

**Biomarcatori e Tecniche di Diagnostica per Immagini
nella Valutazione dell'Infiammazione in Oncologia**



***Overview sui meccanismi biologici legati all'infiammazione
delle patologie tumorali e rilevanza clinica***

Gennaro Ciliberto e Paola Nisticò

IRCCS – Istituto Nazionale Tumori «Regina Elena»

22 Febbraio 2019

Istituto Superiore di Sanità



RUDOLF VIRCHOW

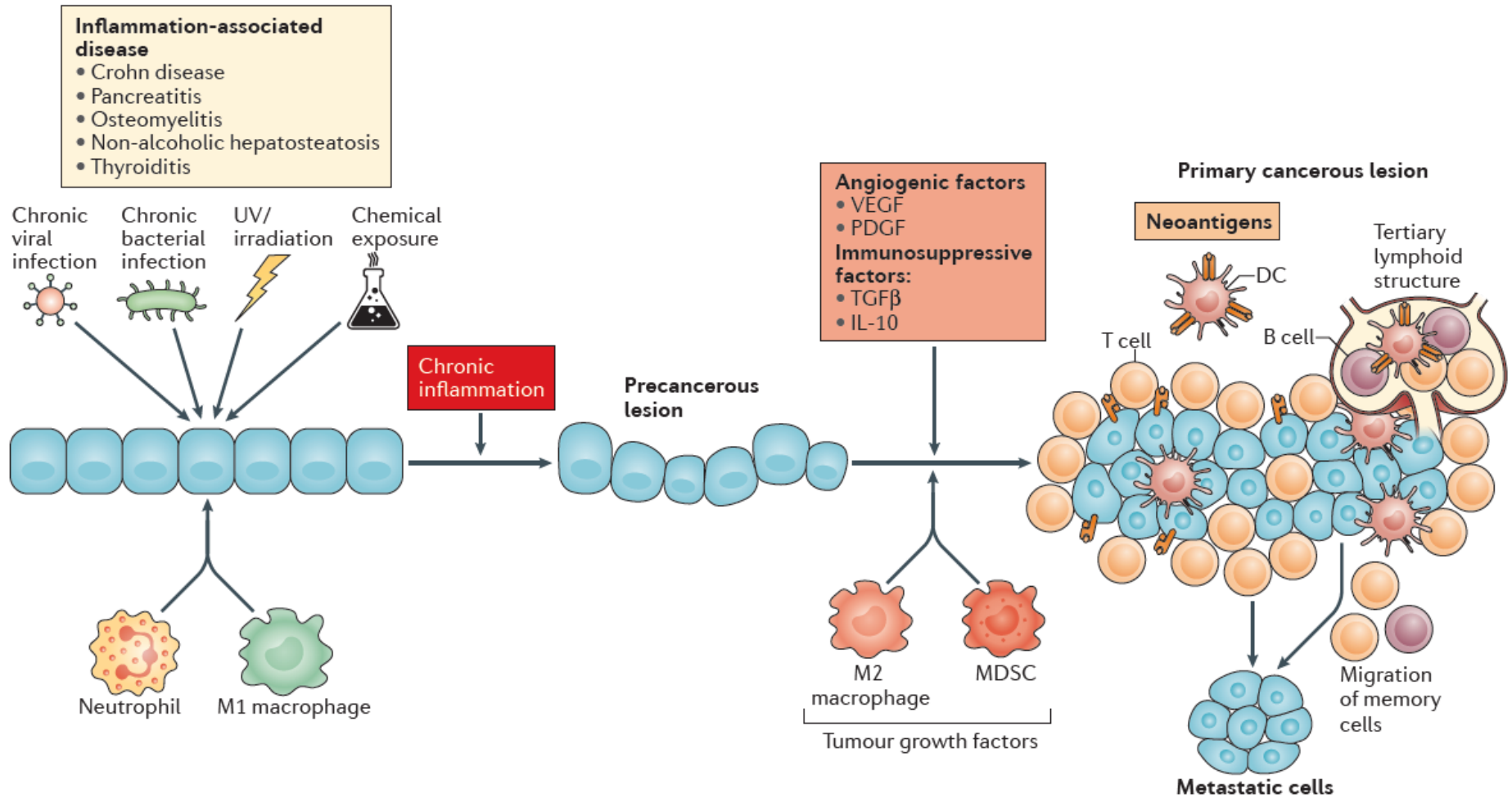


In 1863, Virchow first postulated that cancer originates at sites of **CHRONIC INFLAMMATION**

A functional link exists between inflammation and cancer

- **Increased cancer incidence in individuals affected by chronic inflammatory disorders**
- **Reduced cancer incidence in patients treated with long term anti-inflammatory drugs**
- **The anti-inflammatory therapy with canakinumab targeting the interleukin-1 β innate immunity pathway could significantly reduce incident lung cancer and lung cancer mortality (Trial of canakinumab in 10061 patients) *Ridker, et al The Lancet 2017***

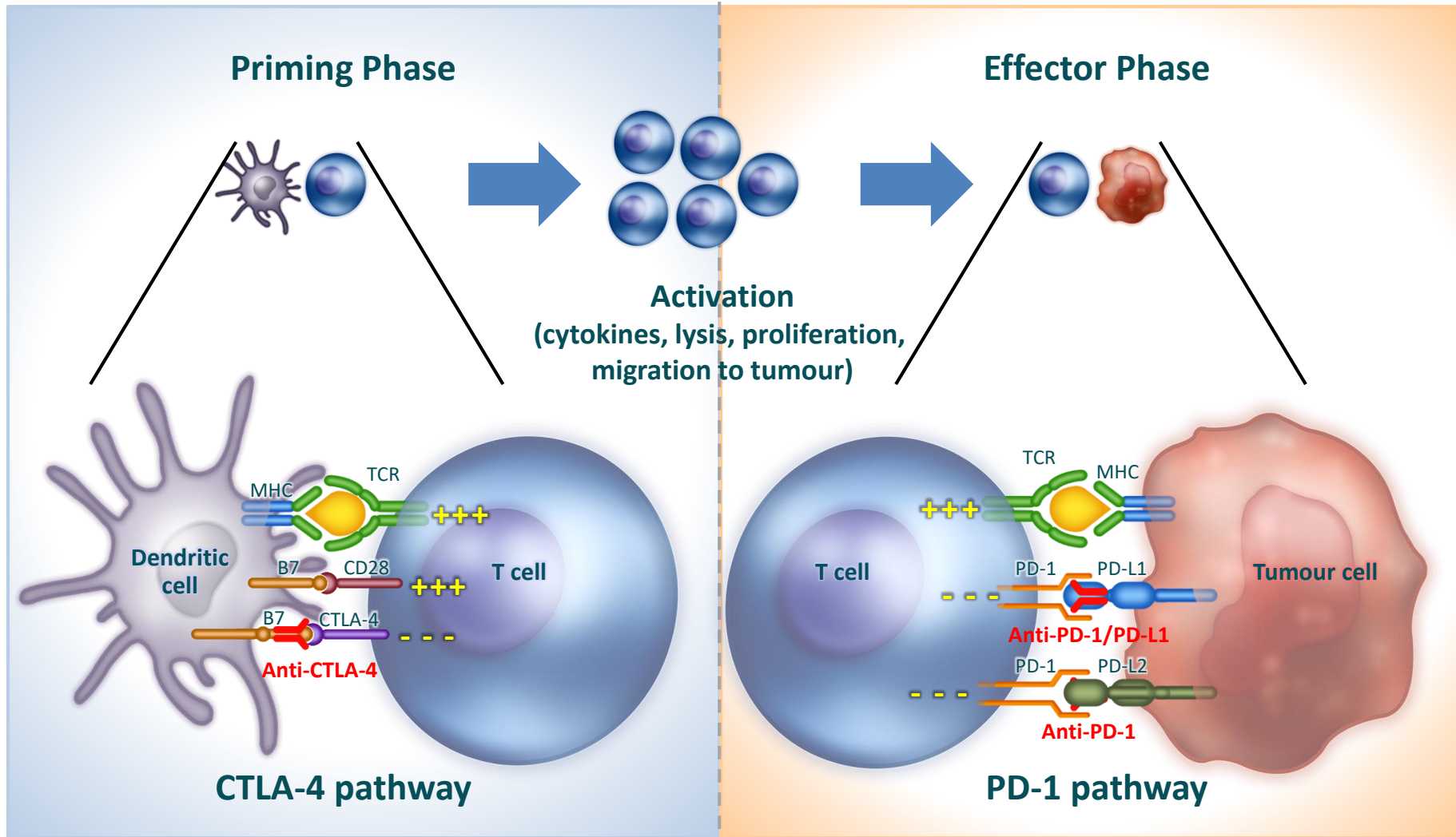
Tumorigenesis goes *hand in hand* with chronic inflammation at the tumor sites



Key Concept

Cancer has to be conceived as a complex organ, not anymore as a mass composed just of tumor cells

This concept is at the basis of the success (or lack thereof) of ICI



**Intrinsic features of Tumor Cells do
not explain alone response to ICI**

**The case of Tumor
Mutational Burden**

Tumor mutational burden predicts ONLY PARTIALLY response to ICI

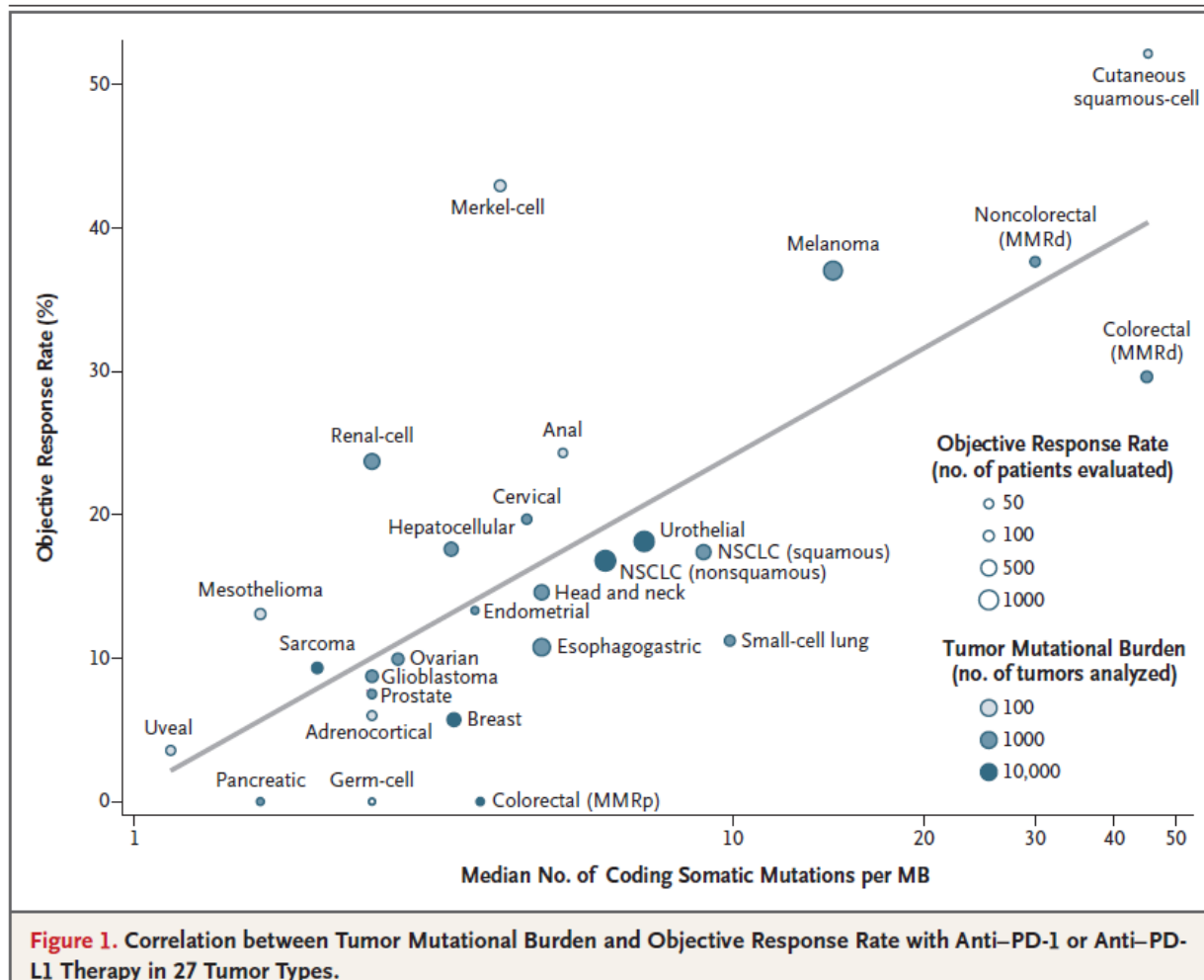
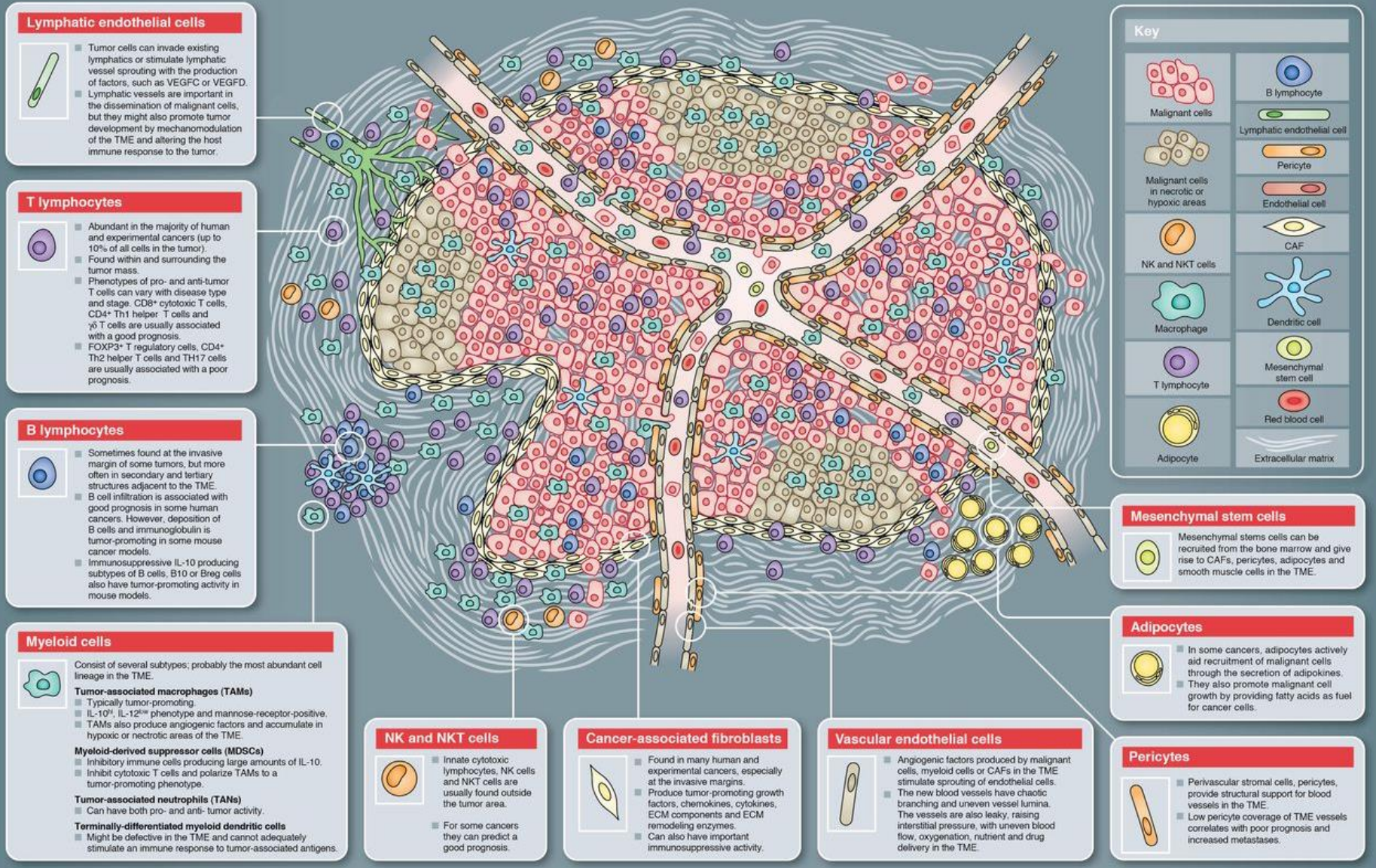


Figure 1. Correlation between Tumor Mutational Burden and Objective Response Rate with Anti-PD-1 or Anti-PD-L1 Therapy in 27 Tumor Types.

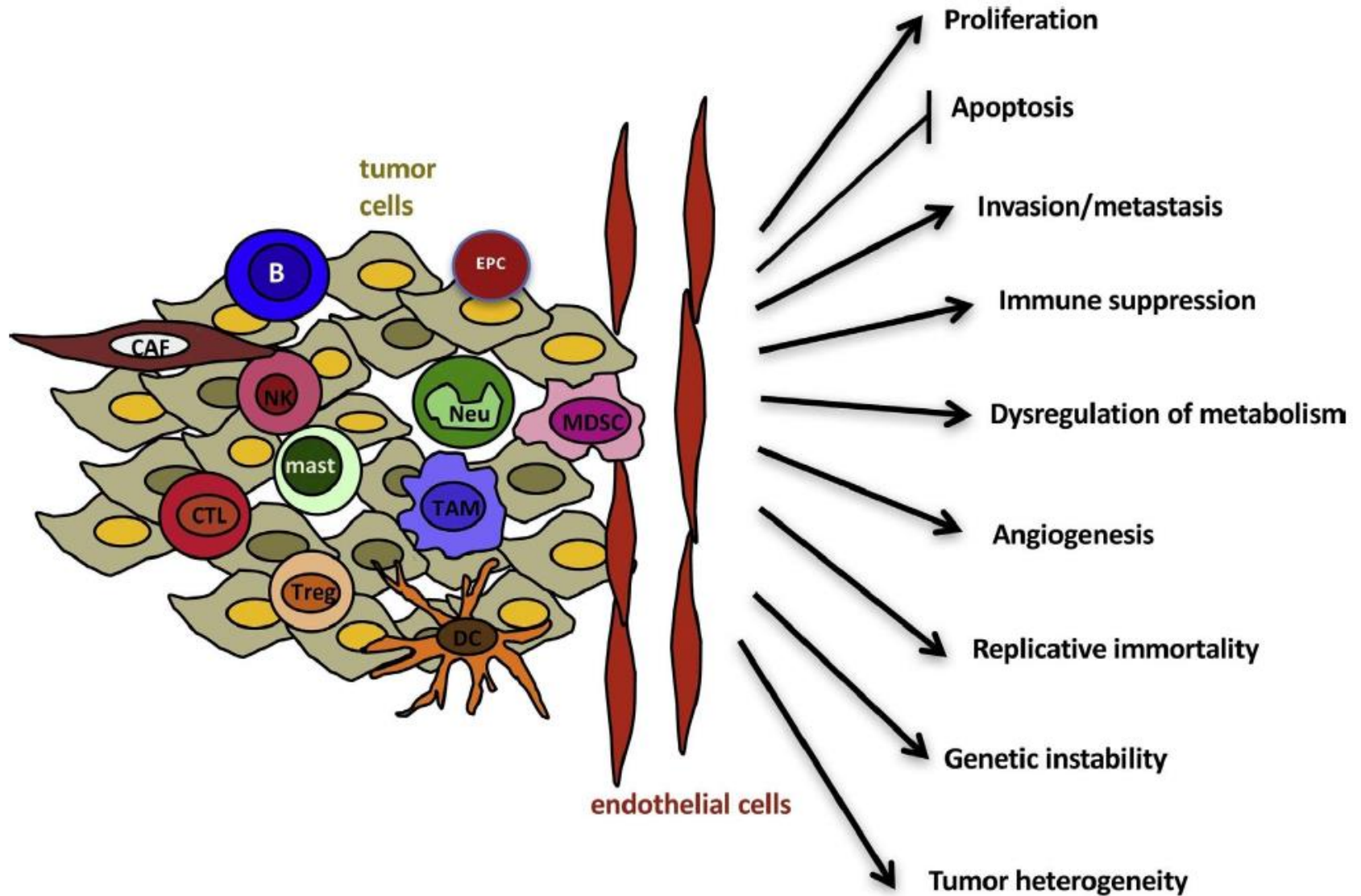
**Intrinsic features of Tumor Cells do
not explain alone response to ICI**

**Response to ICI depends on something
more: the interplay with the tumor
microenvironment**

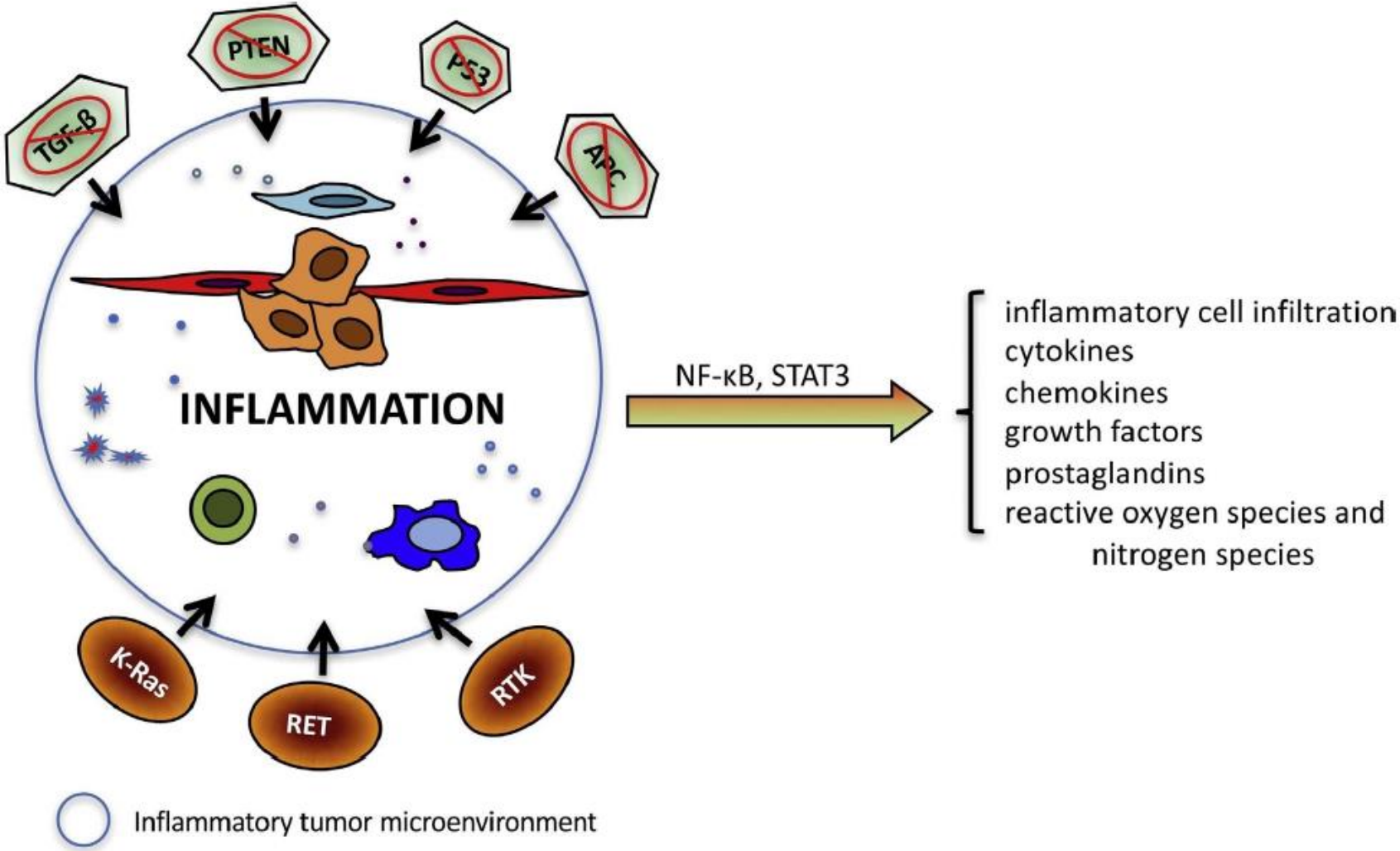
Cancer cells are embedded in a complex tumor microenvironment: a highly trafficked network



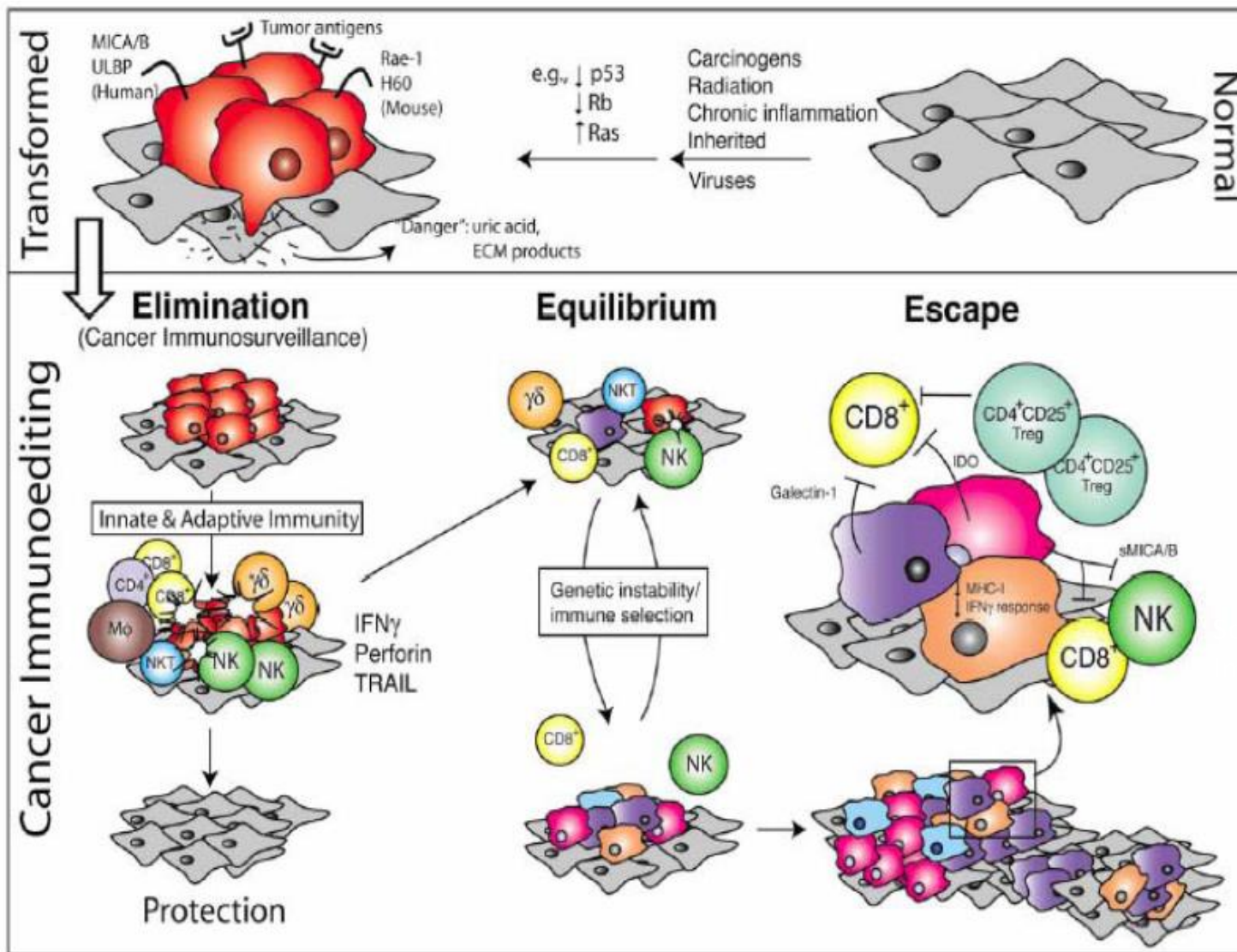
Players in the TME and the outcome resulting from tumor-associated inflammation



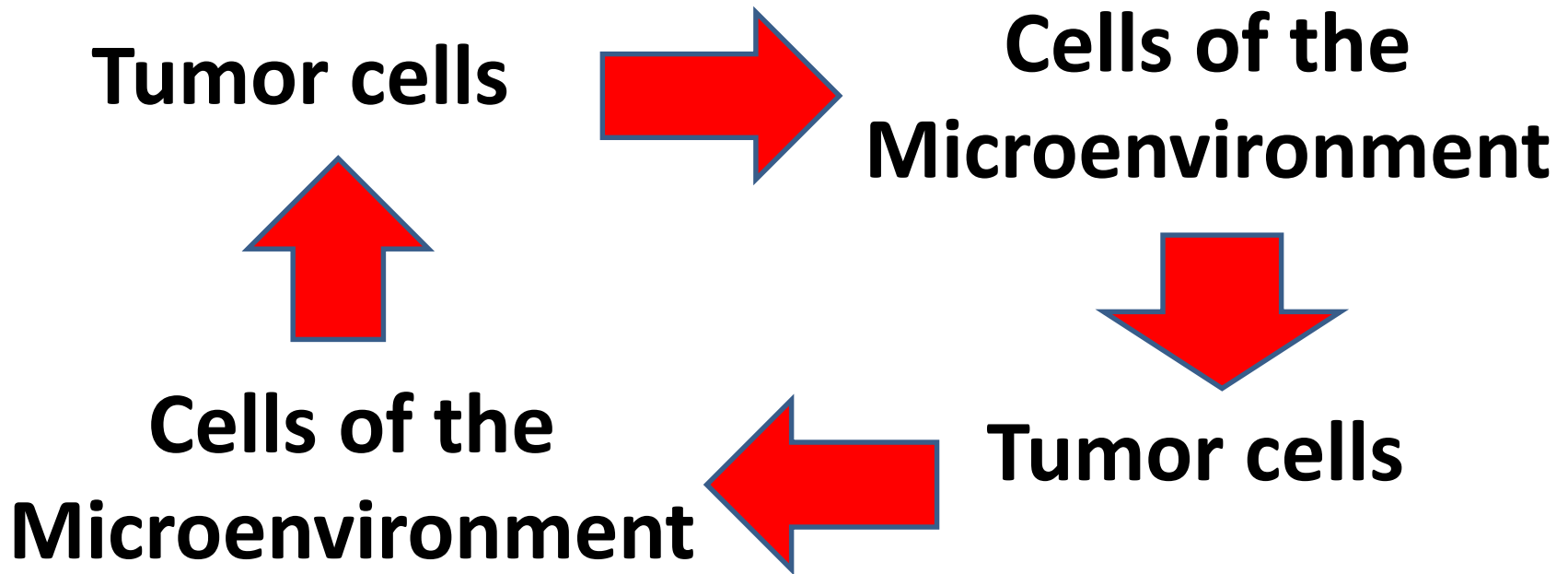
Loss of tumor suppressors and/or activation of oncogenes induces the inflammatory microenvironment



Cancer growth is shaped by Immunoediting: a dynamic interplay with the tumor microenvironment



A dynamic interplay



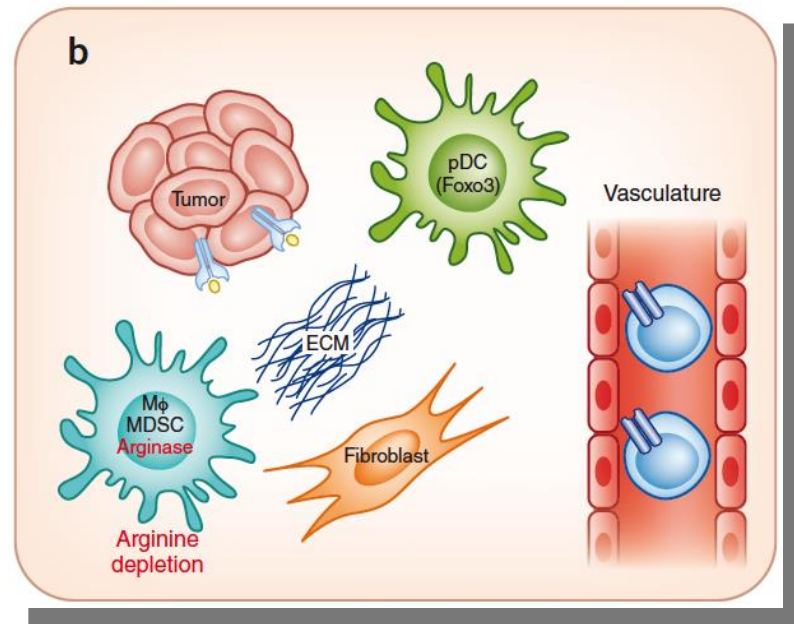
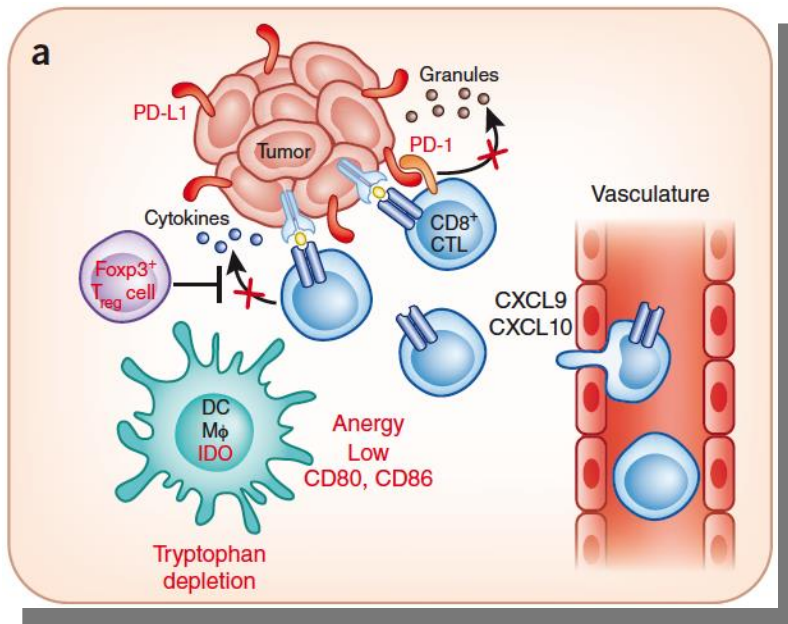
Hot tumors and cold tumors

Innate and adaptive immune cells in the tumor microenvironment

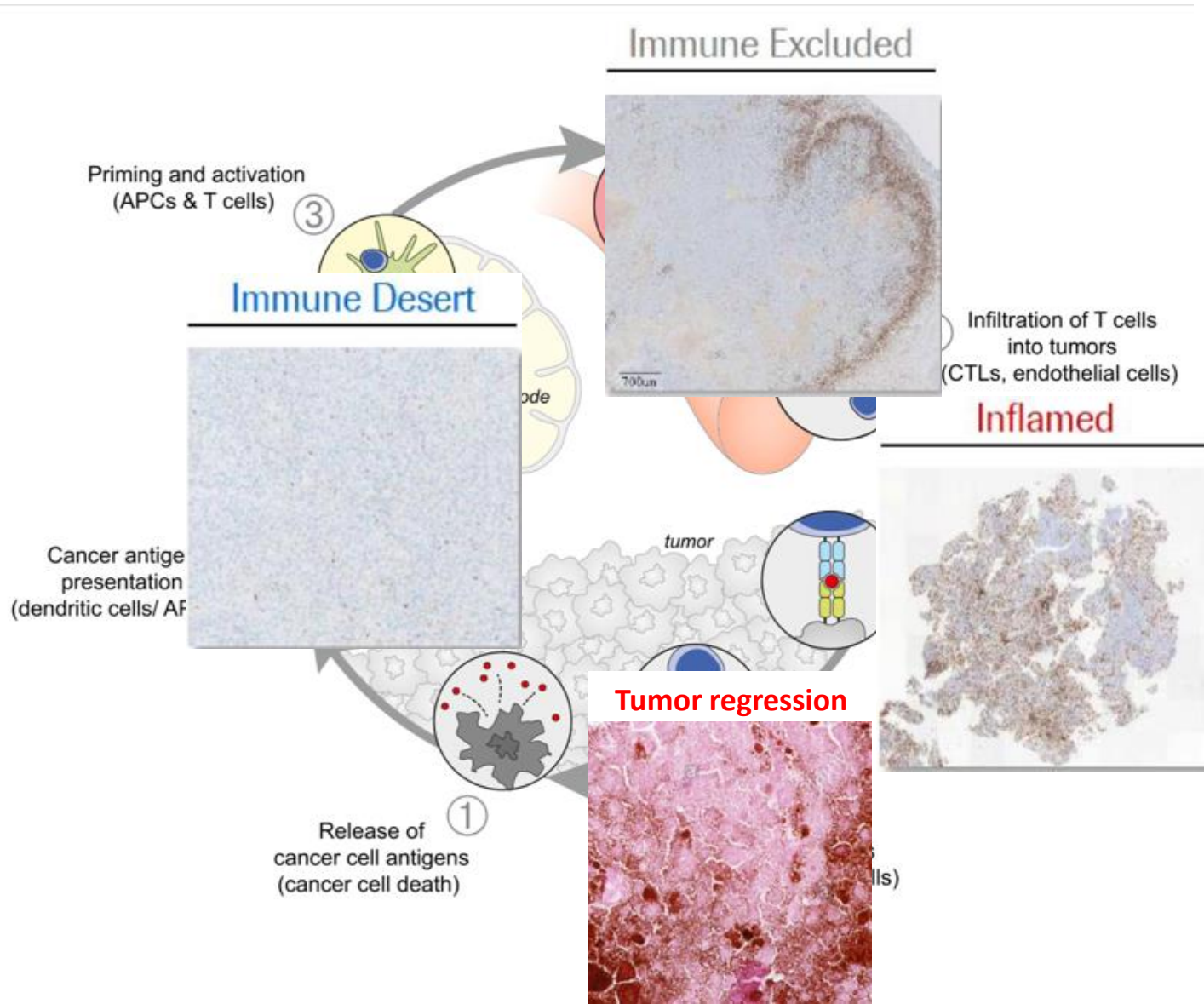
Thomas F Gajewski, Hans Schreiber & Yang-Xin Fu

inflamed tumor

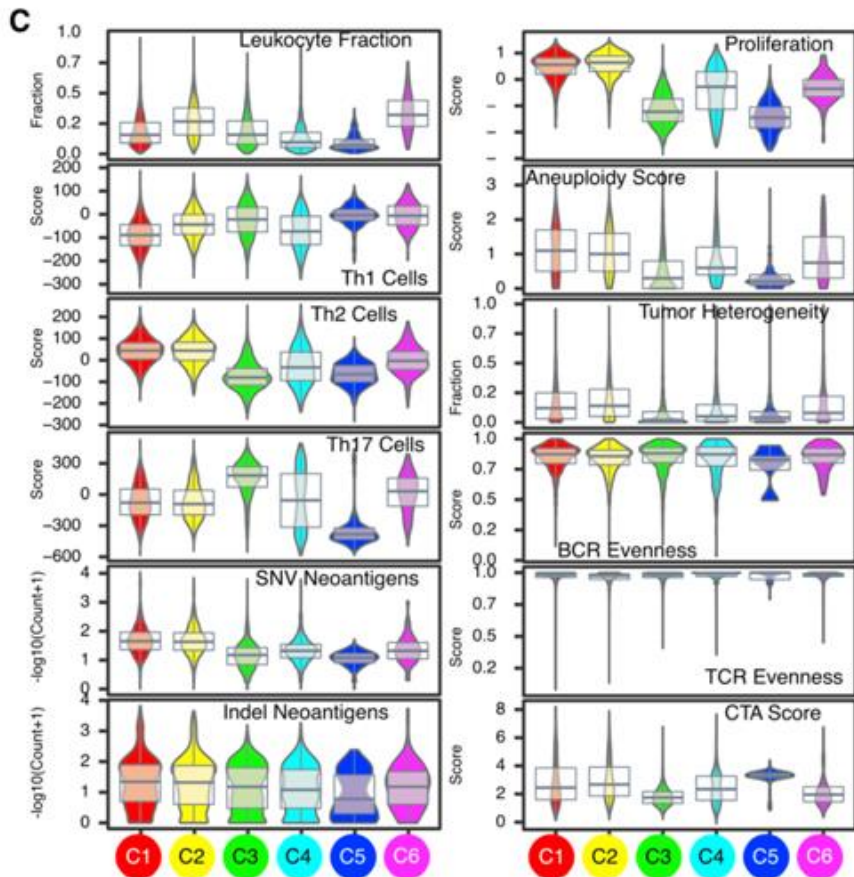
Non-inflamed tumor



Marina Corral Spence



Evolving technologies highlight higher levels of complexity



Immunity

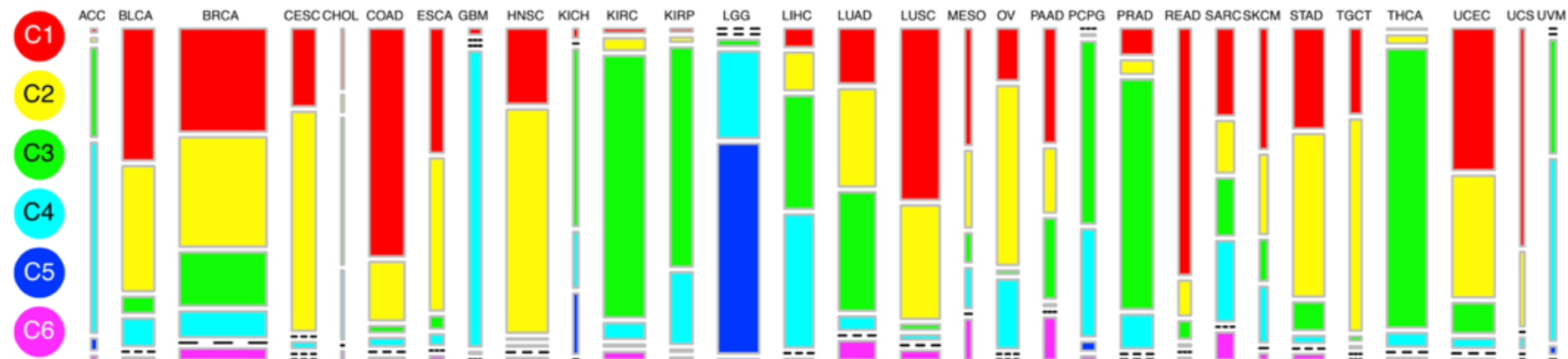
The Immune Landscape of Cancer

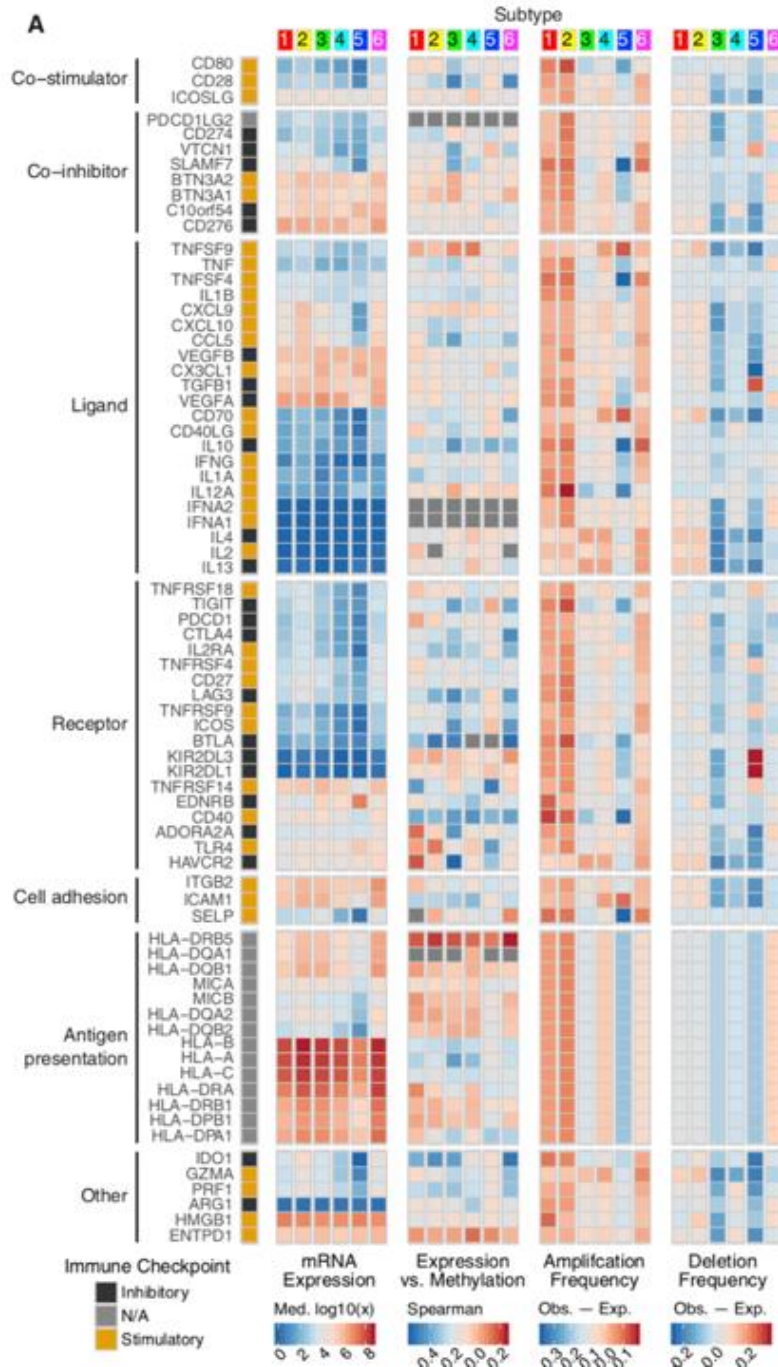
Thorsson et al. Immunity 2018

| | Macrophage: lymphocyte | Th1:Th2 | Proliferation | Intratumoral heterogeneity | Other |
|------------------------|------------------------|------------|---------------|----------------------------|------------------------------------|
| Wound healing | Balanced | Low | High | High | |
| IFN- γ dominant | Lowest | Lowest | High | Highest | Highest M1 and highest CD8 T cells |
| Inflammatory | Balanced | High | Low | Lowest | Highest Th17 |
| Lymphocyte depleted | High | Minimal Th | Moderate | Moderate | |
| Immunologically quiet | Highest | Minimal Th | Low | Low | Highest M2 |
| TGF- β dominant | High | Balanced | Moderate | Moderate | Highest TGF- β signature |

Evolving technologies highlight higher levels of complexity

| | Macrophage: lymphocyte | Th1:Th2 | Proliferation | Intratumoral heterogeneity | Other |
|----------------------------------|------------------------|------------|---------------|----------------------------|------------------------------------|
| C1 Wound healing | Balanced | Low | High | High | |
| C2 IFN- γ dominant | Lowest | Lowest | High | Highest | Highest M1 and highest CD8 T cells |
| C3 Inflammatory | Balanced | High | Low | Lowest | Highest Th17 |
| C4 Lymphocyte depleted | High | Minimal Th | Moderate | Moderate | |
| C5 Immunologically quiet | Highest | Minimal Th | Low | Low | Highest M2 |
| C6 TGF- β dominant | High | Balanced | Moderate | Moderate | Highest TGF- β signature |





Differential expression
of immunomodulatory
molecules among
different immunotypes

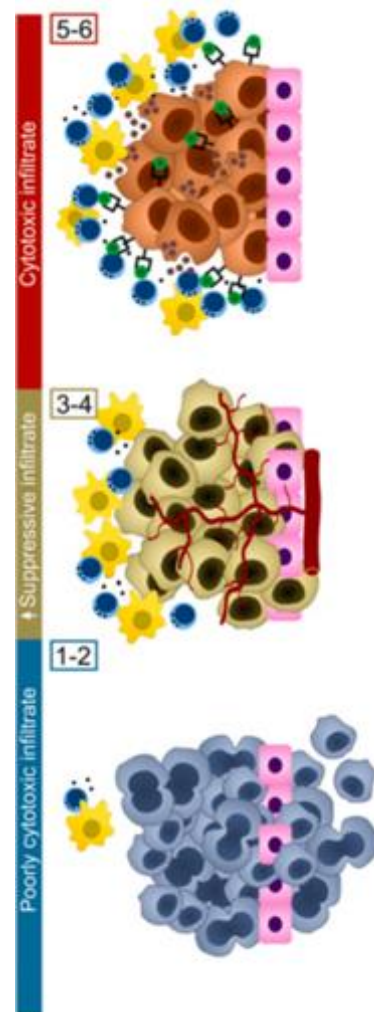
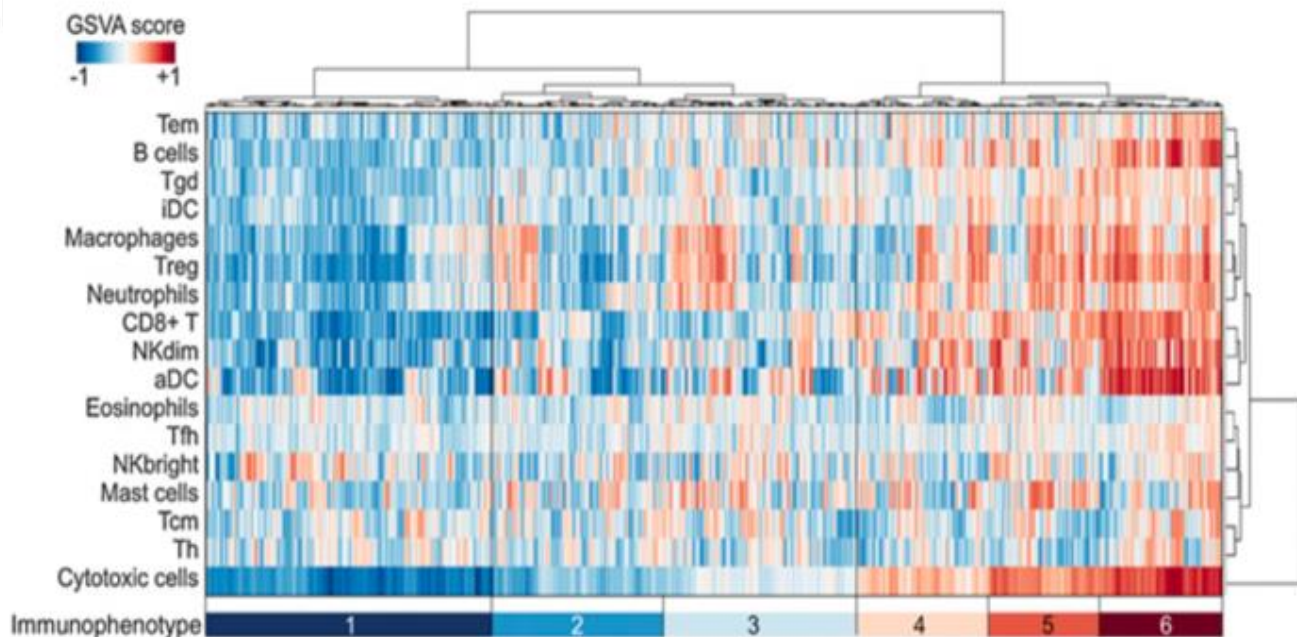
Evolving technologies highlight higher levels of complexity

Translational Cancer Mechanisms and Therapy

Clinical
Cancer
Research

A Pan-cancer Landscape of Interactions between Solid Tumors and Infiltrating Immune Cell Populations

David Tamboreno^{1,2}, Carlota Rubio-Perez^{1,2}, Ferran Muiños², Radhakrishnan Sabarinathan², Josep M. Piulats², Aura Muntasell⁴, Rodrigo Dienstmann^{5,6}, Nuria Lopez-Bigas^{1,2,7}, and Abel Gonzalez-Perez^{1,2}



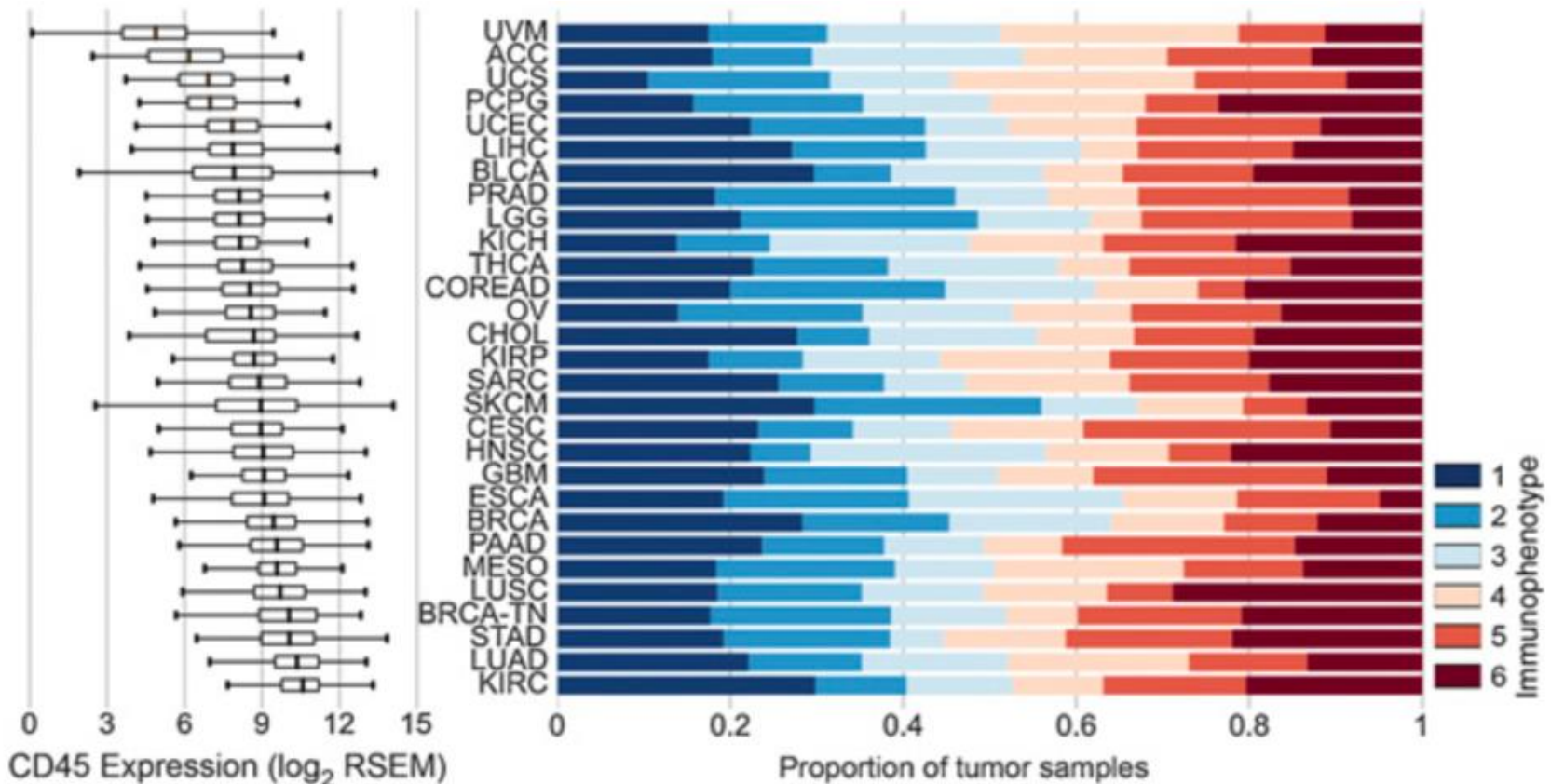
Evolving technologies highlight higher levels of complexity

Translational Cancer Mechanisms and Therapy

Clinical
Cancer
Research

A Pan-cancer Landscape of Interactions between Solid Tumors and Infiltrating Immune Cell Populations

David Tamborero^{1,2}, Carlota Rubio-Perez^{1,2}, Ferran Muiños², Radhakrishnan Sabarinathan², Josep M. Piulats³, Aura Muntasell⁴, Rodrigo Dienstmann^{1,6}, Nuria Lopez-Bigas^{1,2,7}, and Abel Gonzalez-Perez^{1,2}

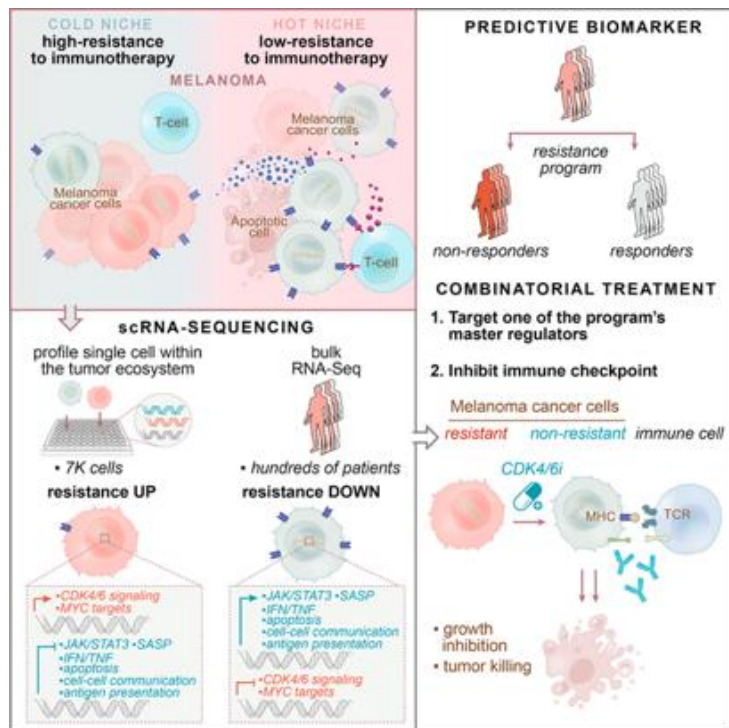


We are starting to understand the origin of microenvironment immunotypes

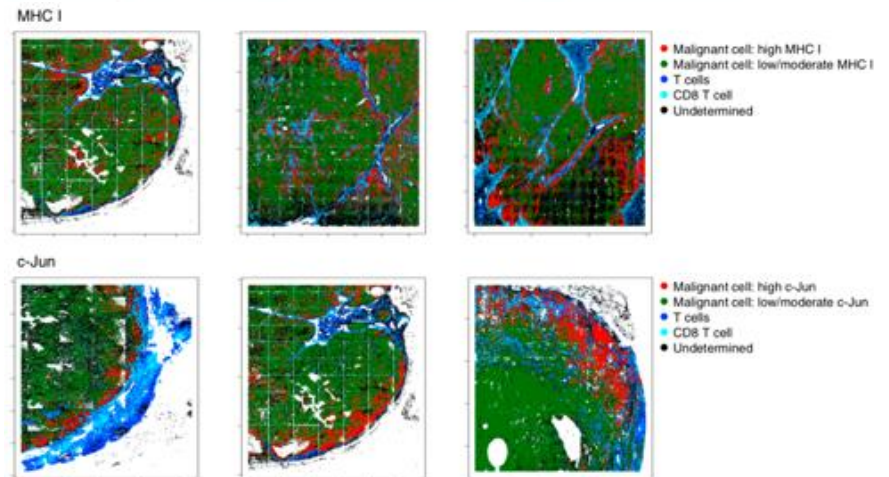


A Cancer Cell Program Promotes T Cell Exclusion and Resistance to Checkpoint Blockade

Jerby-Arnon et al Cell 175, 984–997, November 1, 2018 © 2018 Elsevier Inc.



Malignant Cells in T Cell-Depleted Niches Express Features of the Resistance Program *In Situ*



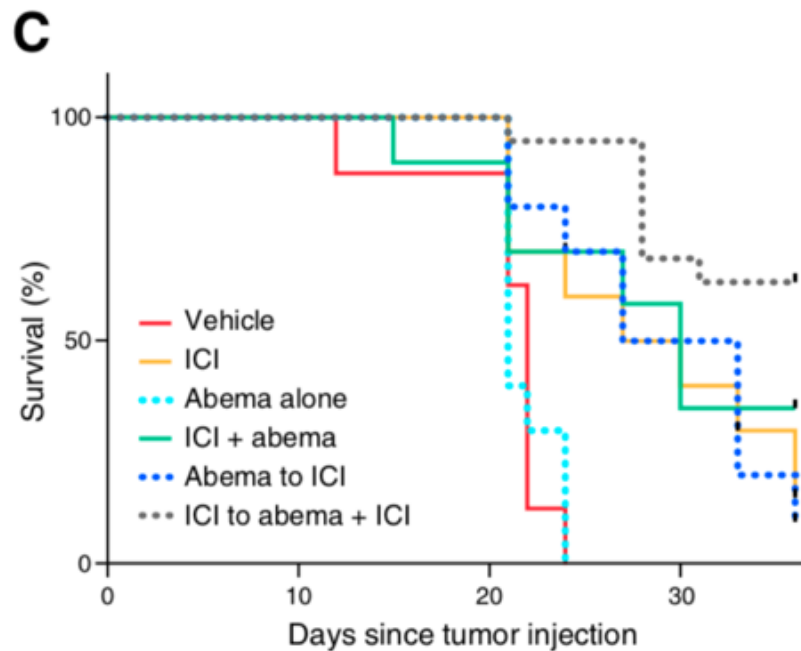
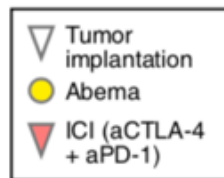
The Resistance Program Is Expressed Prior to Treatment and Is Enhanced following Immunotherapy in Resistant Lesions

The Resistance Program Predicts ICI Responses in Melanoma Patients

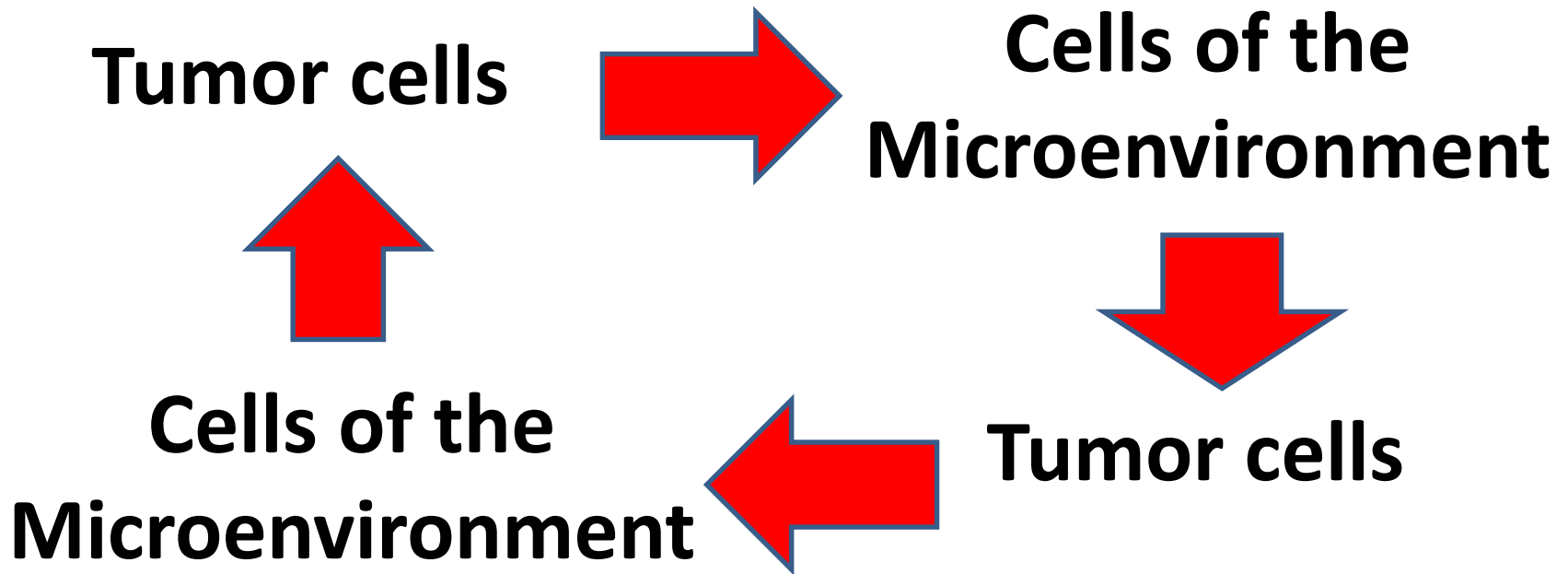
We are starting to understand the origin of microenvironment immunotypes

The Resistance Program Is Coherently Controlled by CDK4/6

CDK4/6 Inhibitors Repress the Resistance Program in Melanoma Cells

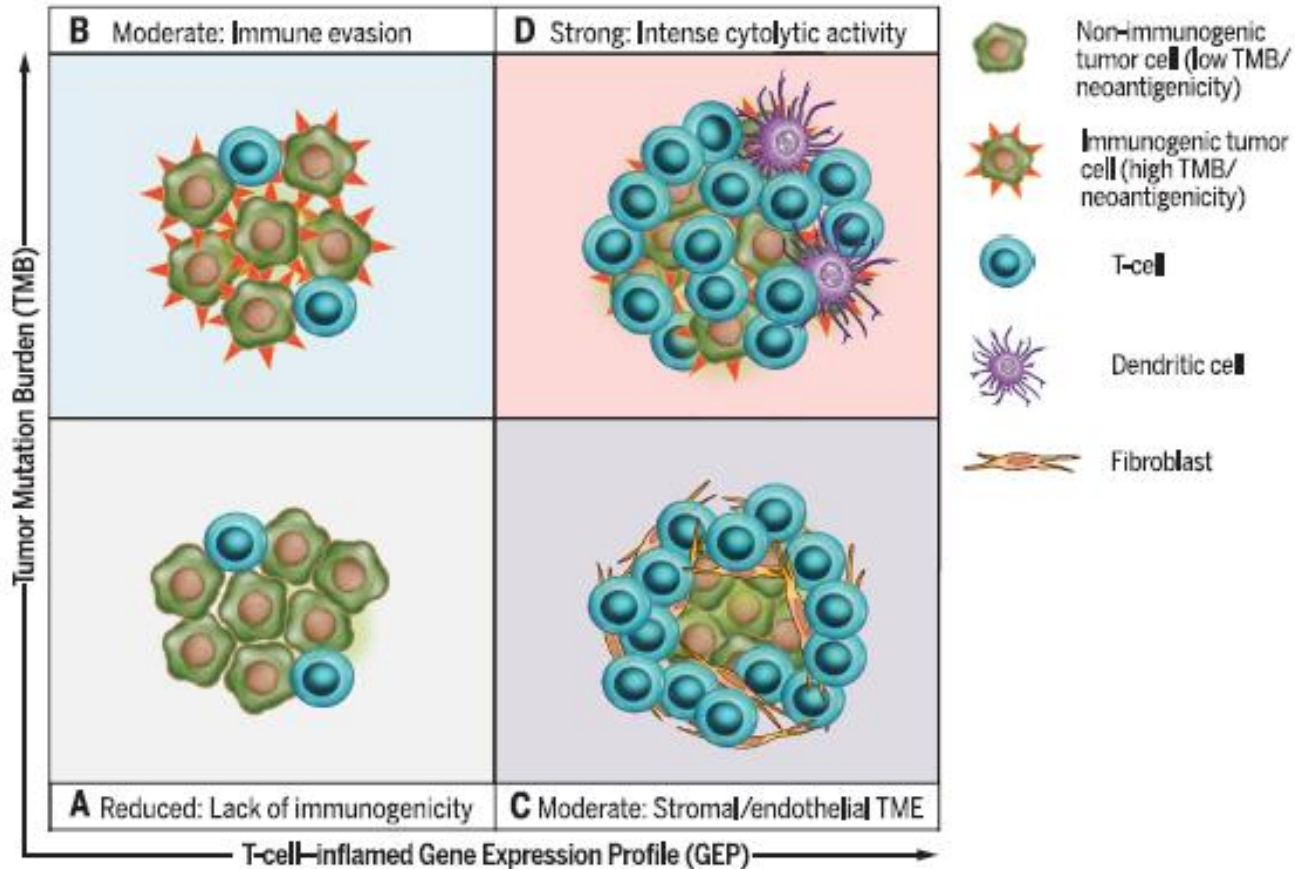


A dynamic interplay



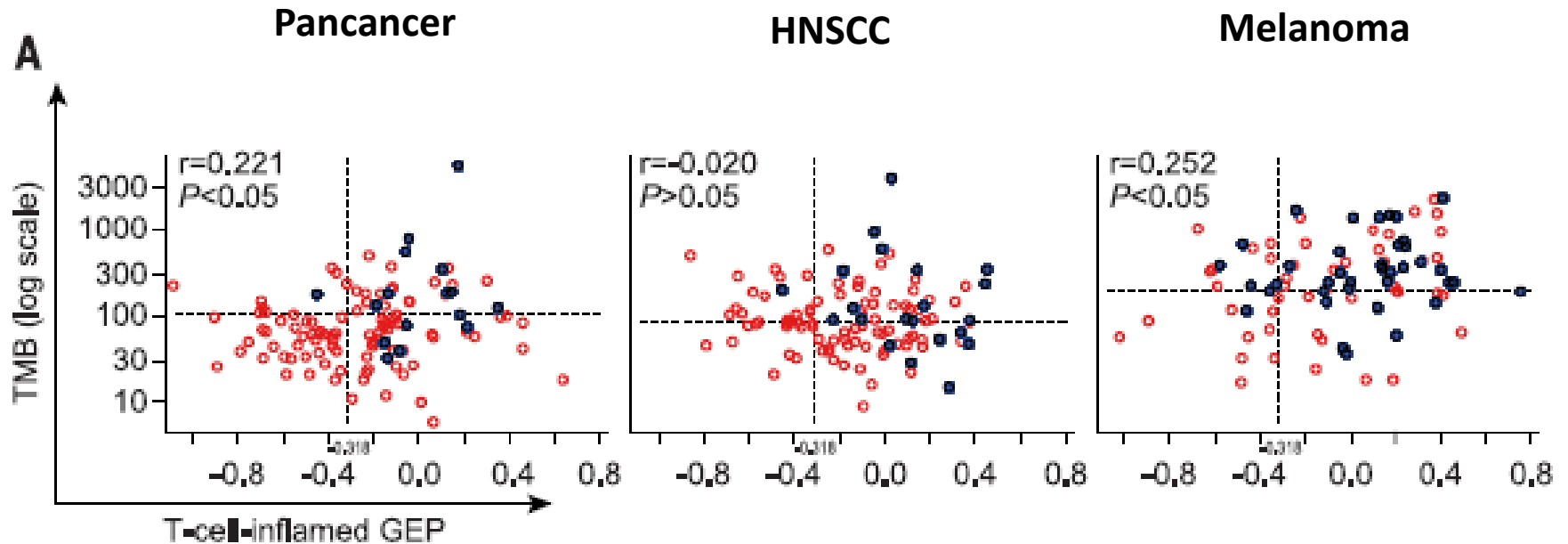
Potential diagnostic and therapeutic implications

Diagnostic Implications: combinations of biomarkers increases the predictive value of ICI



Biomarker-defined responses to pembrolizumab monotherapy identify targetable-resistance biology. (A) Tumors have low TMB and low neoantigenicity and lack a

Diagnostic Implications: combinations of biomarkers increases the predictive value of ICI

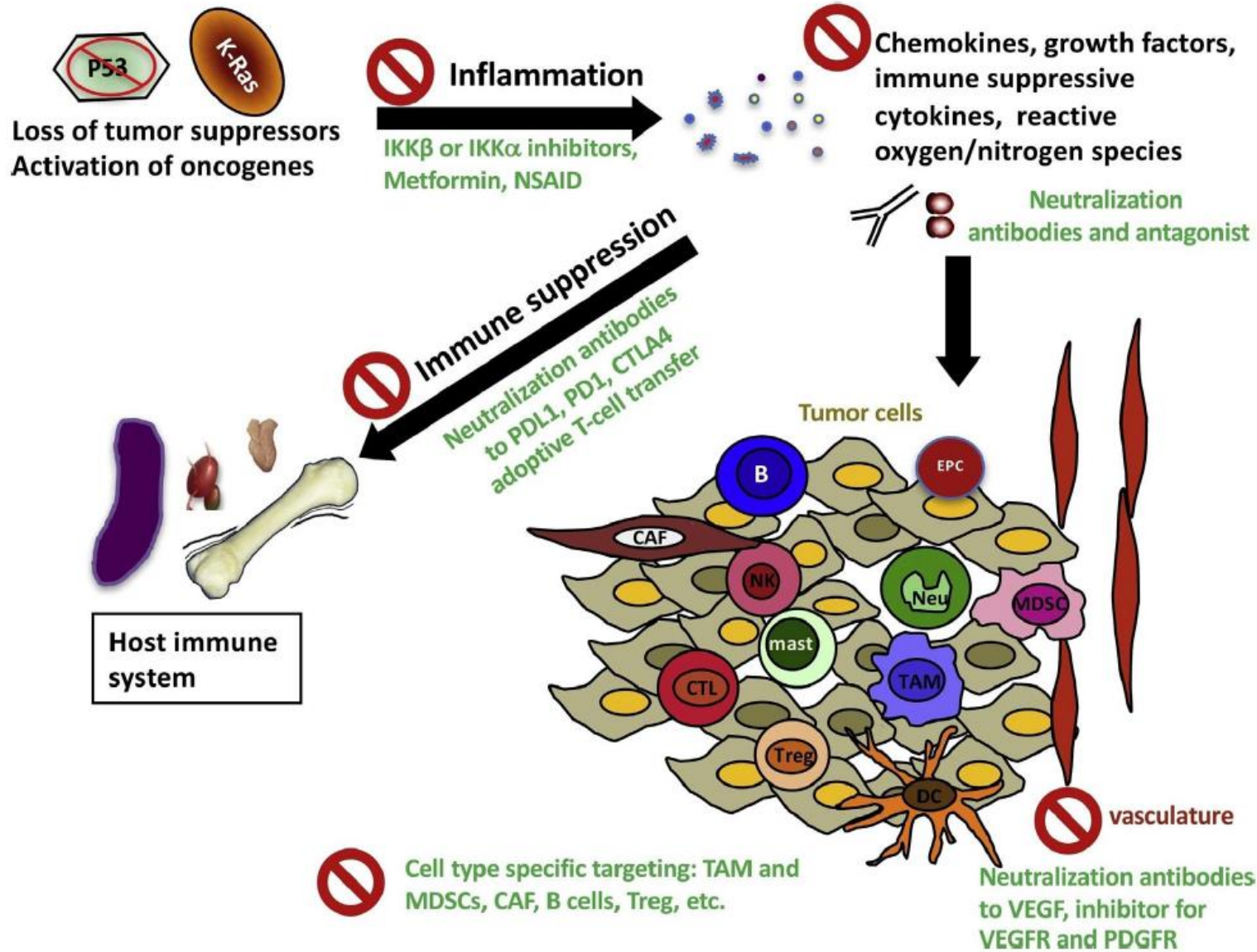


B

Alliance Against Cancer (ACC)

- ACC Immunotherapy
- Several centers involved
- Goal of the project: define a predictive signature of response to ICI through multiplex profiling of fast progressors vs long term responders

Therapeutic Implications



Suggested Reading

The evolving landscape of
biomarkers for checkpoint inhibitor
immunotherapy

Jonathan J. Havel ^{1,2,4}, *Diego Chowell*^{1,2,4} and *Timothy A. Chan* ^{1,2,3*}