

Ankersmit Laboratory

for Diagnosis & Regeneration In Thoracic Diseases & Applied Immunology

RESEARCH BRIEF

Analysis of Immune Signatures in Longitudinal Tumor Samples Yields Insight into Biomarkers of Response and Mechanisms of Resistance to Immune Checkpoint Blockade

Pei-Ling Chen^{1,2}, Whijae Roh¹, Alexandre Reuben³, Zachary A. Cooper^{1,3}, Christine N. Spencer¹, Peter A. Prieto³, John P. Miller¹, Roland L. Bassett⁴, Vancheswaran Gopalakrishnan³, Khalida Wani⁵, Mariana Petaccia De Macedo⁵, Jacob L. Austin-Breneman³, Hong Jiang³, Qing Chang¹, Sangeetha M. Reddy⁶, Wei-Shen Chen^{1,2}, Michael T. Tetzlaff², Russell J. Broaddus², Michael A. Davies⁷, Jeffrey E. Gershenwald³, Lauren Haydu³, Alexander J. Lazar^{2,5}, Sapna P. Patel⁷, Patrick Hwu⁷, Wen-Jen Hwu⁷, Adi Diab⁷, Isabella C. Glitza⁷, Scott E. Woodman⁷, Luis M. Vence⁸, Ignacio I. Wistuba⁵, Rodabe N. Amaria⁷, Lawrence N. Kwong⁵, Victor Prieto², R. Eric Davis⁹, Wencai Ma⁹, Willem W. Overwijk⁷, Arlene H. Sharpe¹⁰, Jianhua Hu⁴, P. Andrew Futreal¹, Jorge Blando⁵, Padmanee Sharma^{8,11}, James P. Allison⁸, Lynda Chin¹, and Jennifer A. Wargo^{1,3}

Cancer Discovery - Research Letter 08/2016



Hallmarks of cancer



Modified from Hanahan et al., Cell 2011



The cancer-immunity cycle





T cell response regulation

- Multiple co-stimulatory and inhibitory interactions for regulation
- "Checkpoint-Inhibitors"
- In lymph nodes or in peripheral tissue
- Bidirectional communication between APC/cancer cells and T cells



Pardoll et al., Nature Reviews 2012



Limitations

- Response rates 8-44%
- Those that do not response: severe AEs
- Costs: \$300.000 / patient (Ipi + Nivo Combo)



Andrews et al., Health Economics 2015 Boutros et al., Nature Reviews 2016 Schadendorf et al., J Clin Oncol 2015 Topalian et al., N Engl J Med 2012 Topalian et al., J Clin Oncol 2014 Wolchok et al., Ann Oncol 2013



Aim of the study

- Lack of predictive biomarkers
- Genomic, immune predictors investigated in pretreatment biopsies ¹⁻⁷
- CD8, CD4, PD1, PD-L1 densities in pretreatment biopsies ^{8,9}
- Mutational load, neoantigen structure ^{10, 11}
- Different effects of CIs on transcriptional profiles of TILs, JAK STAT ¹²

→ no robust results

 \rightarrow Need for comprehensive analyses of longitudinal tumor samples

¹Topalian et al., N Engl J Med 2012 ²Andtbacka et al., J Clin Oncol 2015 ³Larkin et al., N Engl J Med 2015 ⁴Postow et al., N Engl J Med 2015 ⁵Wolchok et al., N Engl J Med 2013 ⁶Topalian et al., J Clin Oncol 2014 ⁷Rizvi et al., Science 2015 ⁸Tumeh et al., Nature 2014 ⁹Taube et al., Clin Cancer Res 2014 ¹⁰Van Allen et al., Science 2015 ¹¹Snyder et al., N Engl J Med 2014 ¹²Gubin et al., Nature 2014



Patients and methods





Patients and methods

- Biopsies:
- Of the most safely accessible sites at different time points
- IHC: (88 FFPE)
- CD3, CD4, CD8, FOXP3, Granzyme B, CD57, CD20, CD45RO, LAG3, PD-1, PD-L1, CD14, CD33, CD68, CD163, CD206
- \rightarrow all were calculated by positive cells/mm²
- \rightarrow PD-1: H score (0-300) = +cells with membrane staining (%) / intensity of staining
- Immunofluorescence: (19 samples)
- DAPI, CD8, CD68 (potential myeloid-T cell interaction?)



Patients and Methods

- NanoString analyses: (54 samples following immune profiling)
- 1. FFPE \rightarrow RNA extraction (1µg tissue RNA / string assay)
- 2. Mixed with specific NanoString code set mix \rightarrow hybridization overnight
- 3. Loading on nCounter Prep Station (binding + washing)
- 4. Scanning / analyzing / data collection
- Gene expression profiling (GEP) analysis: 795-probe codeset
- Immune related genes; common cancer signaling related genes



Immune profiling in early on-treatment biopsies is predictive of response to CTLA-4 blockade







MEDICAL UNIVERSITY OF VIENNA

Immune profiling in early on-treatment biopsies is highly predictive of response to PD-1 blockade











Resuirs











+++



CD3

С







PD-1



D

Gene expression profiling in longitudinal tumor biopsies is predictive of response









NanoString paired analysis







Fold change



Results - Summary

- Early-on CTLA-4 blockade: CD8 (R vs NR)
- Early-on PD-1 blockade: CD8, CD4, CD3, PD-1, PD-L1, LAG3 (R vs NR)
- <u>GEP:</u>
- Pre-CTLA-4; Early-on CTLA-4; Pre-PD-1: no differences in R vs NR
- Early-On PD-1: >400 up-expressed genes in R vs NR
- Pre-PD-1 versus Early-on PD-1: 370 dynamic changes in gene expression
- **Paired biopsies:** (Pre-PD-1 versus Early-On PD-1)
- Stratification of patients in NR and R based on the the GEP-changes

IMMUNE SIGNATURES IN TUMOR BIOPSIES EARLY-ON TREATMENT ARE HIGHLY PREDICTIVE OF RESPONSE TO CIS



Discussion

- Only a fraction of patients benefits from immunotherapy
- Current approaches focus on assessing immune markers in pretreatment tissue
- Immune signatures in biopsies should be evaluated early after treatment initiation rather than in pre-treatment tissue
- Immune signatures in early on treatment rather a consequence of immune response to checkpoint-inhibitor than of therapeutic response?



Personal opinion

- Fluently written, easy to understand
- Clinical relevance? (Early on treatment?)
- Early-on biopsies 1.4 months too early to assess response?
- GEP (37 genes) in each patient during treatment?
- Dynamic GEP only in 13 patients?
- In MM easy to perform re-biopsy, but for NSCLC? And other tumors?
- Liquid biopsy?

