

# Major Questions for Development of Immunotherapies Requiring Animal Models

Workshop on Challenges, Insights, and Future Directions for Mouse and Humanized Models  
in Cancer Immunology and Immunotherapy

SITC

Washington, DC

November 10, 2016

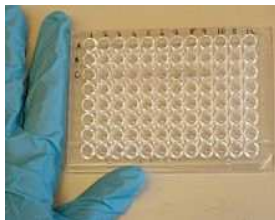
Biological molecule

Role in Anti-Tumor Immunity  
Pharmacology



Role in Immunity

Relevance to  
Human Immunity



Toxicity



Human Trial:

- Toxicity
- Dose/Schedule
- Immunologic effects
- Effect on immune Tumor Microenvironment
- Anti-Tumor effect

Host genetics → Lifetime environmental exposures → TCR repertoire →

Patient  
Presenting  
for Treatment

Tumor evolution  
Metastases  
Evolution of Tumor-Host immune relationship  
Microbiome

Carcinogenesis:  
Mutations  
Altered gene expression  
Chronic inflammation

Tumor microenvironment and Host Anti-tumor immune response

T-cells

- How many?
- What type?
- Recognize tumor antigens?
- Breadth of antigen recognition (one, a few, many)
- Affinity of TCR for peptide-MHC complex
- Functional state
- Differentiated state
- Expression of inhibitory receptors
- Metabolic state and access to glucose and Oxygen
- Where located?

Tumor

- Antigens/neo-antigens
- Density of peptide/MHC complexes
- Expression of inhibitory ligands
- Expression of stimulatory ligands
- Production of inhibitory cytokines
- Production of other inhibitory substances
- Expression of chemokines
- Innate resistance to lytic mechanisms

Stroma/Other Immune Cells

- Treg
- MDSC
- Monocytes/macrophages/APC
- B-cells
- NK and NKT cells
- Tumor Vasculature
- Fibroblasts
- Secreted Inhibitory Factors
- Complement

Other

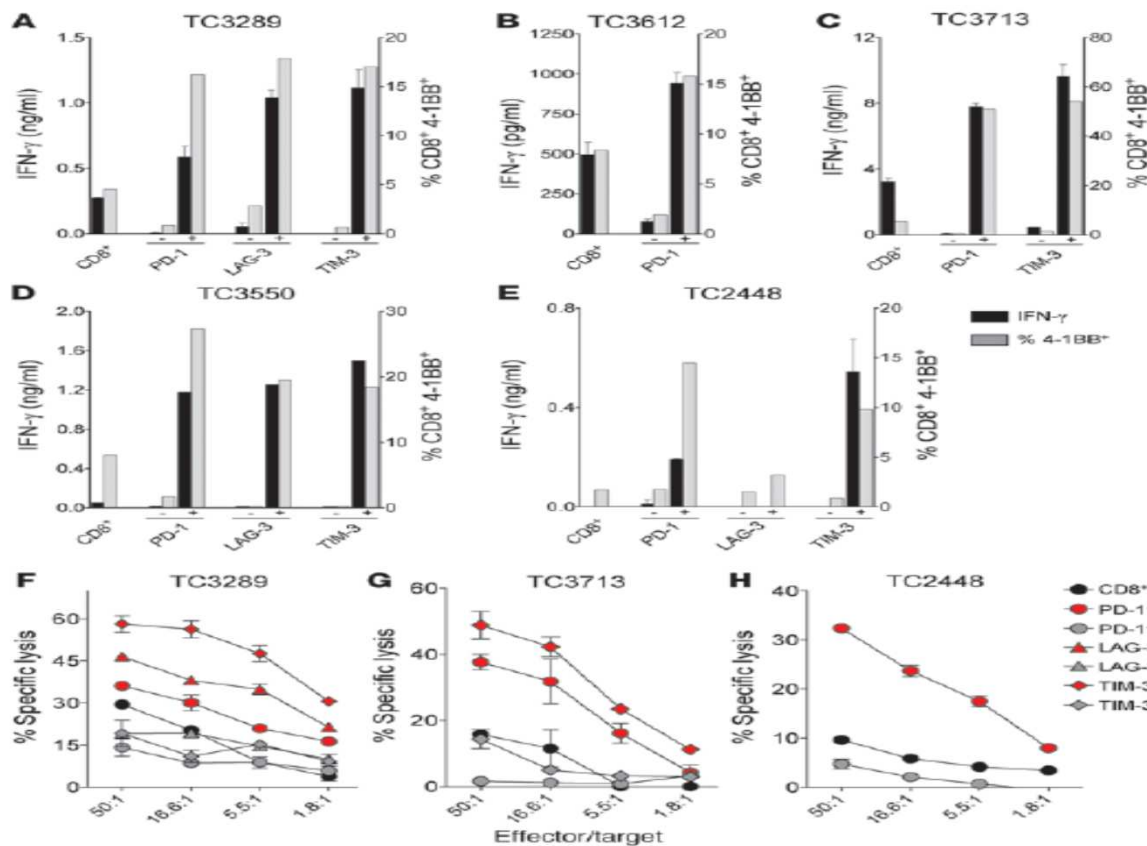
- Microbiome
- Lymph nodes
- Blood

$\Sigma$

Immune Intervention

Outcome

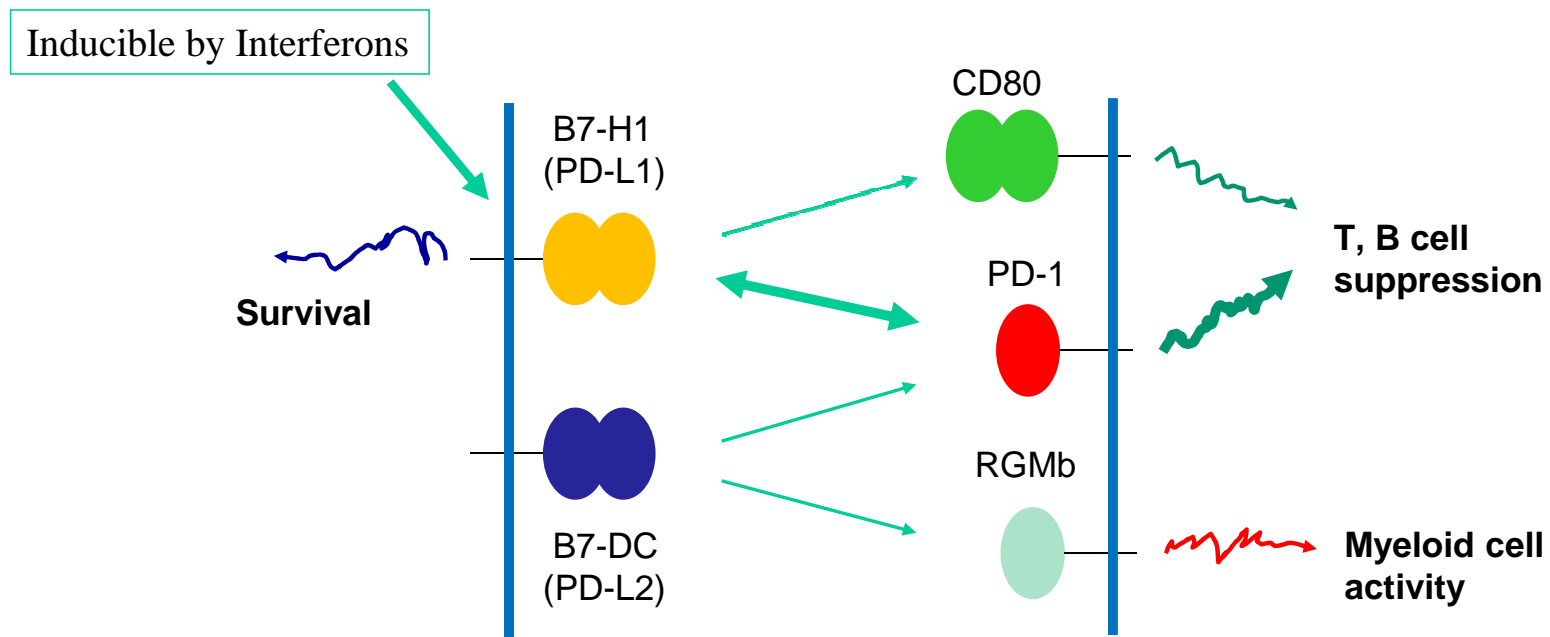
Tumor-specific T cells are contained in the PD-1+ TIL population and are functional after in vitro culture



**Figure 3**

Recognition and lysis of autologous tumor by CD8<sup>+</sup> TILs sorted based on PD-1, LAG-3, and TIM-3 expression. Bulk CD3<sup>+</sup>CD8<sup>+</sup> TILs were sorted to high purity from FrTu3289, FrTu3612, FrTu3713, FrTu3550, and FrTu2448 based on positive or negative expression of PD-1, LAG-3 and/or TIM-3, and expanded in vitro for 15 days. (A–E) Response of fresh tumor-derived TILs to their respective autologous tumor cell lines, TC3289 (A), TC3612 (B), TC3713 (C), TC3550 (D) and TC2448 (E). Reactivity was assessed by measuring IFN- $\gamma$  release (duplicates, mean  $\pm$  SD) and frequency of 4-1BB upregulation. (F–H) Cytolytic activity of fresh tumor-derived TILs in response to their respective autologous tumor cell lines, TC3289 (F), TC3713 (G), and TC2448 (H). Percentage of specific lysis at different effector/target ratios is shown as mean  $\pm$  SD.

# The PD-L1/PD-1 Pathway



Slide courtesy of Lieping Chen

# Spectrum of PD-1/PD-L1 Antagonist Activity

## Active

- **Melanoma**
- **Renal cancer (clear cell and non-clear cell)**
- **NSCLC – adenocarcinoma and squamous cell**
- Small cell lung cancer
- **Head and neck cancer**
- Gastric and gastroesophageal junction
- **MMR-repair deficient tumors (colon, cholangiocarcinoma)**
- **Bladder**
- Triple negative breast cancer
- Ovarian
- Hepatocellular carcinoma
- Thymoma
- Mesothelioma
- Cervical
- **Hodgkin lymphoma**
- Diffuse large cell lymphoma
- Follicular lymphoma
- T-cell lymphoma (cutaneous T-cell lymphomas, peripheral T-cell lymphoma)
- **Merkel cell**

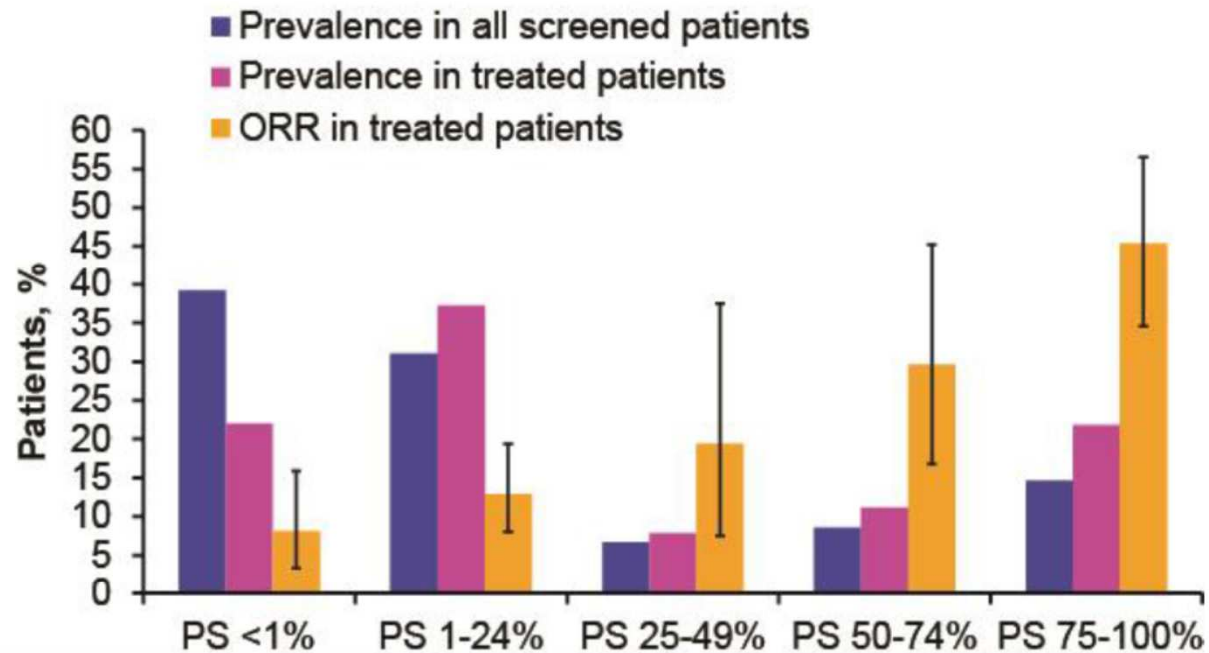
## Minimal to no activity

- Prostate cancer
- MMR+ (MSS) colon cancer
- Myeloma
- Pancreatic cancer

## Major PD-1/PD-L1 antagonists

- Nivolumab (anti-PD-1)
- Pembrolizumab (anti-PD-1)
- Atezolizumab (MPDL3280, anti-PD-L1)
- Durvalumab (anti-PD-L1)
- Avelumab (anti-PD-L1)

# Pembrolizumab ORR in NSCLC by PD-L1 Tumor Expression



All screened patients, n (%)	323 (39.2)	255 (31.0)	55 (6.7)	71 (8.6)	120 (14.6)
All treated patients, n (%)	87 (22.0)	147 (37.2)	27 (6.8)	39 (9.9)	72 (18.2)
ORR in treated patients, n (%) [95% CI]	7 (8.1) [3.3-15.9]	19 (12.9) [8.0-19.4]	6 (19.4) [7.5-37.5]	13 (29.6) [16.8-45.2]	39 (45.4) [34.6-56.5]



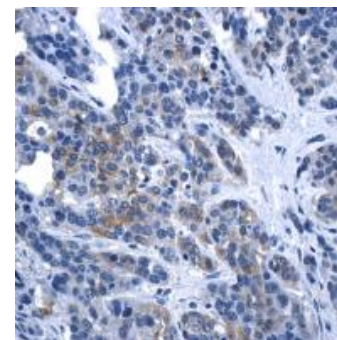
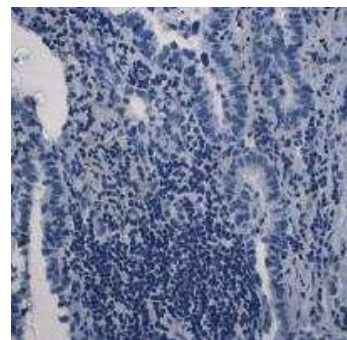
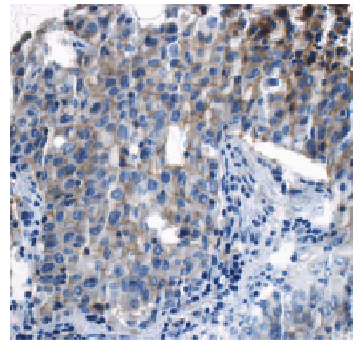
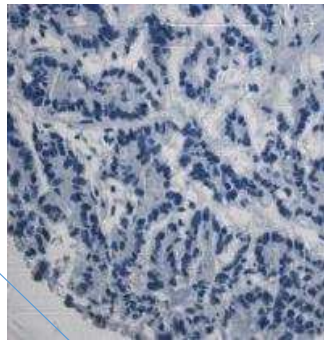
## Presence of PD-L1 or TILs<sup>1</sup>

PD-L1-/TIL-

PD-L1+/TIL+

PD-L1-/TIL+

PD-L1+/TIL-



Schalper and Rimm,  
Yale University

NSCLC

45%  
Type 1  
45%

17%  
Type 2  
41%

26%  
Type 3  
13%

12%  
Type 4  
1%

**Table 2.** Correlation of B7-H1 expression by melanocytes with the presence of immune cell infiltration.

Histology	Total	Number of cases/total cases (%)				P*
		B7-H1 <sup>++</sup>		B7-H1 <sup>-</sup>		
		TIL <sup>++</sup>	TIL <sup>-</sup>	TIL <sup>+</sup>	TIL <sup>-</sup>	
Benign nevi	40	14/14 (100)	0/14 (0)	4/26 (15)	22/26 (85)	<0.0001
Primary melanomas (in situ or invasive)	54	19/19 (100)	0/19 (0)	15/35 (43)	20/35 (57)	<0.0001
Metastases	56	23/24 (96)	1/24 (4)	7/32 (22)	25/32 (78)	<0.0001
All	150	56/57 (98)	1/57 (2)	26/93 (28)	67/93 (72)	<0.0001

\*Fisher's exact test, two-sided, was conducted on the 2 × 2 matrix defined by B7-H1 (±) expression and TIL (±) for each lesion type.  
#Including mild, moderate, and severe lymphocyte infiltrates and their associated histiocytes/macrophages.

†More than 5% melanocytes with membranous expression on IHC.

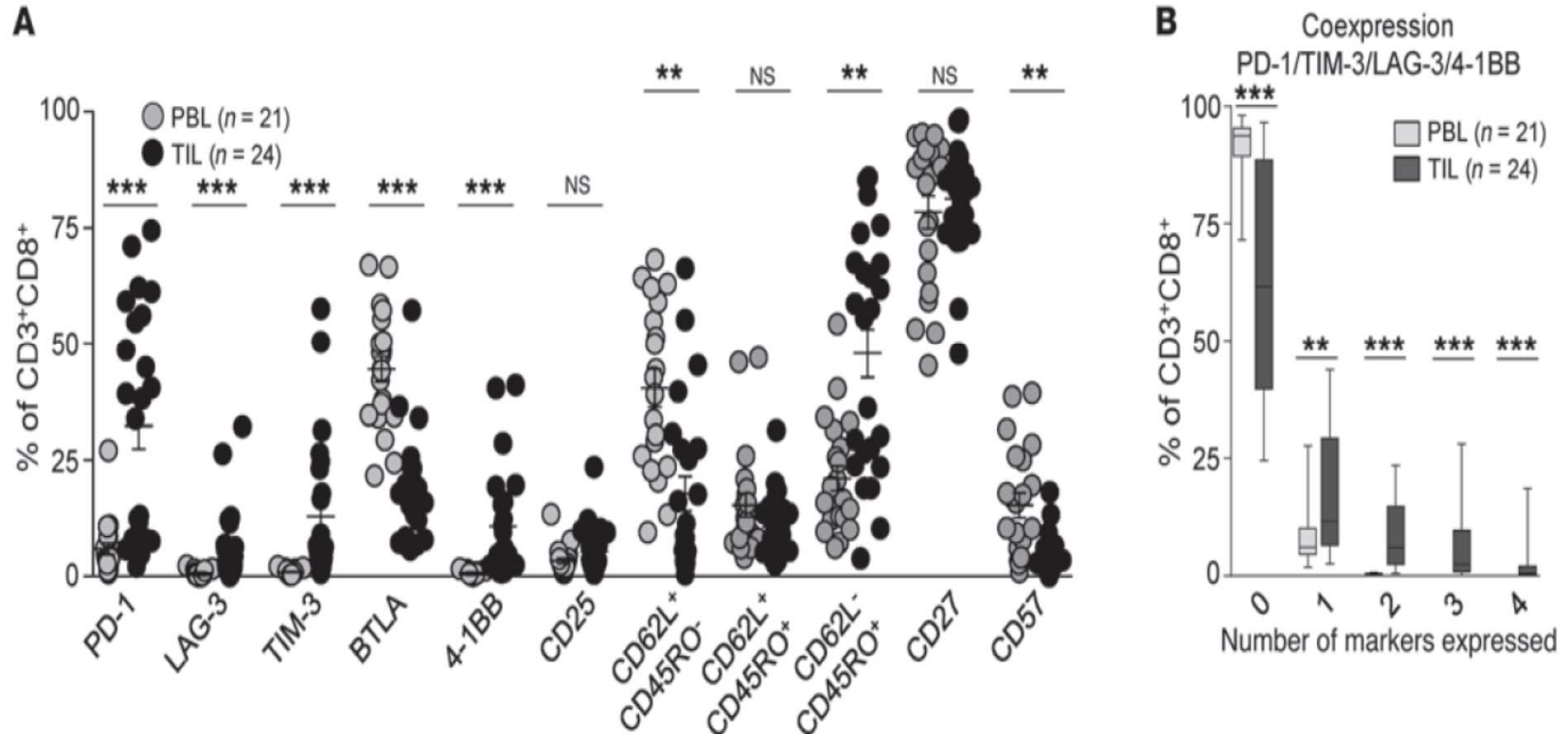
Melanoma

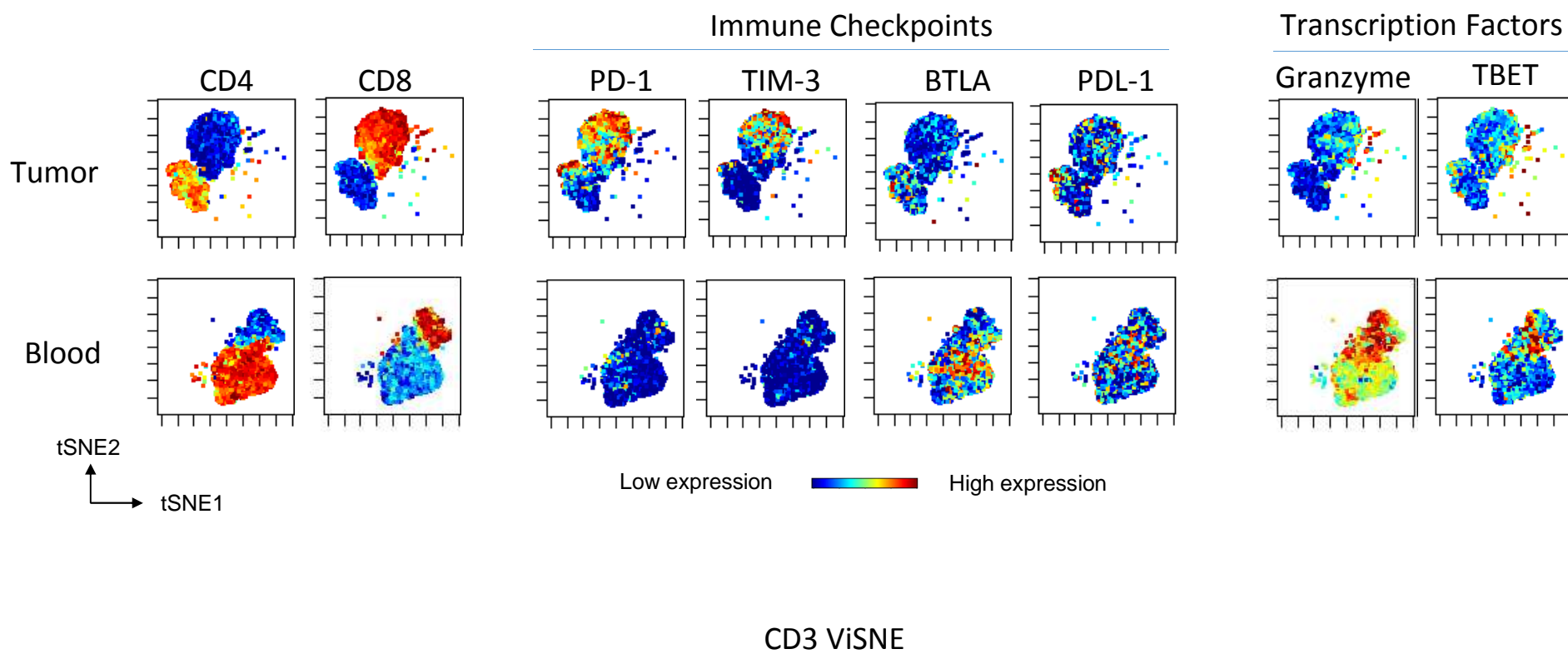
Taube et al




# Melanoma TIL – Expression of Co-inhibitory and Co-stimulatory Receptors (Gros et al)

The Journal of Clinical Investigation <http://www.jci.org> Volume 124 Number 5 May 2014





Provided by Kavita Dhodapkar, Yale University

Antigen Presenting Cell or Tumor	T-lymphocyte	Function (excluding Treg)
Peptide-MHC	T cell receptor	Signal 1
CD80/CD86 (B7.1, B7.2)	CD28/CTLA-4 	Stimulatory/ <i>inhibitory</i>
CEACAM-1	CEACAM-1	<i>inhibitory</i>
CD70	CD27	stimulatory
LIGHT	HVEM	stimulatory
HVEM	BTLA, CD160	<i>inhibitory</i>
<b>PD-L1 (B7-H1)</b> 	<b>PD-1</b> and CD80	<i>Inhibitory</i> (Th1)
PD-L2 (B7-DC)	PD1 and ?	<i>Inhibitory</i> (Th2) or stimulatory
OX40L	OX40	stimulatory
4-1BBL	CD137	stimulatory
CD40	CD40L	Stimulatory to DC/APC
B7-H3	?	<i>Inhibitory</i> or stimulatory
B7-H4	?	<i>inhibitory</i>
PD-1H (Vista)	?	<i>inhibitory</i>
GAL9	TIM-3	<i>inhibitory</i>
MHC class II	LAG-3	<i>inhibitory</i>
B7RP1	ICOS	stimulatory
MHC class I	KIR	<i>Inhibitory</i> or stimulatory
GITRL	GITR	stimulatory
CD48	2B4 (CD244)	<i>inhibitory</i>
HLA-G, HLA-E	ILT2, ILT4; NKG2a	<i>inhibitory</i>
MICA/B, ULBP-1, -2, -3, and -4+-	NKG2D	<i>Inhibitory</i> or stimulatory
CD200	CD200R	<i>inhibitory</i>
CD155	<i>TIGIT</i> /CD226	<i>Inhibitory</i> /stimulatory

### Other Inhibitory Factors

IDO

Treg

MDSC

Macrophages

TGF-beta

IL-10?

VEGF

Checkpoint Inhibitors



LAG3, TIM3, TIGIT, B7-H3, B7-H4, PD-1H (Vista), CD200, CEACAM1, KIR

MDSC  
Type 2 macrophages



HDACi, MER-TKi, CCR2i, CSF-1Ri, CKITi, ibrutinib,  
Anti-CD47 ('Don't Eat Me Signals' )

Treg



Anti-CCR4, anti-CTLA-4

Inhibitory  
Cytokines



Antibodies and small molecule inhibitors of TGF-beta or  
its receptors

Hypoxia/Adenosine

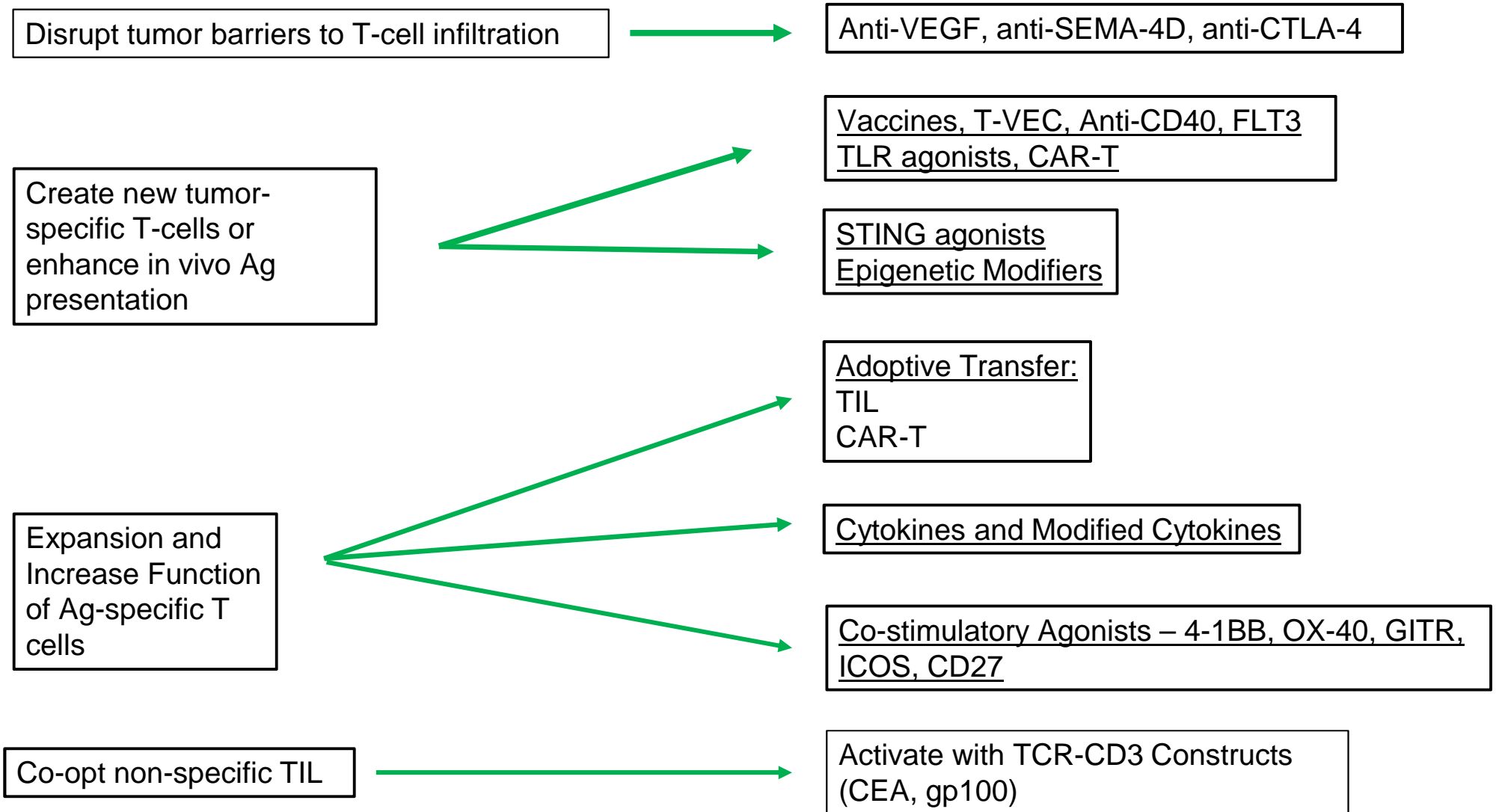


Adenosine 2AR inhibitors  
Anti-CD39, anti-CD73

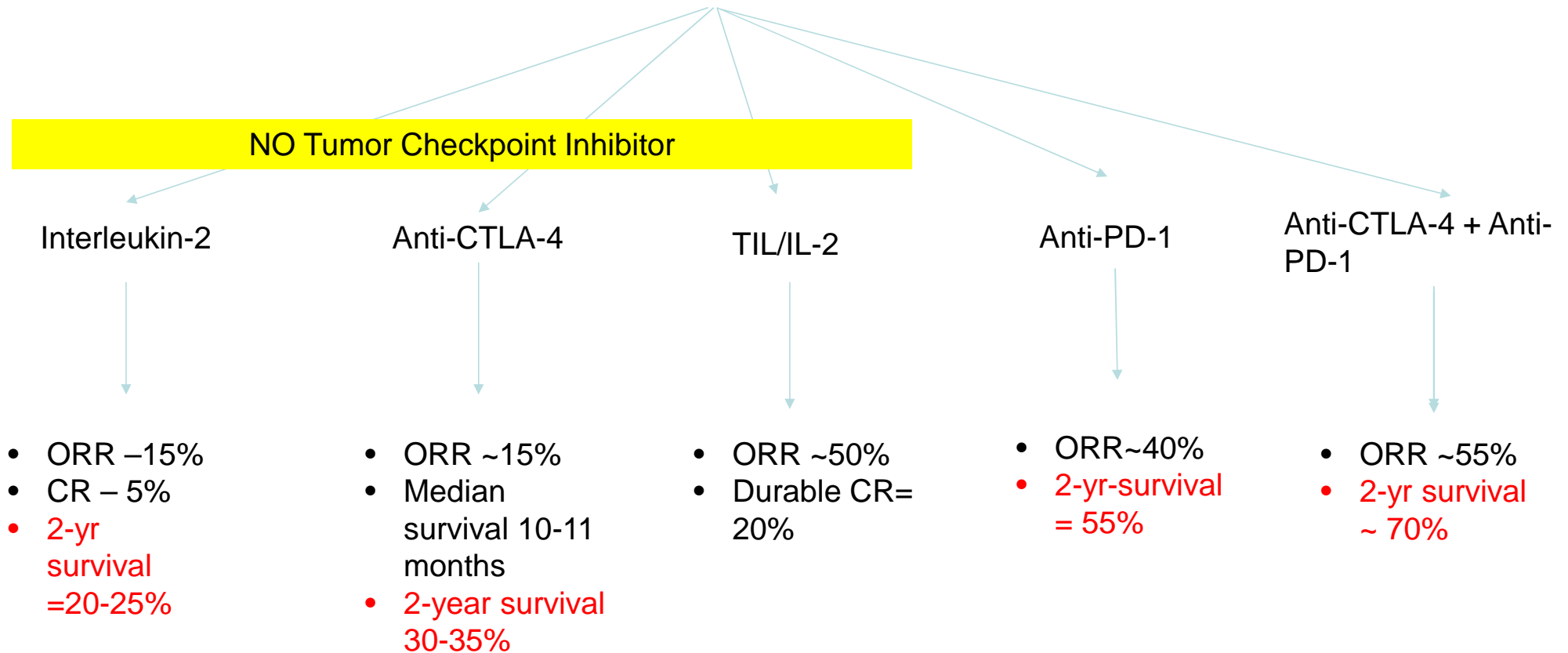
Metabolic Inhibitors and  
Prostaglandins



IDO inhibitors, Cox2 inhibitors



# Metastatic Melanoma



# A few critical questions for the future

- What are the relevant anti-tumor T-cells recognizing?
  - mutations, stem cell Ag, developmental Ag, tissue specific Ag
  - Are certain relevant tumor antigens ignored during tumor development?
- If no or minimal T cells in tumor, why?
  - not present in body, or can't get in?
- How does tumor biology control/determine anti-tumor immune response?
  - Type and number of mutations, signaling pathways (Sting, WNT/beta catenin, RTK, PI3K/AKT, RAS-RAF, IFN-g, epithelial to mesenchymal transition), vascularity and production of angiogenic cytokines, effects on stromal cells, antigen processing and presentation, resistance to lysis, pattern of organ metastases
- How do host factors influence developing an active anti-tumor immune response
  - Genetic polymorphisms, microbiome
- What are the critical T-cell features to mediate anti-tumor effect and what controls them? What is missing?
  - Proliferation and regenerative capacity, cytotoxicity, cytokine production, chemokine production, vascular effects, migration, ability to kill serially, formation of long-term memory (which subsets are most effective)
  - Magnitude of T-cell response?
  - T-cell receptor affinity for target peptide/MHC complex
  - Breadth of response (multiple antigens)?
- What are the most critical mechanisms for inhibiting T-cell activity within the tumor?
  - Checkpoints, hypoxia, glucose metabolism, essential amino acids, Treg, MDSC/type 2 macrophages, cytokines
- What other signals do T-cells need to function optimally and how best to deliver (even after checkpoint blockade)?
  - Cytokines, co-stimulatory ligands, functional APC
  - Multi-valent, requirement for Fc, cis versus trans, duration
- Are other types of cells (NK, NKT, monocytes, eos, granulocytes) or antibodies important?



Host genetics → Lifetime environmental exposures → TCR repertoire →

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for Treatment

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Metastases  
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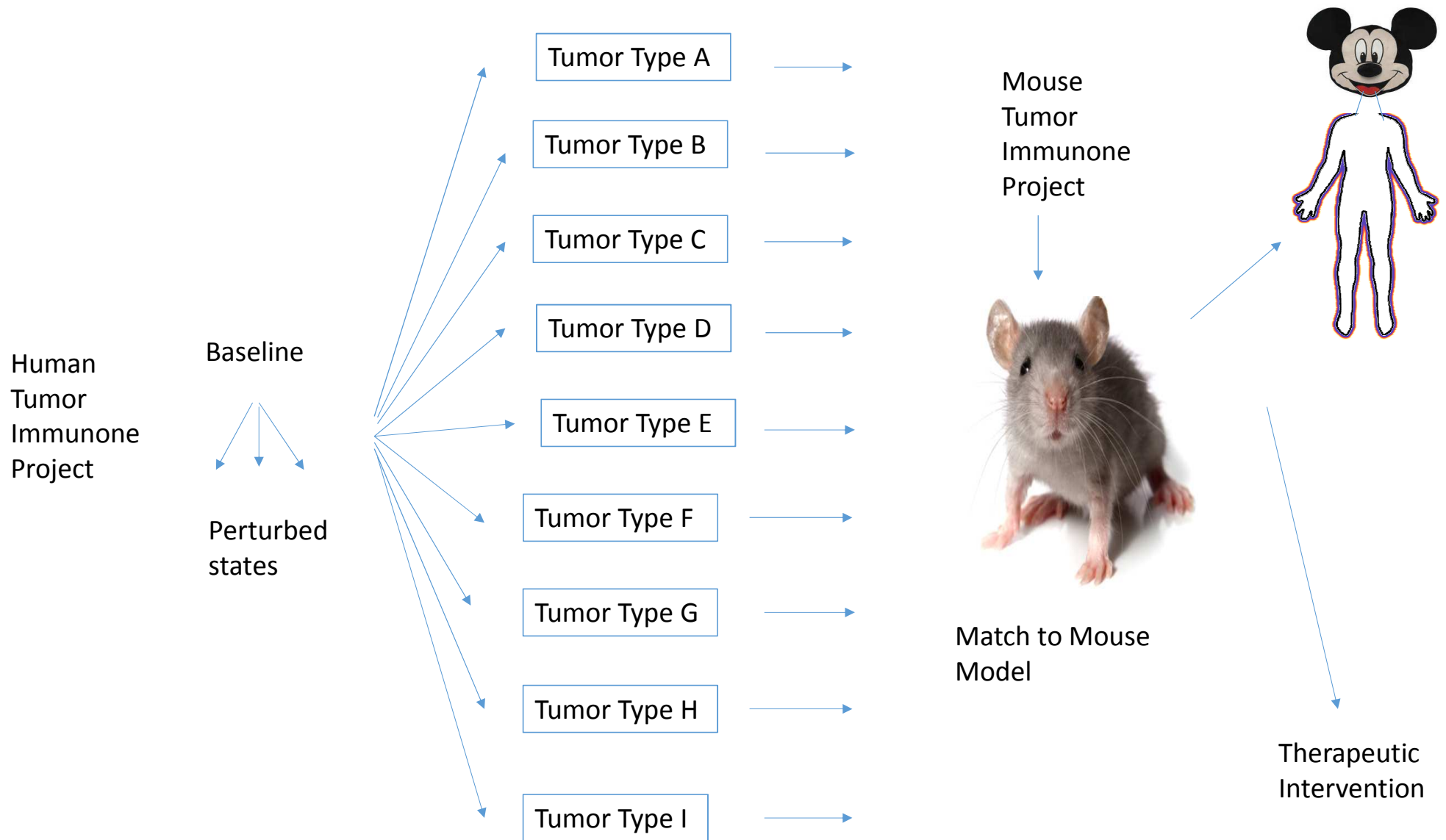
Other

- Microbiome
- Lymph nodes
- Blood

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Immune Intervention

Outcome



# Mechanisms of Toxicity for Immune Checkpoint Blockade or Co-Stimulatory Agents

- Mostly unknown
- May be epitope dependent (4-1BB)
- Cross-reactivity of Ab with normal tissue
- Activation of prior subclinical auto-immunity (recognition of self-Ag)
  - Prior genetic predisposition
  - Epitope spread
  - Cross-reactivity of tumor and normal tissue Ag
  - Increased effector cell function (Th1, Th2, Th17, other)
  - Reduced Treg function
- Cytokines may play role in pathology
- Role of antibody-dependent toxicity (serologic responses)
- Role of microbiome