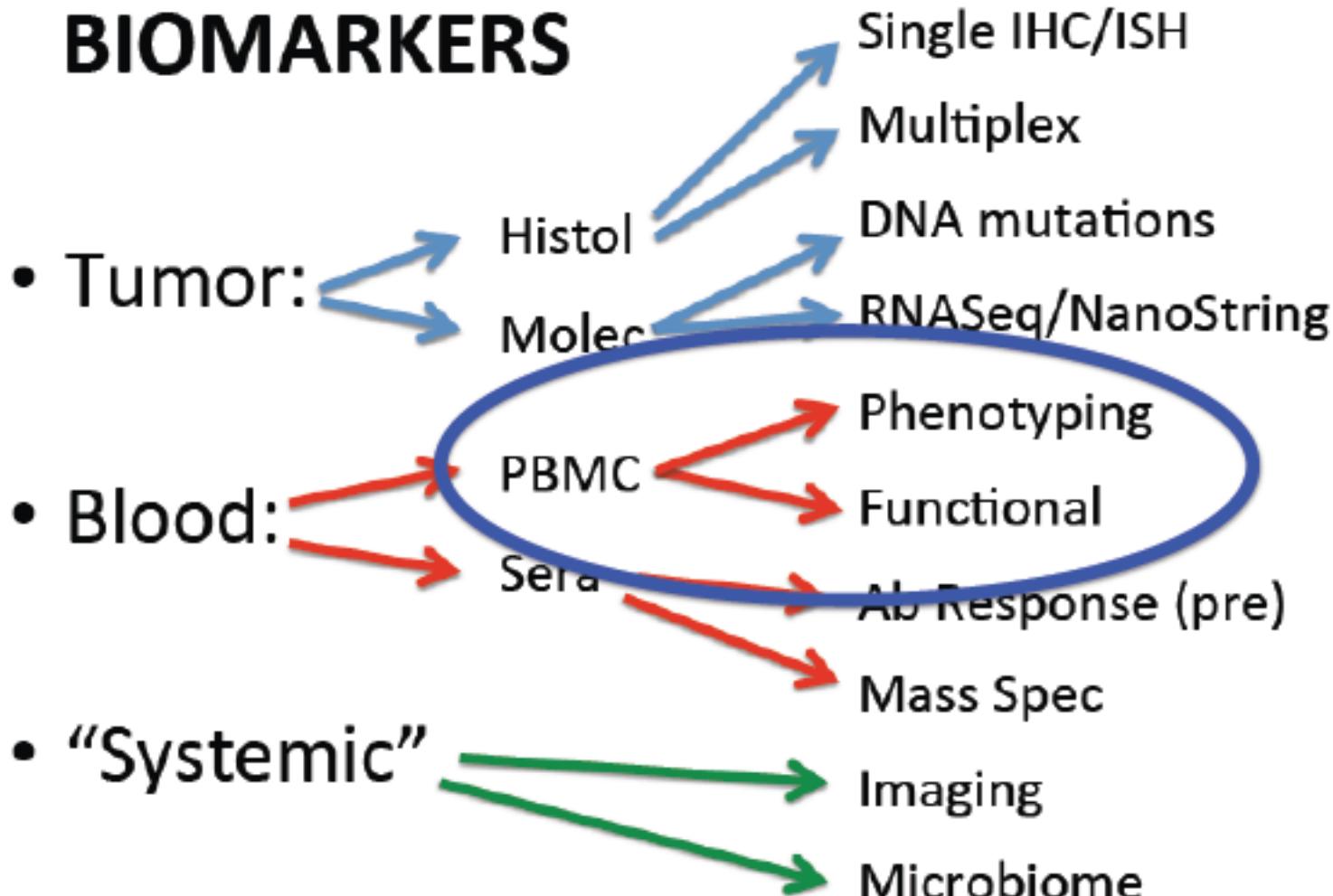
***Currently some CLIA
platforms – LDT (PE)***

Not to distant FUTURE

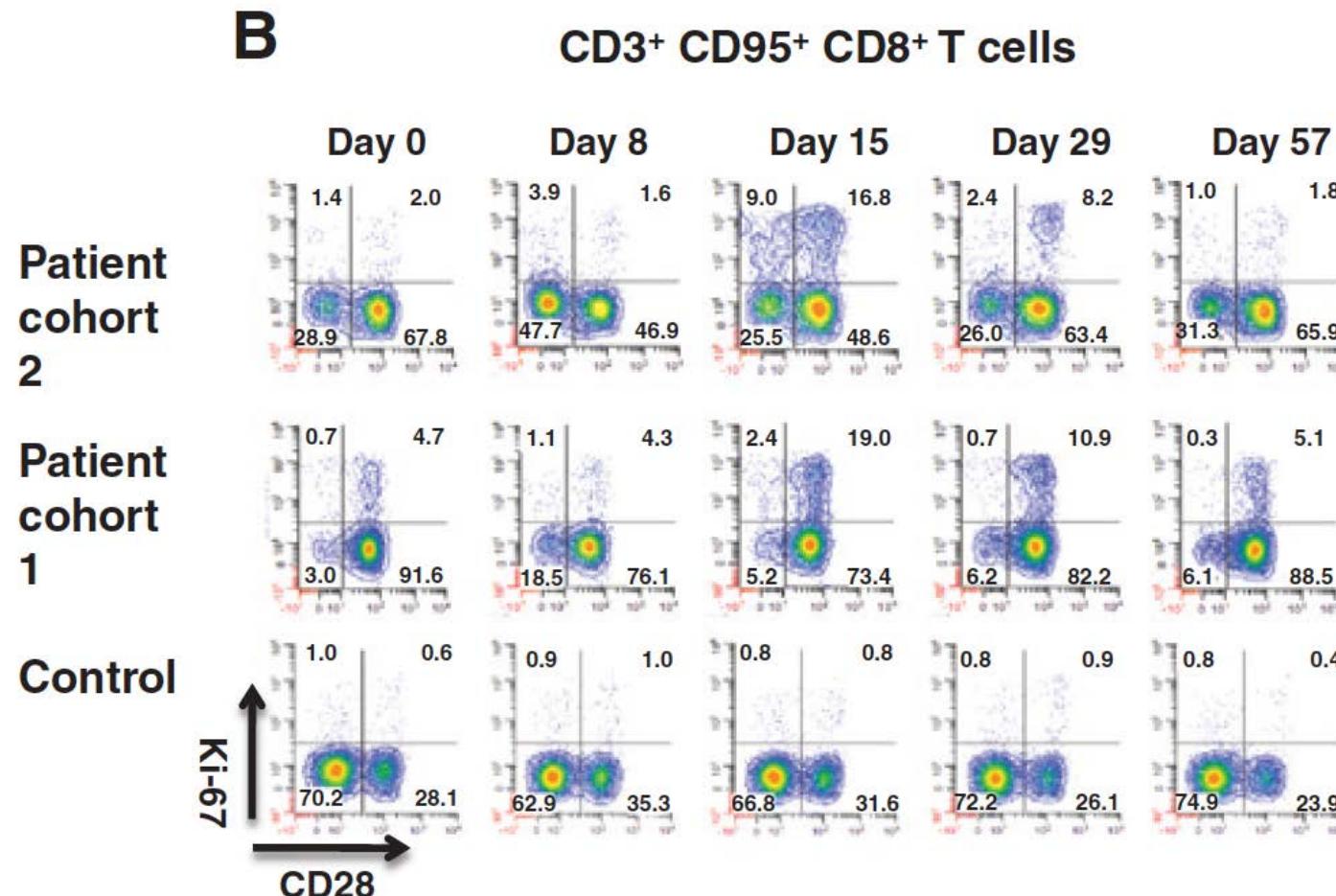
- ***10-25? markers***
***(PD-L1, TIM3,
VISTA, other)***

Tailor Therapy

BIOMARKERS



Blood – Phenotypical changes to treatment: Anti-OX40 administration induces qualitative changes in cycling CD8 T cells



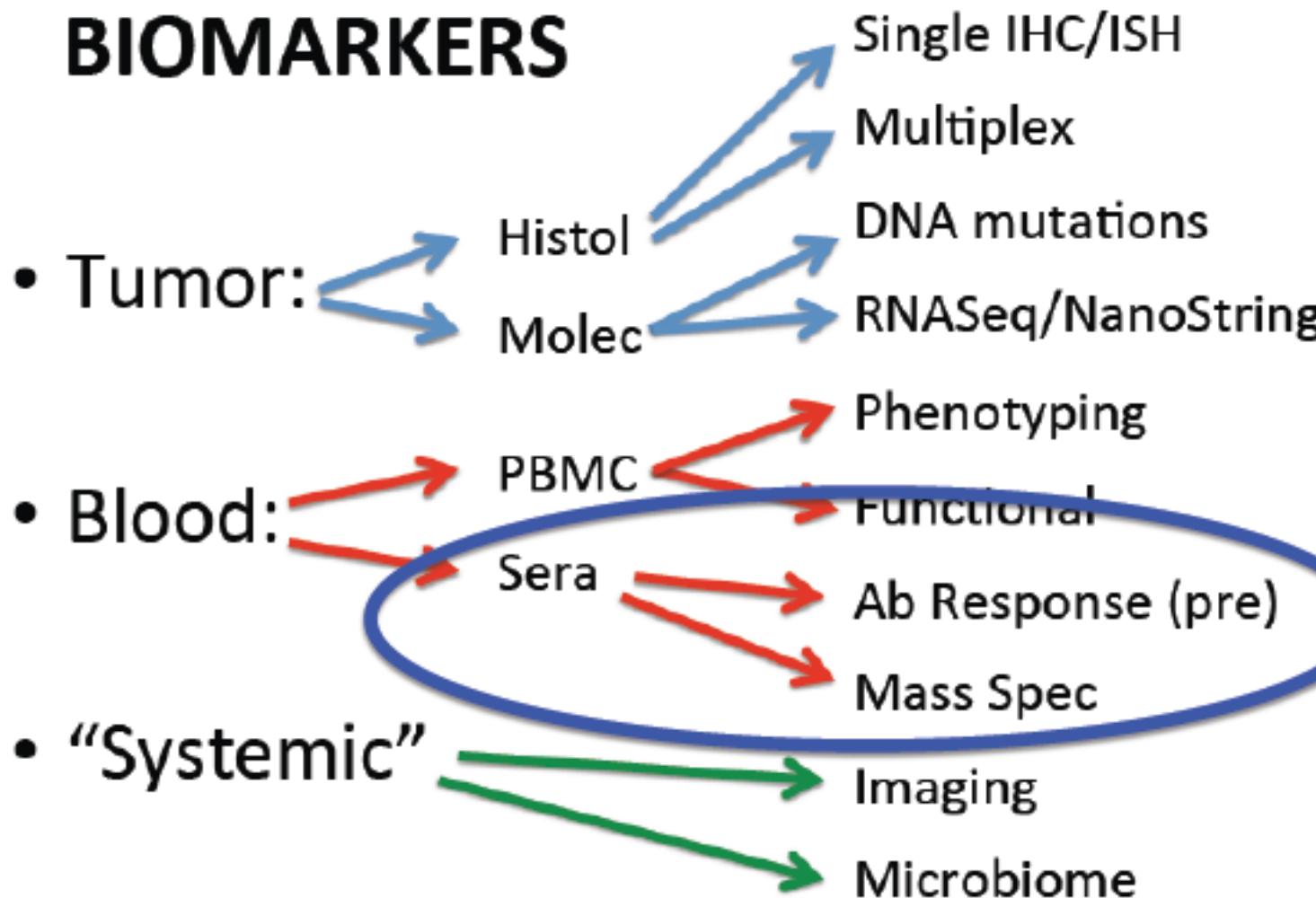
Blood – Functional : Pre-existing or Treatment induced Immunity

Idea: Immunoscore in the “Blood”

Method: Use DC-Targeted microvesicles containing viral ag or >100 over-expressed cancer proteins / NCI-Prioritized Ags / CRA

- **Demonstrated to viral antigen (CMV)**
Ye, W. J. Transl. Med 12:100, 2014
- **Evidence also in Prostate Cancer**
van de Ven, R., Manuscript in preparation

BIOMARKERS



Sera: Abs or Other Proteins

Hypotheses:

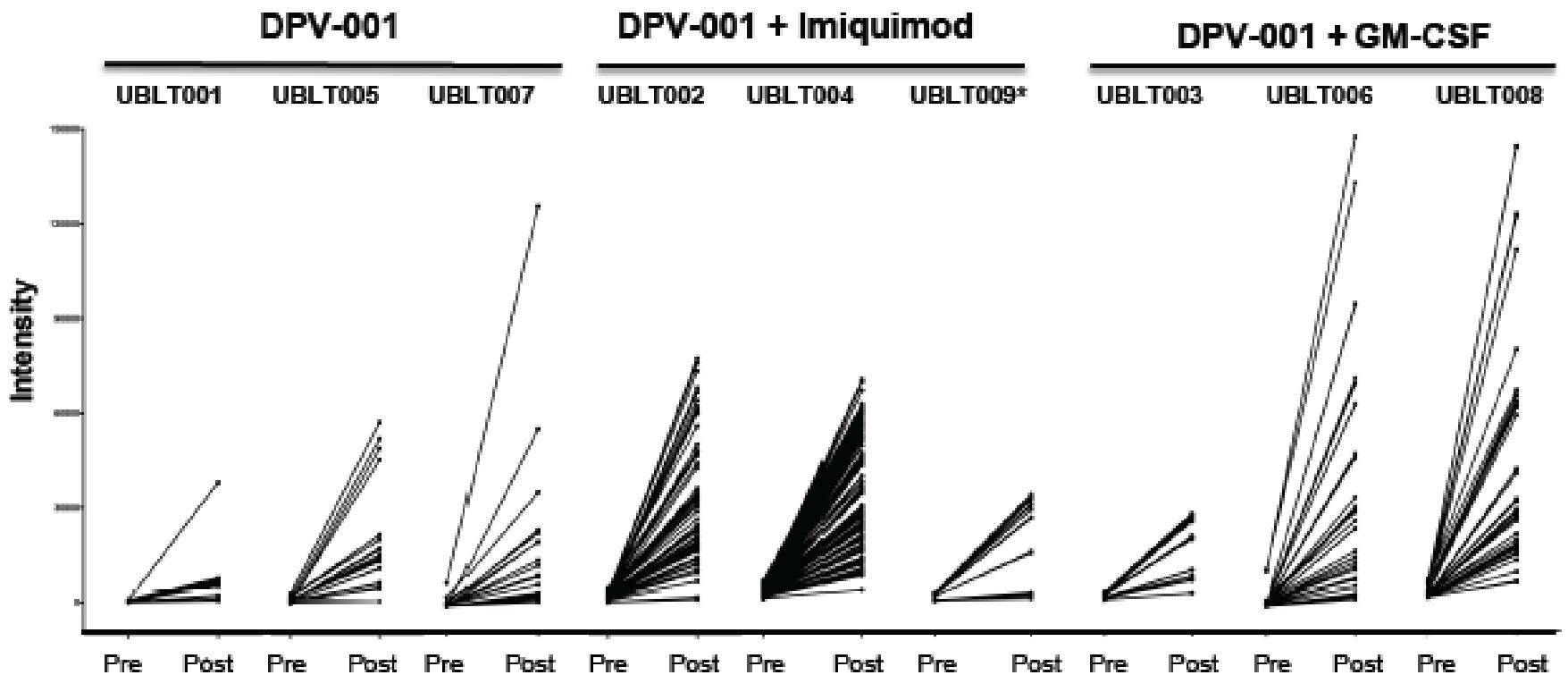
- Abs to “certain” targets ID patients with “therapeutic” immunity
- Inflammatory/other proteins in sera ID patients with ongoing anticancer immunity

Method:

- Protein Arrays
- Mass Spec / Deep Maldi Approach

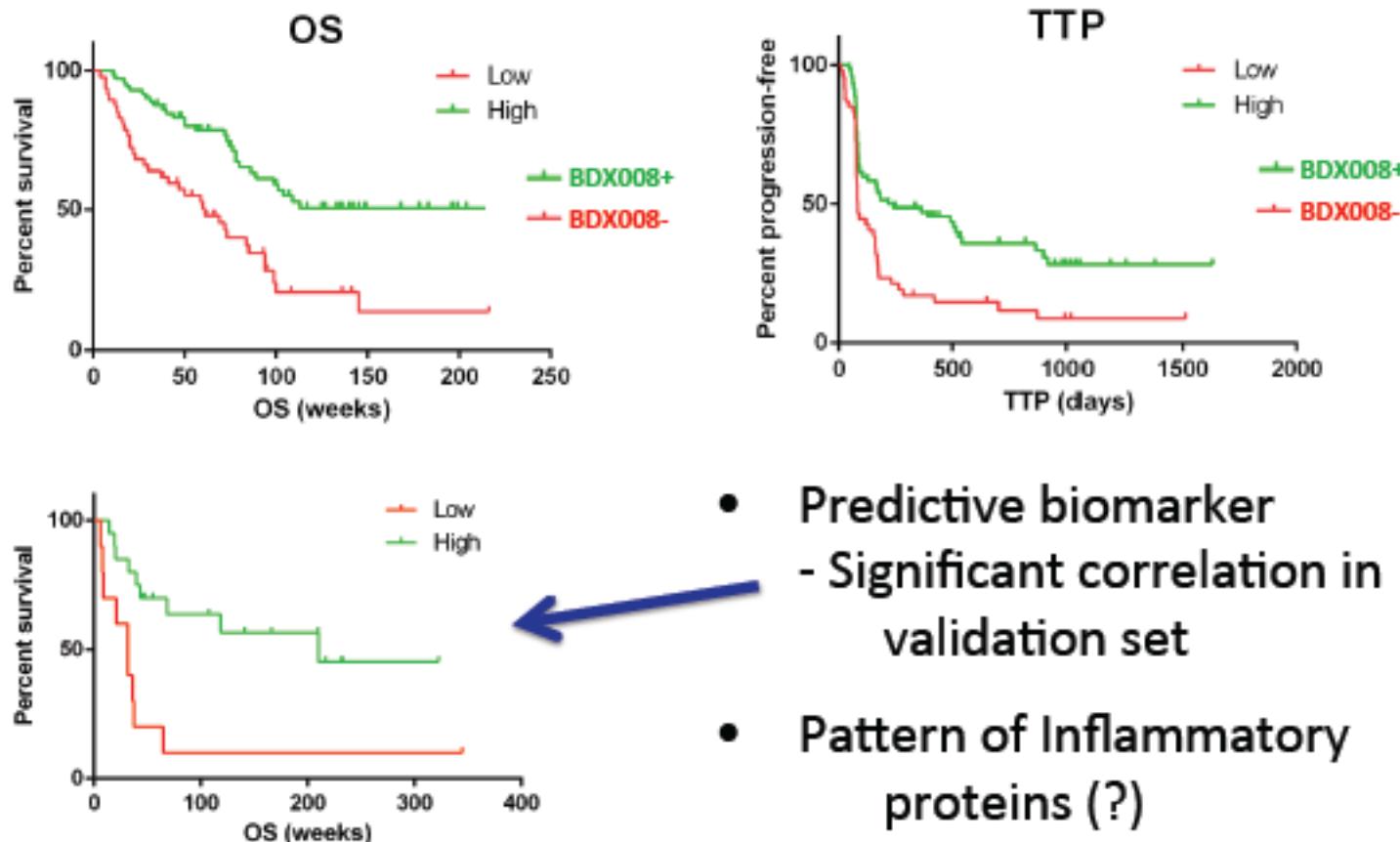
Vaccinated patients make strong immune response (10 fold) to cancer antigens.

- Majority against non-mutated epitopes



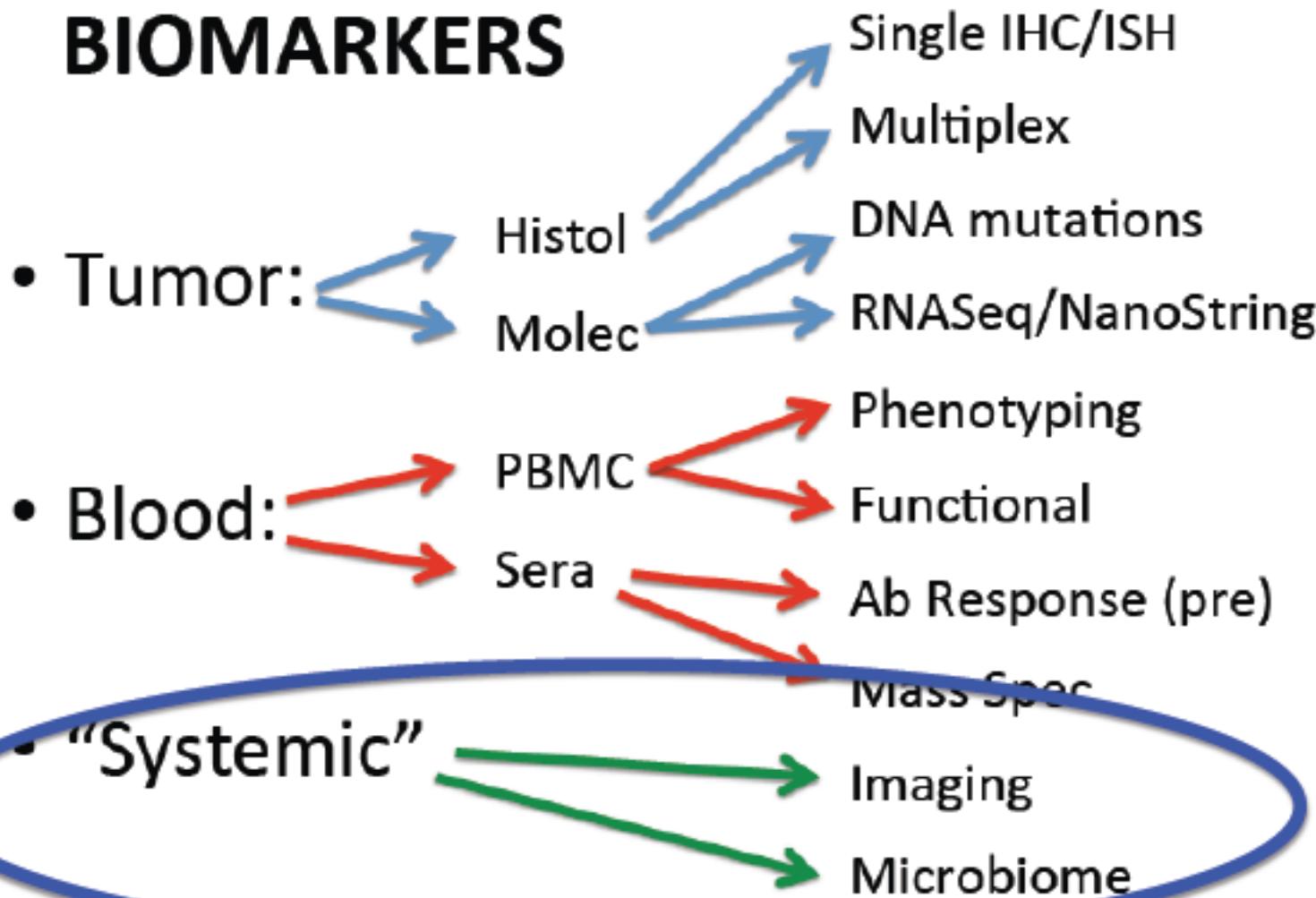
Sanborn, R., Journal for ImmunoTherapy
of Cancer 2015, 3(Suppl 2):P435

Sera: Mass Spec / Deep Maldi Approach Melanoma – Anti-PD-1 Cohort



Weber, J.S. Journal for ImmunoTherapy of Cancer
2015, 3(Suppl 2):P103 (4 November 2015)

BIOMARKERS



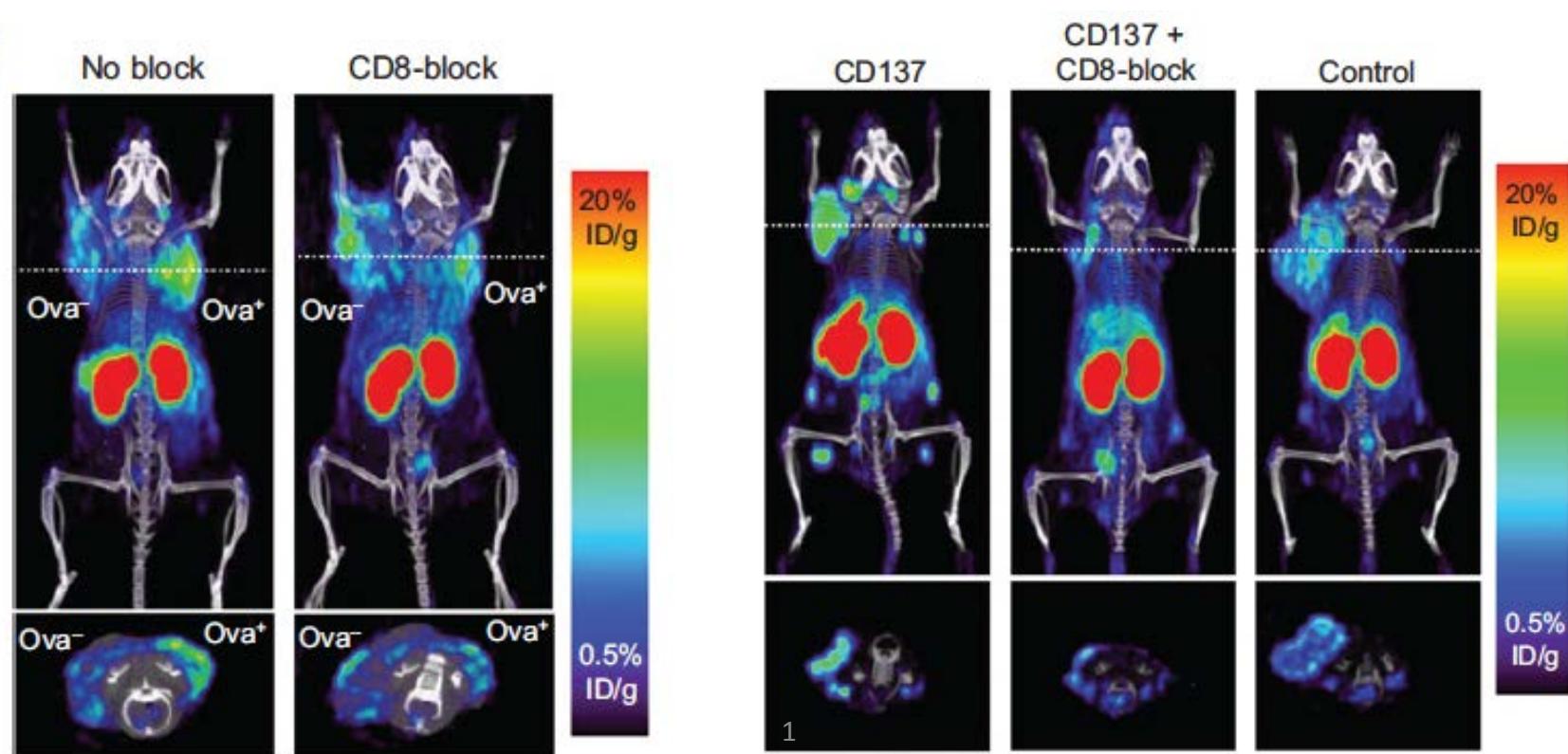
Future:



An Effective Immuno-PET Imaging Method to Monitor CD8-Dependent Responses to Immunotherapy

OnlineFirst November 16, 2015

Richard Tavaré^{1,2}, Helena Escuin-Ordinas³, Stephen Mok², Melissa N. McCracken², Kirstin A. Zettlitz^{1,2}, Felix B. Salazar^{1,2}, Owen N. Witte^{2,4,5,6}, Antoni Ribas^{2,3,7,8,9}, and Anna M. Wu^{1,2,7}

B



“Never, ever, think outside the box.”

Must Evaluate the MICROBIOME!!

- Call to archive “relevant” microbiome of all patients enrolled on clinical trials
- Need to develop a TCGA for Microbiome of patients on “Immunotherapy” trials

Earle A. Chiles Res. Inst.

Fox Lab:

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David Messenheimer

Tyler Hulett

Zip Feng

Hong-Ming Hu

Keith Bahjat

Andy Weinberg

Michael Gough

William Redmond

Marka Crittenden

Walter Urba

Carlo Bifulco

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Alison Conlin

David Page

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Bryan Bell

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Chichung Wang

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Nat. Tumor Inst.

Napoli

Paolo Ascierto

Definiens

ACD Bio

Emily Park

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Patients and their families

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Robert Franz