

# What Relevant Immune Monitoring and Correlative Study Criteria Do We Use?

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# Disclosure

The following relationships exist related to this presentation:

None

The following relationships exist:

Advisory Board participant: EMD Serono, Novartis, Ziopharm

Special Advisor: Kite Pharma

# Correlations between Immune Monitoring and Clinical Outcome

Vaccination of vulvar intraepithelial neoplasia patients with long peptides from HPV16 {Welters '10}. Investigators tested lymphocyte proliferation and cytokine production to immunizing antigens as well as circulating regulatory T cells (Treg). Their study showed that the patients with the **strongest proliferation, positive IFN $\gamma$  and IL-5 production, and lowest Treg were those with complete responses (CR)** to therapy.

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A final example of an immune measure that several independent groups have correlated to clinical outcome is “epitope spreading” or “**determinant spreading**”. This acquisition of T cell and antibody reactivity to shared antigens or epitopes other than those used in a vaccine, has been observed in melanoma {Ranieri '00; Butterfield, '03; Ribas '04; Butterfield '08}, renal cancer {Wierecky '06 } and breast cancer {Disis '02 }.

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# Immunologic Biomarkers in Cancer

## *Organizers:*

**Lisa H. Butterfield, PhD**

*University of Pittsburgh*

**Mary (Nora) L. Disis, MD**

*University of Washington*

**Samir Khleif, MD**

*National Cancer Institute*

**Francesco Marincola, MD**

*National Institutes of Health*

**Magdalena Thurin, MD**

*National Institutes of Health*

## **Participating Organizations:**

- Association for Immunotherapy of Cancer (CIMT)
- Biotherapy Development Association (BDA)
- Cancer Vaccine Consortium (CVC) of the Cancer Research Institute (CRI)
- Food and Drug Administration (FDA)
- Italian Network for Tumor Biotherapy (NIBIT)
- Japanese Society of Cancer Immunology (JSCI)
- National Cancer Institute (NCI)
- National Institutes of Health (NIH)
- Nordic Center for Development of Antitumour Vaccines (NCV-network)

## Recommendations from the iSBTc-SITC/FDA/NCI Workshop on Immunotherapy Biomarkers

Lisa H. Butterfield<sup>1</sup>, A. Karolina Palucka<sup>4,5</sup>, Cedrik M. Britten<sup>23,24</sup>, Madhav V. Dhodapkar<sup>6</sup>, Leif Håkansson<sup>28</sup>, Sylvia Janetzki<sup>7</sup>, Yutaka Kawakami<sup>29</sup>, Thomas-Oliver Kleen<sup>9</sup>, Peter P. Lee<sup>10</sup>, Cristina Maccalli<sup>31</sup>, Holden T. Maecker<sup>11</sup>, Vernon C. Maino<sup>12</sup>, Michele Maio<sup>33</sup>, Anatoli Malyguine<sup>13</sup>, Giuseppe Masucci<sup>27</sup>, Graham Pawelec<sup>25</sup>, Douglas M. Potter<sup>2</sup>, Licia Rivoltini<sup>32</sup>, Lupe G. Salazar<sup>19</sup>, Dolores J. Schendel<sup>26</sup>, Craig L. Slingluff, Jr.<sup>20</sup>, Wenru Song<sup>21</sup>, David F. Stronck<sup>15</sup>, Hideaki Tahara<sup>30</sup>, Magdalena Thuri<sup>18</sup>, Giorgio Trinchieri<sup>14</sup>, Sjoerd H. van Der Burg<sup>34</sup>, Theresa L. Whiteside<sup>3</sup>, Jon M. Wigginton<sup>8</sup>, Francesco Marincola<sup>16</sup>, Samir Khleif<sup>17</sup>, Bernard A. Fox<sup>22</sup>, and Mary L. Disis<sup>19</sup>

### Variability:

1. Patient
2. Blood draw
3. Processing/cryo/thaw
4. Cellular product
5. Assay choice
6. Assay conduct
7. Assay analysis
8. Data reporting
9. Next cool new assay

### Recommendations:

1. Save DNA/RNA/cells/tumor; include healthy donor control
2. Standardized procedures
3. Standardized procedures
4. Functional assays to characterize/develop potency
5. Standardized, functional
6. SOP
7. Appropriate biostatistical methods
8. Full details, controls, QA
9. Sufficient blood/tissue to interrogate the samples *now*, as well as *later*, to generate new hypotheses.



# ImmunoAssay Proficiency Panel Program

**The program's objectives are:**

- 1) to offer an external validation program, and**
- 2) to enhance assay harmonization.**



**The Proficiency Panel Program seeks to identify:**  
**issues and deficiencies of current assay practices;**  
**sources for assay variability within and among institutions;**  
**protocol details that optimize assay performance.**

Define criteria for and support harmonization & validation of the assay for individual member laboratories.

Determine the value of resulting criteria in the cancer immunotherapy community.

Offer training programs to the cancer immunotherapy community to enhance assay performance and comparability between laboratories.

Establish immune assays as standard monitoring tools, if feasible.

**ELISPOT** -- Dr. Sylvia Janetzki, ZellNet Consulting

**ICS** -- Dr. Lisa McNeil, Wyeth

**Multimer** (Tetramer/Pentamer) Staining -- Dr. Pedro Romero, Ludwig Institute for Cancer Research, Lausanne, Switzerland, and Dr. Cedrik Britten, University of Leiden, The Netherlands

**Luminex** -- Dr. Michael Kalos, Univ. Pennsylvania

Central laboratory services, cells, and shipping are provided by the **Immunology Quality Assurance Center (IQAC)** of the Duke Human Vaccine Institute under the leadership of Dr. Thomas Denny.



Immunity 37, July 27, 2012

# T Cell Assays and MIATA: The Essential Minimum for Maximum Impact

C.M. Britten,\* S. Janetzki, L.H. Butterfield, G. Ferrari, C. Gouttefangeas, C. Huber, M. Kalos, H.I. Levitsky, H.T. Maecker, C.J.M. Melief, J. O'Donnell-Tormey, K. Odunsi, L.J. Old, T.H.M. Ottenhoff, C. Ottensmeier, G. Pawelec, M. Roederer, B.O. Roep, P. Romero, S.H. van der Burg, S. Walter, A. Hoos, and M.M. Davis.

Lack of conformity on reporting data hampers the ability to objectively evaluate data.

**Minimal Information About T cell Assays (“MIATA”):** we propose to generate recommendations on the minimum information required to allow an objective and thorough interpretation of published results from immunological T cell assays.  
Any recommendation in the proposed Modules is strictly based on supporting published data.

**Long-term goal of MIATA:** adapt the reporting framework for annotations of immune monitoring data sets from human studies for a central database, possibly in the context of a Human Immunity Project. This may include structured database vocabulary.

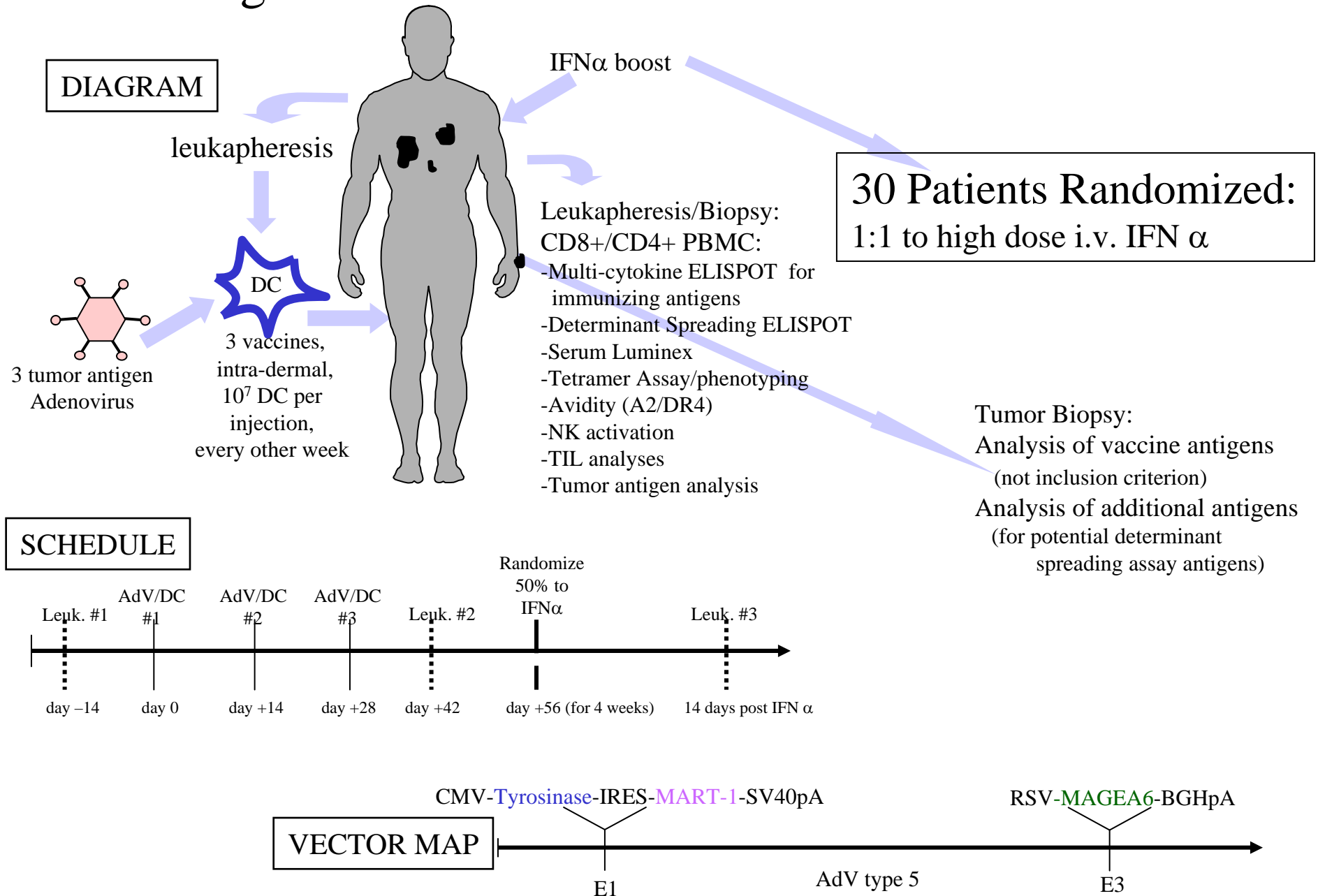


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Putting the recommendations into practice:

# Multi-Antigen-AdV-Transduced DC with IFN $\alpha$ Boost Trial



# Immune Response, Clinical Response and Determinant Spreading

Immune response to the vaccine-encoded antigens can correlate with clinical response, but the magnitude of CD8+ T cells to a specific immunizing peptide rarely defines the clinical responders.

Total *immune score*, *breadth* of immunity and *determinant spreading* may be more critical.

## How to strengthen and broaden immunity:

1. Multiple CD8/class I epitopes (dominant and subdominant)
2. Include cognate CD4 helper stimulation
3. Further improve the DC (subtype, maturation)
4. Promote innate immunity (NK cells)
5. Immunize with multiple antigens
6. Boost the immune response (cytokines)

# In Vivo Cross-Presentation

A recent study testing a multi-peptide vaccine in renal cancer patients demonstrated improved clinical outcome correlated with a larger number of peptides responded to {Walter '12 }. This last observation has been made in previous multi-peptide vaccine trials {Banchereau, '01 }.

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## **Immunologic biomarkers as correlates of clinical response to cancer immunotherapy.**

There is a need to identify immune biomarkers capable of predicting clinical response.....Biomarkers that measure the magnitude of the Type I immune response generated with immune therapy, epitope spreading, and autoimmunity are readily detected in the peripheral blood and, in clinical trials of cancer immunotherapy, have been associated with response to treatment. {Disis, CII, '11 }.

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## **How should this be assessed?**

T cells infiltrating the **tumor** (antigen source)?

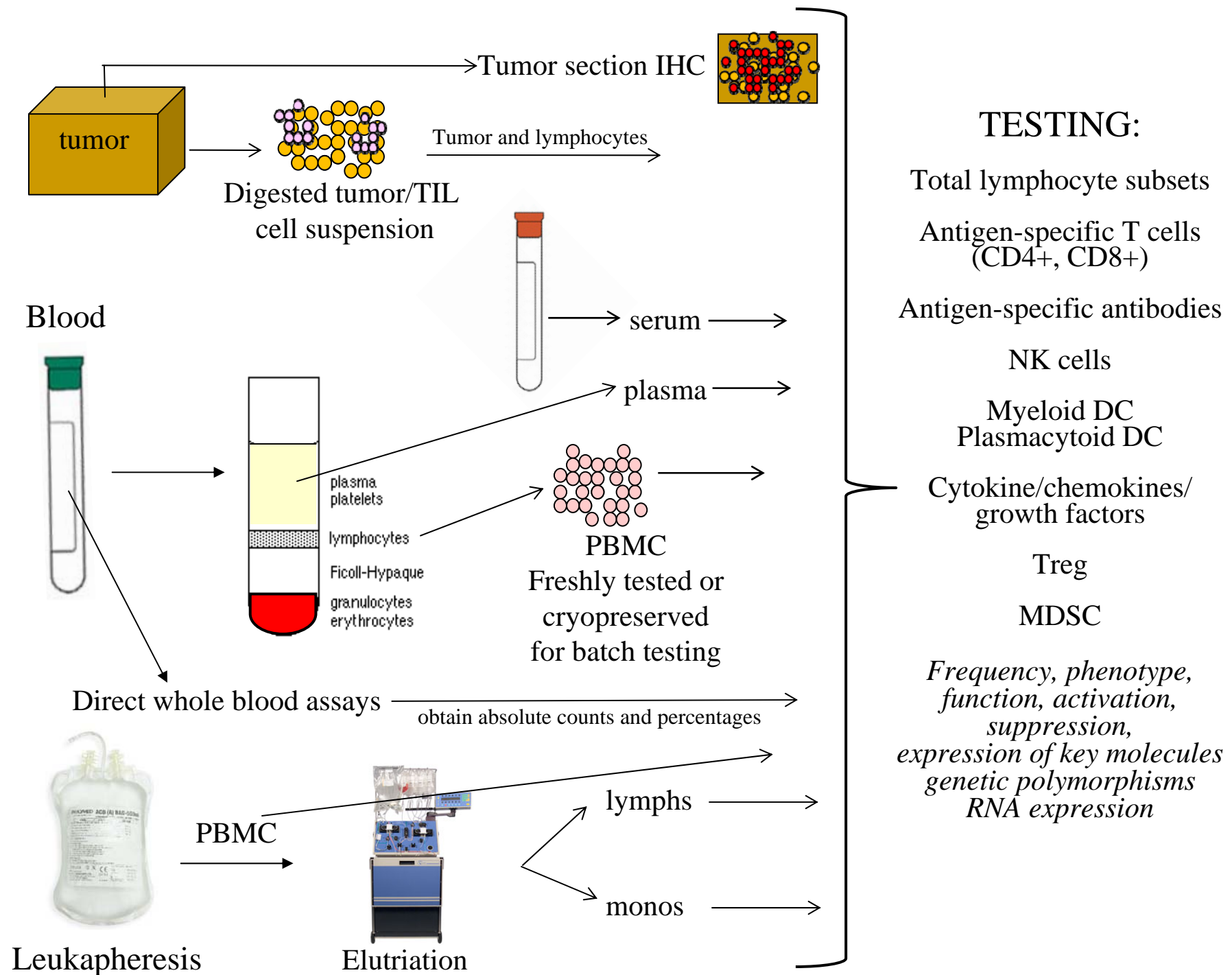
“Inflamed” **tumor** (mRNA/protein)?

Broad antigen recognition (many CD8+ T cell clones activated)?

Functional cellular + humoral immunity induced?

Reduced immune suppression?

Tumor private/mutated antigen responses?





# **AdVTMM2/DC +/- IFN $\alpha$ clinical trial (30 pt.)**

- a. Clinical outcomes in stage IV patients
- b. Immunologic outcomes (blood and TIL; post-vaccine and post-IFN $\alpha$ )
  - i. CD8+ T cells specific for vaccine antigens
  - ii. CD4+ T cells specific for vaccine antigens
  - iii. NK cell activation (flow), infiltration (TIL flow, IHC)
  - iv. T cell response to “spreading” antigens: gp100, NY-ESO-1, etc.
  - v. T and NK cell responses to autologous tumor (blood and TIL)
  - vi. Antibody responses to melanoma antigens
  - vii. Treg and MDSC regulatory status
- c. AdV responses
  - i. CD4+ hexon T cells (frequency, skewing)
  - ii. CD4+ and CD8+ T cells to the total viral proteins
  - iii. Anti-AdV total antibodies (ELISA)
  - iv. Anti-AdV neutralizing antibodies (flow)

# AdVTMM2/DC +/- IFN $\alpha$ clinical trial

a. Clinical outcomes in stage IV patients

b. Immunologic outcomes (blood and TIL; post-vaccine and post-IFN $\alpha$ )

**primary  
immune  
endpoint**

i. **CD8+ T cells specific for vaccine antigens**

ii. **CD4+ T cells specific for vaccine antigens**

iii. NK cell activation (flow), infiltration (TIL flow, IHC)

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v. T and NK cell responses to autologous tumor (blood and TIL)

vi. Antibody responses to melanoma antigens

vii. Treg and MDSC regulatory status

“exploratory”

“hypothesis-generating”

c. AdV responses

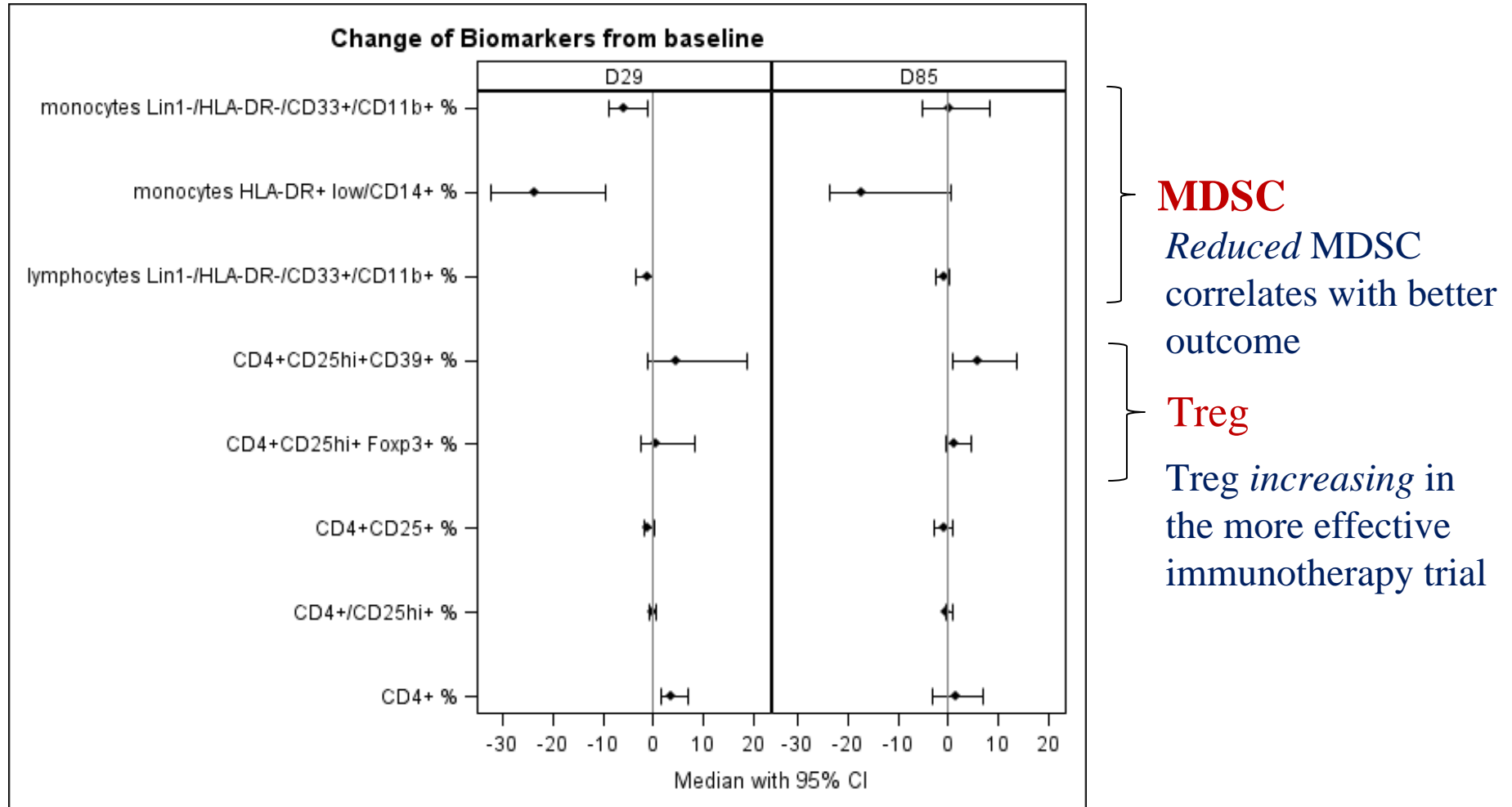
i. CD4+ hexon T cells (frequency, skewing)

ii. CD4+ and CD8+ T cells to the total viral proteins

iii. Anti-AdV total antibodies (ELISA)

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# Exploratory can be important



Forest plot of the multicolor flow cytometry data comparing T-reg and MDSC on PBMC at baseline and following treatment (day 29 and day 85) in patients treated with tremelimumab and interferon- $\alpha$ . Tarhini AA, J Immunother., 2012.

# Mechanism of Response and Biomarkers of Response

## from the AdVTMM2/DC +/- IFN $\alpha$ clinical trial

d. Vaccine Transcriptional Profiling (+/- AdVTMM2)

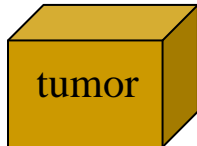
Compare to standard DC vaccine pheno/cytokine/Ag expression



e. Tumor Transcriptional Profiling (pre, post DC, post IFN)

How are tumors modulated by DC and IFN $\alpha$  therapy?

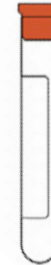
Baseline tumor profile?



f. Peripheral blood signaling

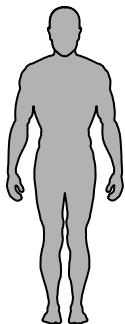
g. Serum profiling (autoimmunity antibodies, LDH, CRP)

Support previously identified serum biomarkers/autoimmunity?

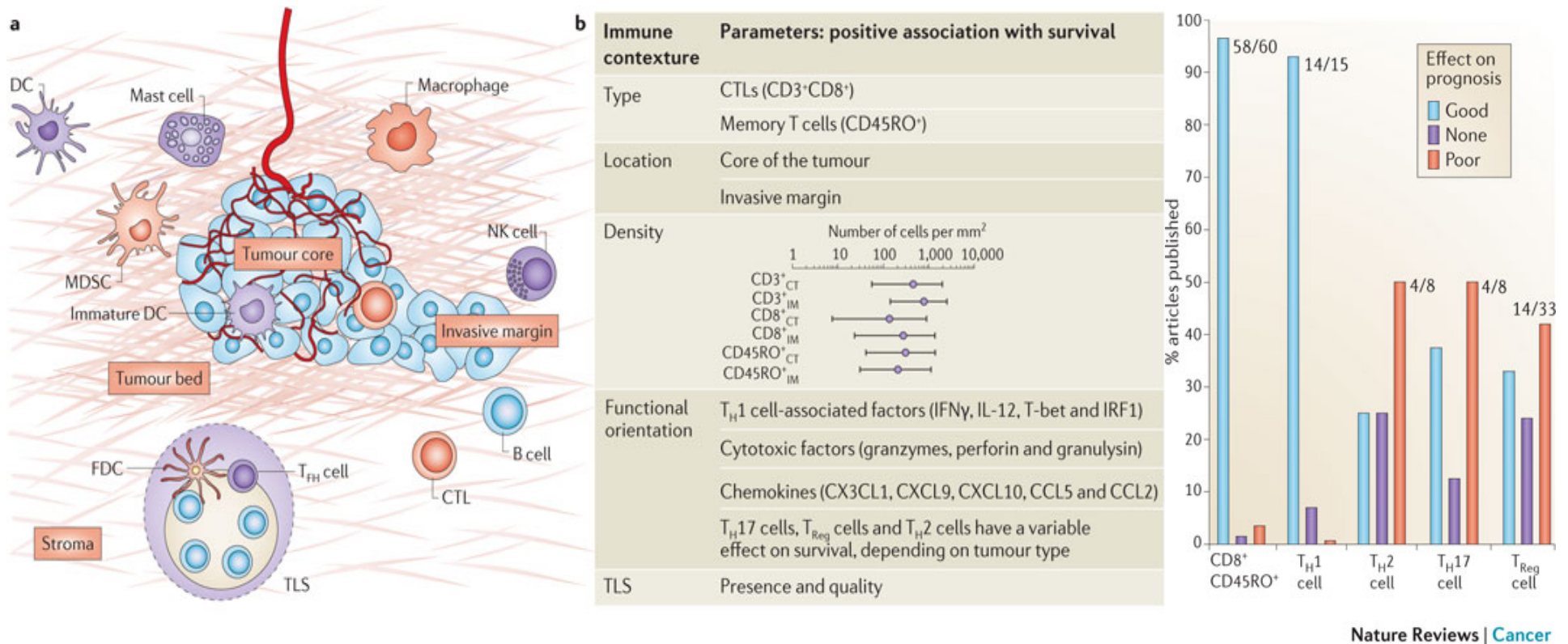


h. Molecular mimicry with mycoplasma

i. Germline DNA SNP analysis



# Immune Score



Nature Reviews | Cancer

Tumor anatomy showing the features of the immune contexture *J. Galon, W. Fridman; Science, 2006.*

## The Immune Score as a New Possible Approach for the Classification of Cancer

Jérôme Galon<sup>1,2,3,4,5\*</sup>, Franck Pagès<sup>1,2,3,4</sup>, Francesco M Marincola<sup>5,6</sup>, Magdalena Thurin<sup>7</sup>, Giorgio Trinchieri<sup>8</sup>, Bernard A Fox<sup>5,9,10</sup>, Thomas F Gajewski<sup>5,11</sup> and Paolo A Ascierto<sup>12,13</sup> **Journal of Translational Medicine 2012**

# Immunologic Monitoring and Cellular Products Laboratory

- The Cellular Products Laboratory (CPL) is dedicated to preparation of products for tumor vaccines and for cellular and gene therapy of cancer.
  - vaccine preparation
  - sterility and safety evaluations of generated products
  - assists in the preparation of INDs
  - operates according to FDA criteria for current good manufacturing practice (cGMP).
- The Immunologic Monitoring Laboratory (IML) is responsible for serial monitoring of immunologic functions in patients with cancer.
  - state-of-the-art immunologic assays
  - rigorous quality control program
  - development of new assays
  - advice on test selection and result interpretation
- The Tissue Procurement Facility (TPF) provides tissue and blood banking support under current Good Tissue Practice (cGTP) criteria.

CLIA certified; inspected by CAP and the state of PA; registered with FDA (3004571535)

FDA Master file: BB-MF-12244; FACT accredited



**UPMC** LIFE  
CHANGING  
MEDICINE

# Services

- Specimen intake, processing, banking
- Cell bank preparation
- Culture of peptide-specific T cells
- Culture of tumor infiltrating lymphocytes (TIL)
- DC preparation (peripheral blood, stem cells)
- Gene transfer: any human cells
- Expansion of stem cells
- Vaccine production for intra-tumor,-nodal, -lymphatic or intra-venous administration
- Peptide lyophilization, testing
- Adenovirus production (not fully GMP, with Vector Core)
- Assay standardization, normal control ranges
- Mixed lymphocyte-tumor cultures
- Cytotoxicity ( $^{51}\text{Cr}$ -release, FLOCA, CD107a).
- Proliferation ( $^3\text{H}$ -thymidine and CFSE)
- Multiparameter flow cytometry (effectors, regulatory (Treg, MDSC) cells)
- Single-cell T-cell assays (ELISPOT, CFC, and tetramer)
- Signaling molecules
- Frequency of apoptotic T cells
- Anti-tumor antibodies
- Cytokine/chemokine analysis (Luminex, ELISA)
- Data entry and analysis



# Standard Operating Procedures (SOPs)

UNIVERSITY OF PITTSBURGH CANCER INSTITUTE  
IMMUNOLOGIC MONITORING AND CELLULAR PRODUCTS LABORATORY

## **PROCEDURE FOR VIRUS INFECTIVITY NEUTRALIZATION ASSAY** **Purpose and Scope**

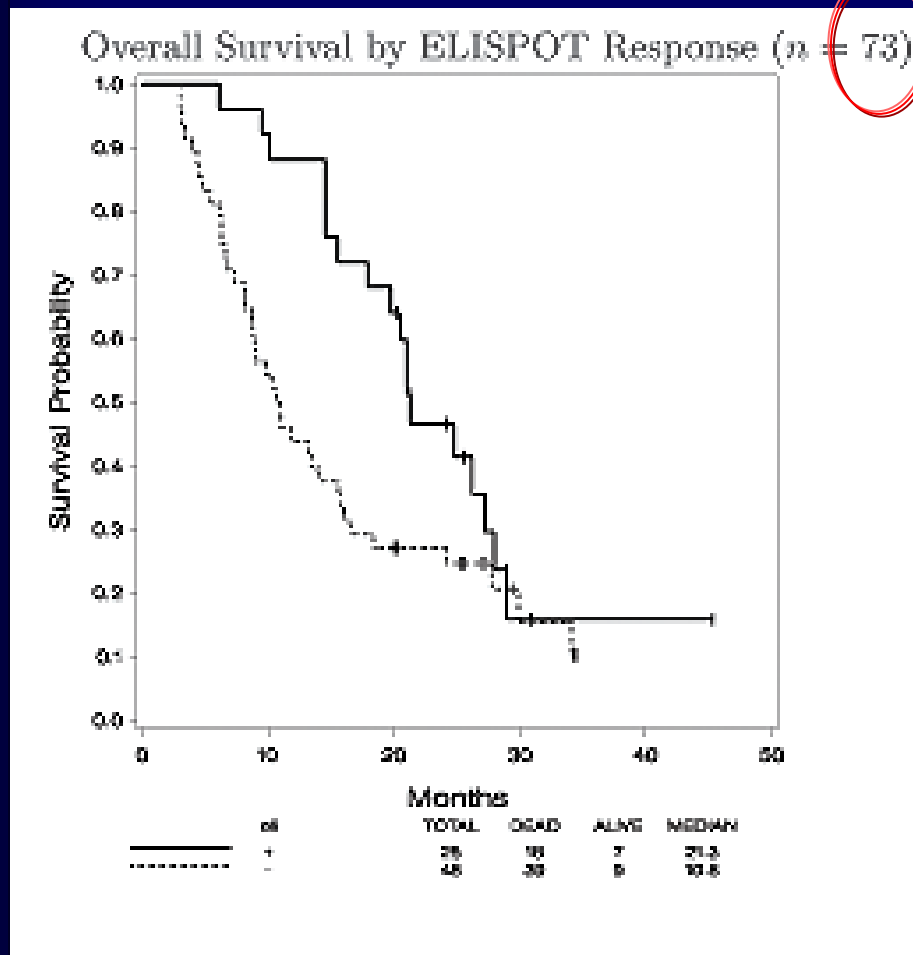
The purpose of this procedure is to test serum for the presence of antibodies to adenovirus which are capable of neutralizing viral infectivity. The assay tests decreasing dilutions of serum for the ability to reduce the efficiency by which an easily transduced cell line (A549) can be transduced with adenovirus encoding eGFP (AdVeGFP). The virus makes A549 cells GFP-positive, which is detected by flow cytometry. Serum samples with high levels of neutralizing antibodies will be able to reduce AdveGFP transduction of A549 cells, even after many serial dilutions.

This procedure is designed to quantitatively measure the inhibition of adenoviral transduction by serotype-specific neutralizing antibodies that are present in the patient serum. The serum dilutions for this assay are 1:4, 1:8, 1:16, 1:32, 1:64, 1:128, 1:256 and 1:512 in addition to a negative control (A549 cells + medium) and a positive control (A549 cells + virus + medium).



# Immune Response Correlates with Overall Survival

Multiple melanoma antigen peptide vaccine  $\pm$  GM-CSF  $\pm$  IFN $\alpha$ 2b



The Kaplan-Meier plot for OS by immune response status is shown for E1696 (Phase II).

*There was a significant difference in OS by immune response status. Immune responders lived longer than the non-immune responders (median OS 21.3 versus 10.8 months,  $p=0.033$ ).*

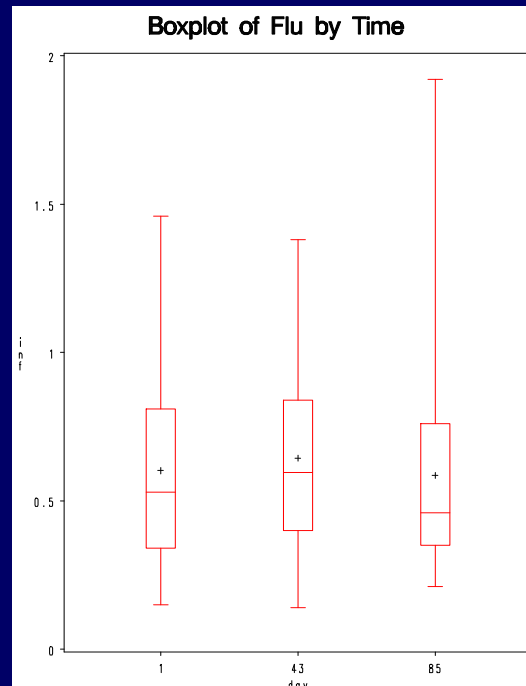
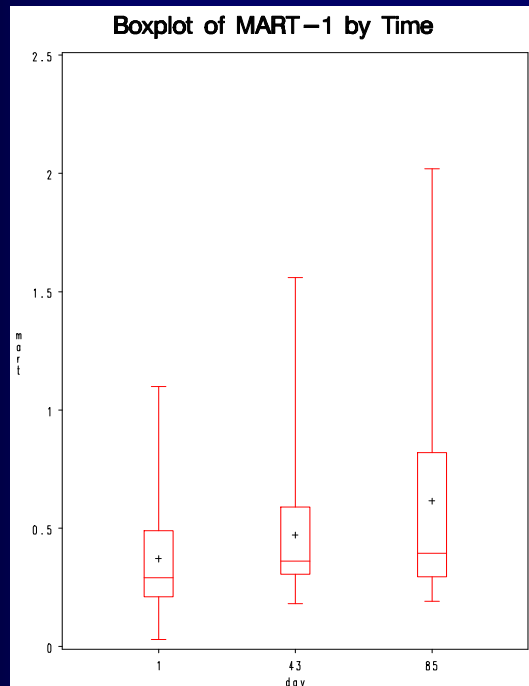
(Kirkwood, J.M., Clin. Cancer Res. 2009)

eastern cooperative oncology group



# Immune Response: E1696

## Melanoma antigen peptide-specific CD8+ T cells



%MART-1  
CD8+ cells: .29% .36% .39%

.53% .53% .43%

%effector  
cells: 10% 16% 18%  
(p=0.048)

17% 17% 16%  
(p = ns)

### MHC Tetramer Analysis:

The frequency of vaccine peptide-specific CD8+ T cells was measured by MHC tetramers, showing significant increases for all 3 melanoma antigen peptides (not Flu).

The MART-1 and gp100-specific cells differentiated towards effector cells with vaccination.

# AdVTMM2/DC +/- IFN $\alpha$ trial pt. #1 DC Vaccine

## •Vaccine Phenotype:

CPL#	HLA-DR+	CD86+	CD80+	CD83+	CCR7+	CD40+	CD11c+
CPL-12-19	98%	95%	93%	88%	28%	95%	96%

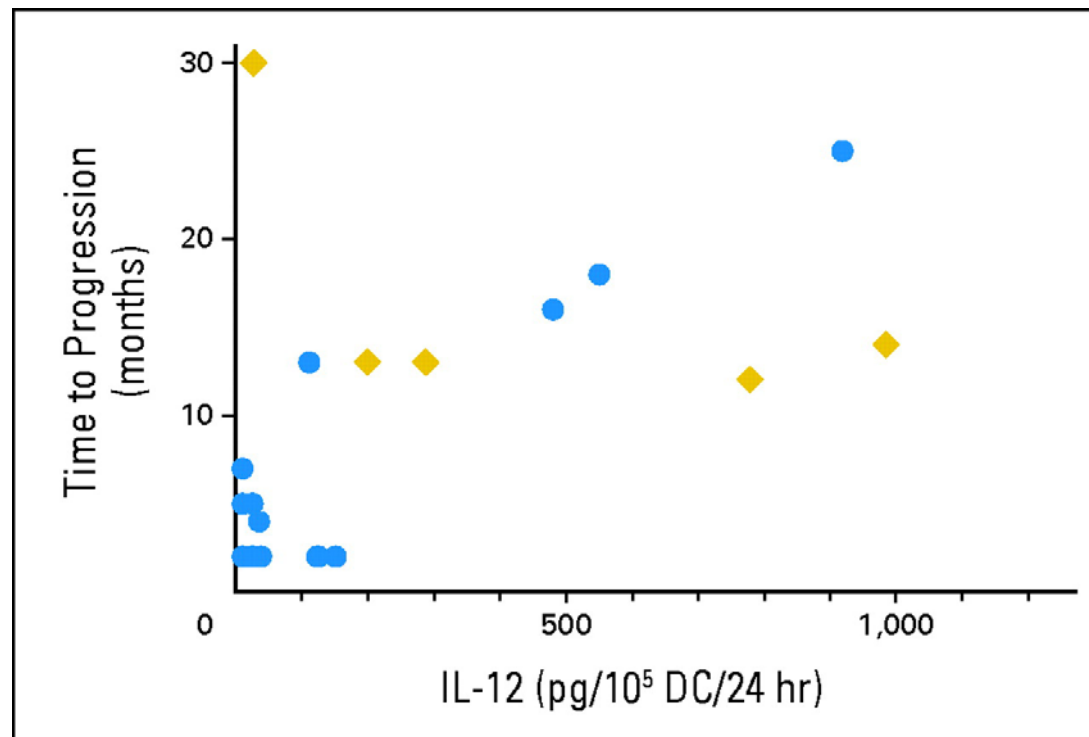
CPL#	SAMPLE	Intracellular: MART-1	TYROSINASE	MAGE-A6
CPL-12-19	mDC (control)	0	2	4
	prematAdVTMM2/DC (vaccine)	6%+	18%+	10%+

## •IL-12p70/IL-10 production by vaccine:

IL-12p70	HD DC	Pt. mDC	Pt. AdV/DC vaccine		IL-10	HD DC	Pt. mDC	Pt. AdV/DC vaccine
DC only	0	2430	<b>1237</b>			0	8	<b>0</b>
DC+CD40L	323	5551	<b>4024</b>			505	130	<b>84</b>
DC+LPS	99	2840	<b>1344</b>			285	23	<b>19</b>
DC+CD40L+LPS	877	6102	<b>3838</b>			1043	411	<b>268</b>

## •Immature DC, matDC, premat/AdV/DC (vaccine): *frozen for mRNA per SOP*

# DC Vaccine IL-12p70 Production and Clinical Outcome



IL-12p70 production levels positively correlated with time to progression.  $P$  .0255 is based on Cox regression followed by likelihood-ratio test. Circles indicate patients who have already experienced disease progression; diamonds represent patients who have not experienced recurrence to date (Okada, H. JCO 2010). Assay standardization (Butterfield, JIT 2008).

# Anti-CTLA-4, Anti-PD-1/PD-L1

- Absolute lymphocyte count (ALC) >1,000, increased ALC
- Serum LDH, CRP, IL-17
- Treg
- MDSC
- NY-ESO-1 cellular and humoral response
- Tumor IDO
- Tumor TIL
- Tumor PD-L1
- Ki67/EOMES in CD4+/CD8+ T cells
- Exhausted T cells (PD-1/TIM3/LAG-3)

# In Vivo Cross-Presentation

## How should this be assessed?

T cells infiltrating the **tumor** (antigen source)?

“Inflamed” **tumor** (mRNA/protein)?

**tumor**

Broad antigen recognition (many CD8+ T cell clones activated)?

Functional cellular + humoral immunity induced?

**effectors**

Tumor private/mutated antigen responses?

Reduced immune suppression?

**suppression**

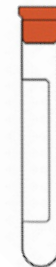
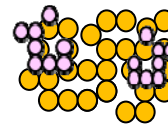
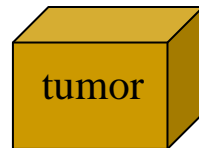
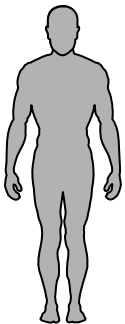
# Recommendations:

## Variability:

1. Patient
2. Blood draw
3. Processing/cryo/thaw
4. Cellular product
5. Assay choice
6. Assay conduct
7. Assay analysis
8. Data reporting
9. Next cool new assay

## Recommendations:

1. Save DNA/RNA/cells/tumor; include healthy donor control
2. Standardized procedures
3. Standardized procedures
4. Functional assays to characterize/develop potency
5. Standardized, functional
6. SOP
7. Appropriate biostatistical methods
8. Full details, controls, QA
9. Sufficient blood/tissue to interrogate the samples *now*, as well as *later*, to generate new hypotheses.





# Acknowledgements



## SITC Immunotherapy Biomarkers Taskforce

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University of Pittsburgh Cancer Institute  
NIH RO1 CA104524, NIH RO1 CA 138635  
NIH 1P50 CA121973  
(Kirkwood, J.M., PI, Skin SPORE)  
The Pittsburgh Foundation  
The Hillman Foundation