

EMERGING BIOMARKERS OF IMMUNE CHECKPOINT INHIBITORS

Rodrigo Dienstmann



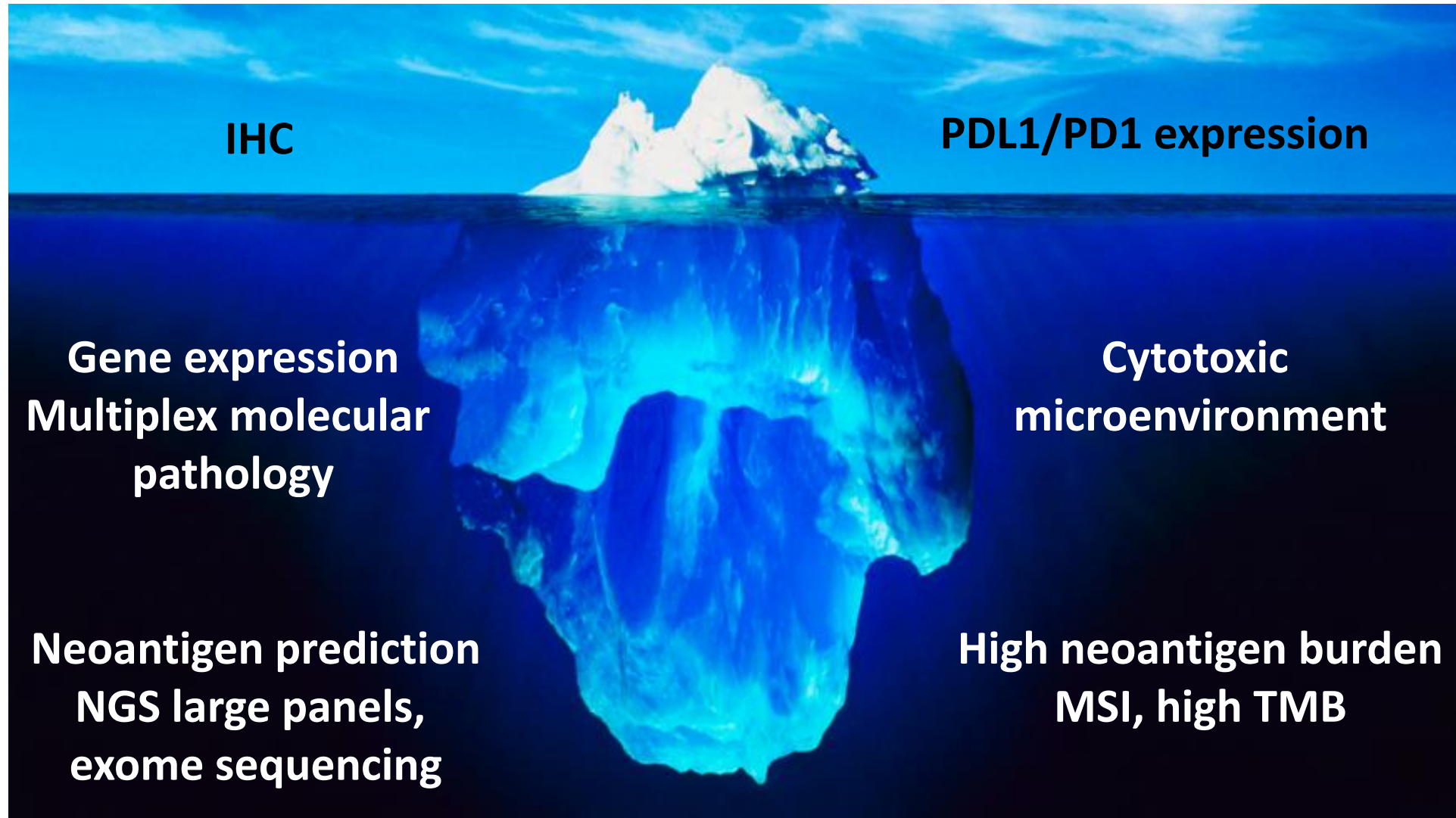
(PERSONAL) CONFLICTS OF INTEREST

Advisory role: Roche, Foundation Medicine
Boehringer-Ingelheim

Speaker's fee: Roche
Amgen
Ipsen
Servier
Sanofi
MSD

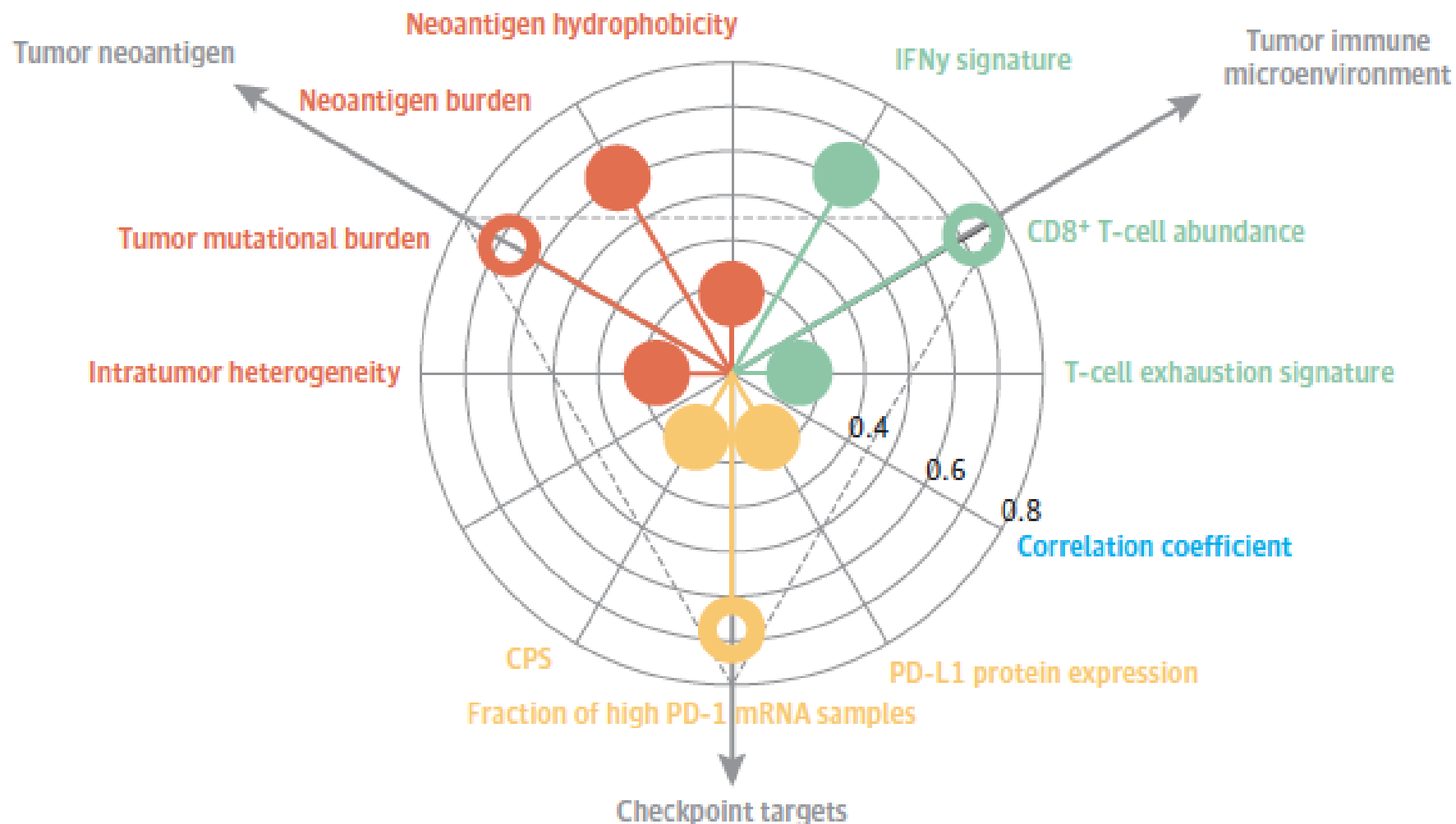
Research grant: Merck
Pierre Fabre
BMS

INTEGRATED BIOMARKER ANALYSIS



MULTI-OMICS PREDICTIVE MARKERS

Triple axis of anti-PD-1/PD-L1 response



MULTI-OMICS PREDICTIVE MARKERS

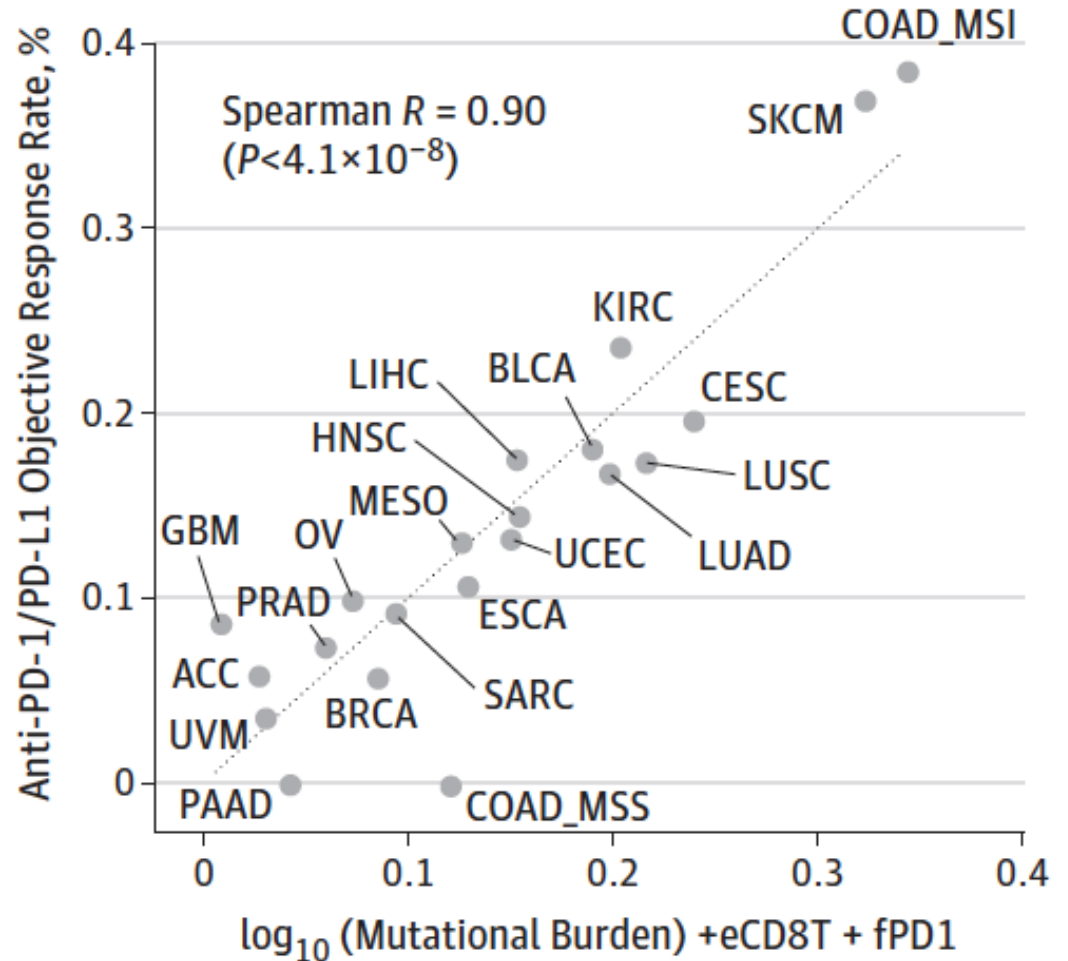


In-silico analysis

TCGA data vs. results clinical trials

Spearman $R = 0.9$

Model with **PDL1 + CD8 + TMB**
explains **80%** of **ORR** across tumor types



MULTI-OMICS PREDICTIVE MARKERS

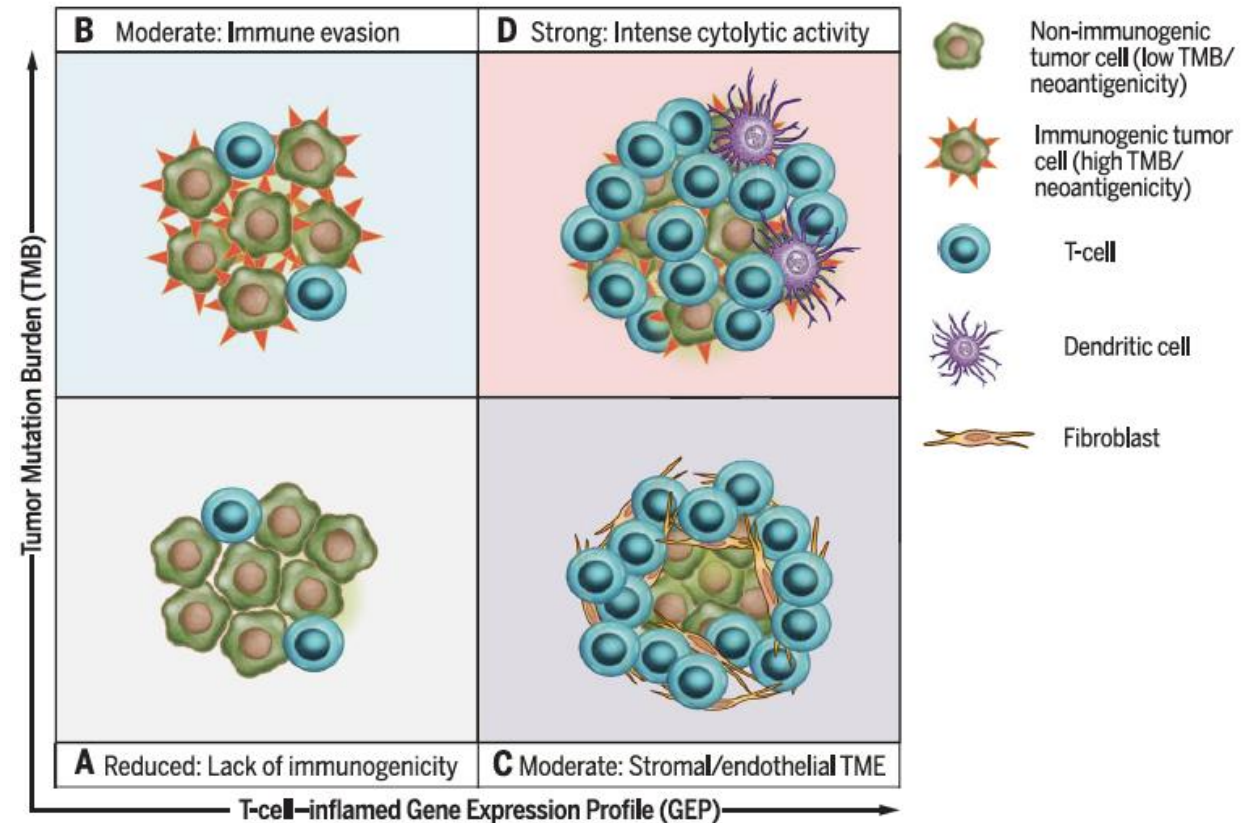


Metanalysis KEYNOTE trials

300 patients, 22 tumors

TMB

GEP cytotoxicity signature (18 genes)



MULTI-OMICS PREDICTIVE MARKERS

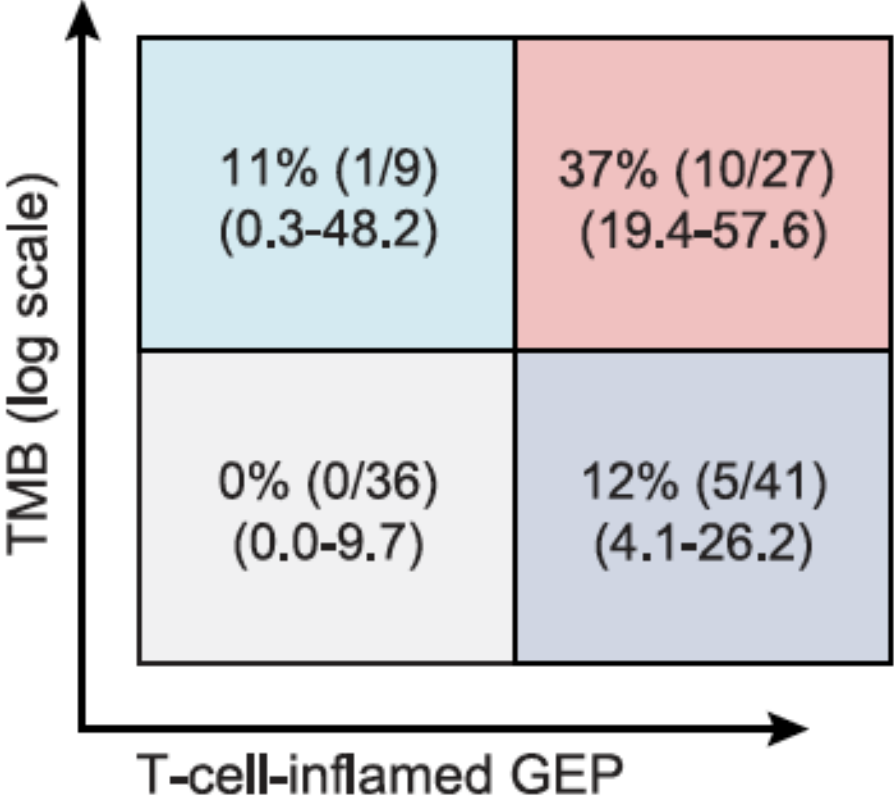


Metanalysis KEYNOTE trials

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 TMB^{lo} GEP^{lo}

 TMB^{lo} GEP^{hi}

 TMB^{hi} GEP^{lo}

 TMB^{hi} GEP^{hi}

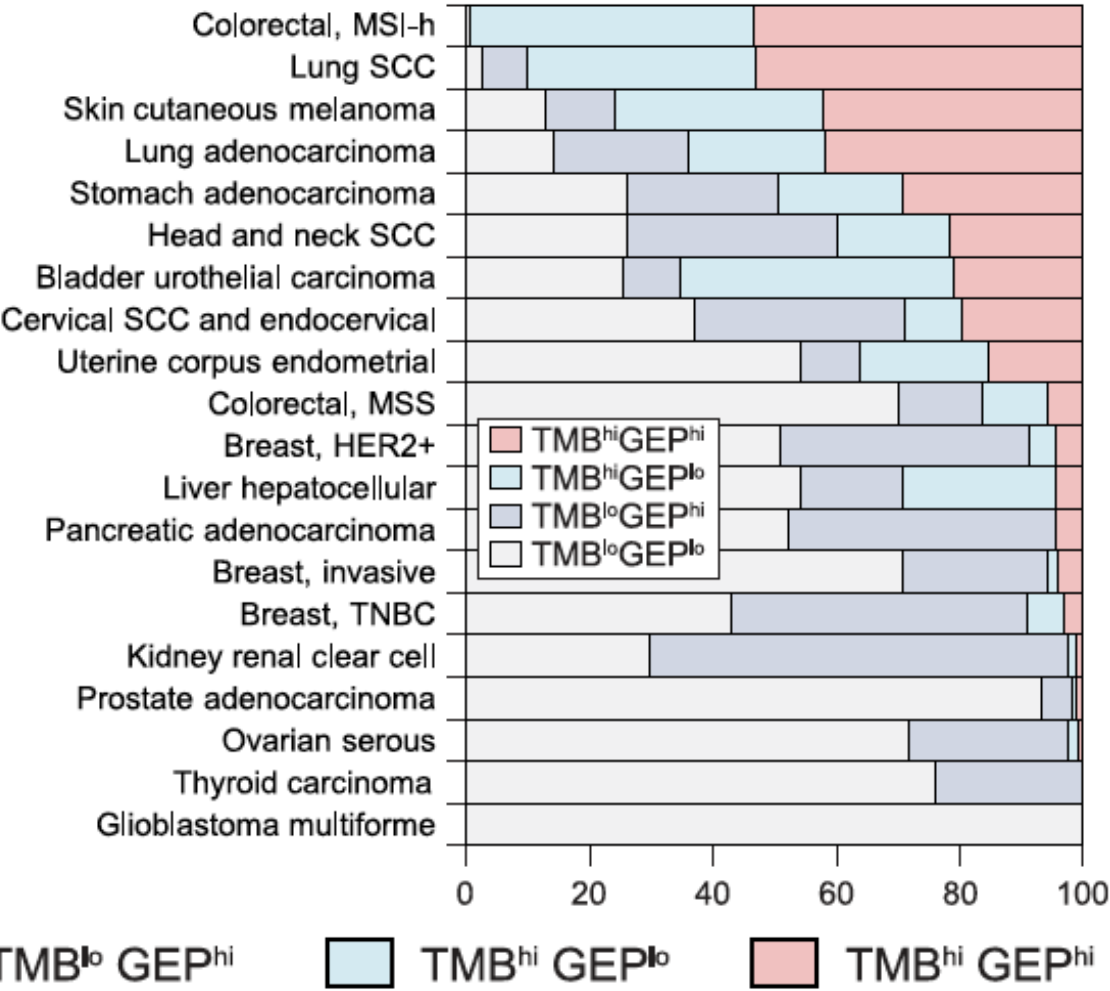
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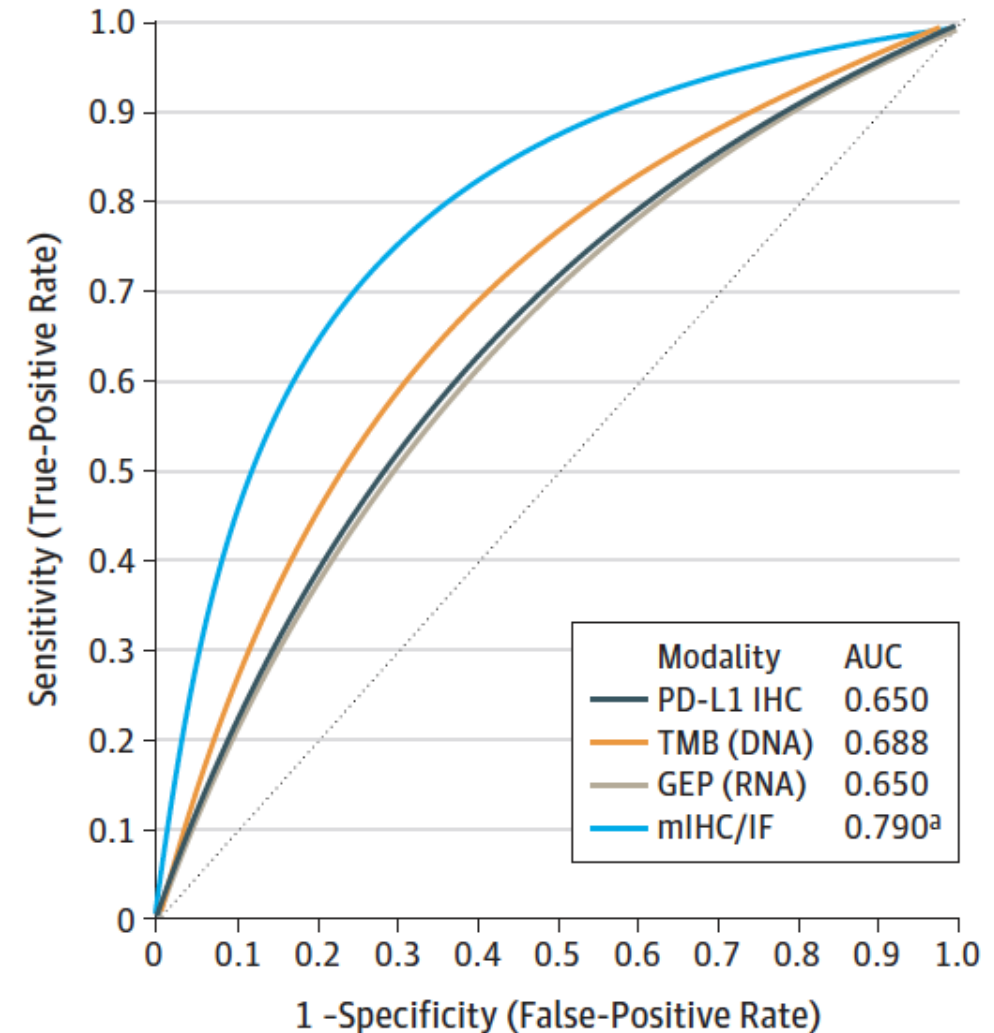


MULTI-OMICS PREDICTIVE MARKERS



Systematic review and metanalysis
45 clinical trials

PD-L1 IHC
TMB
GEP cytotoxicity signature
Multiplex IHC/IF: quantitative CD8/PD-L1



MULTI-OMICS PREDICTIVE MARKERS



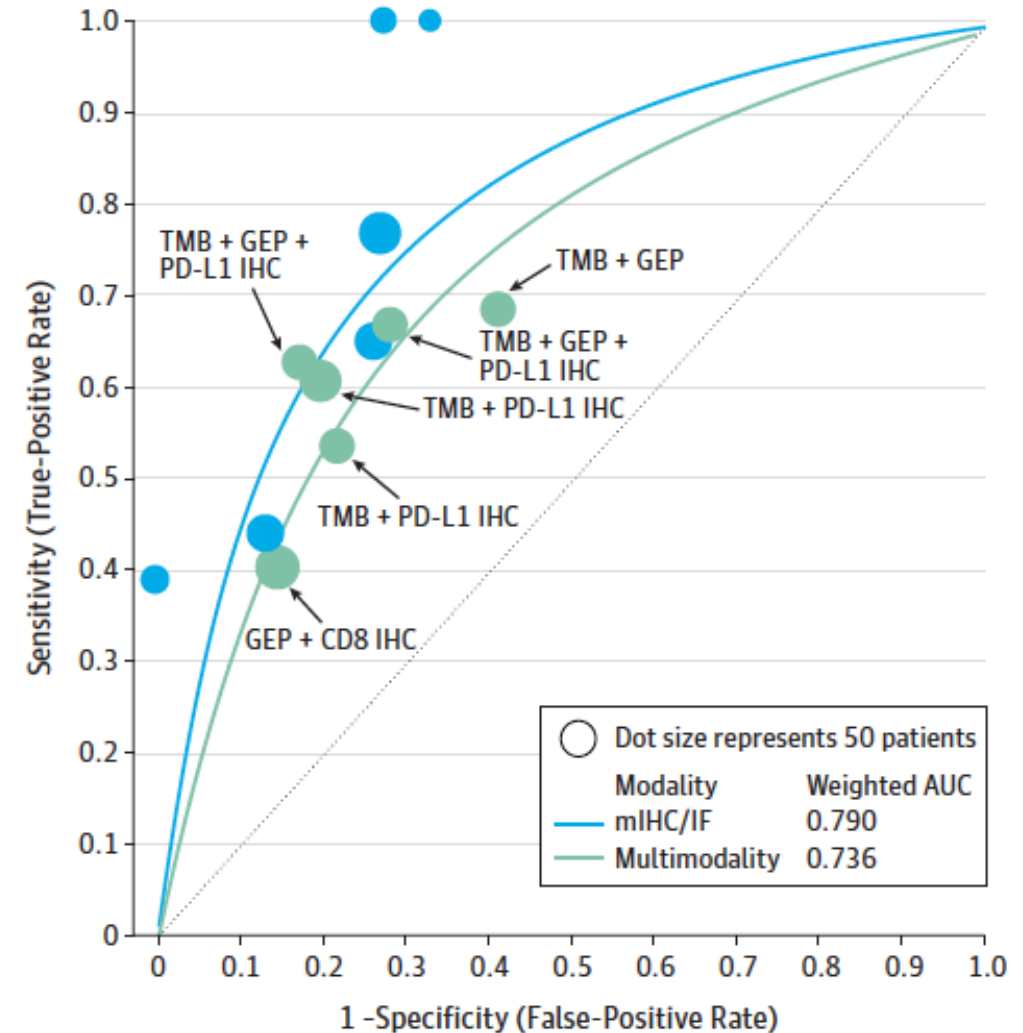
Systematic review and metanalysis 45 clinical trials

PD-L1 IHC

TMB

GEP cytotoxicity signature

Multiplex IHC/IF: quantitative CD8/PD-L1



GENOMIC MARKERS



Response

Clonal mutations, viral neoantigens

Hypermutation – MSI, *POLE* mut, high TMB, DDR alt

Upregulation of *PDL1* (ampl)

Epigenetic events (*PBMR1*, *SMARCA4*, *ARID1A*, *ARID2*...)

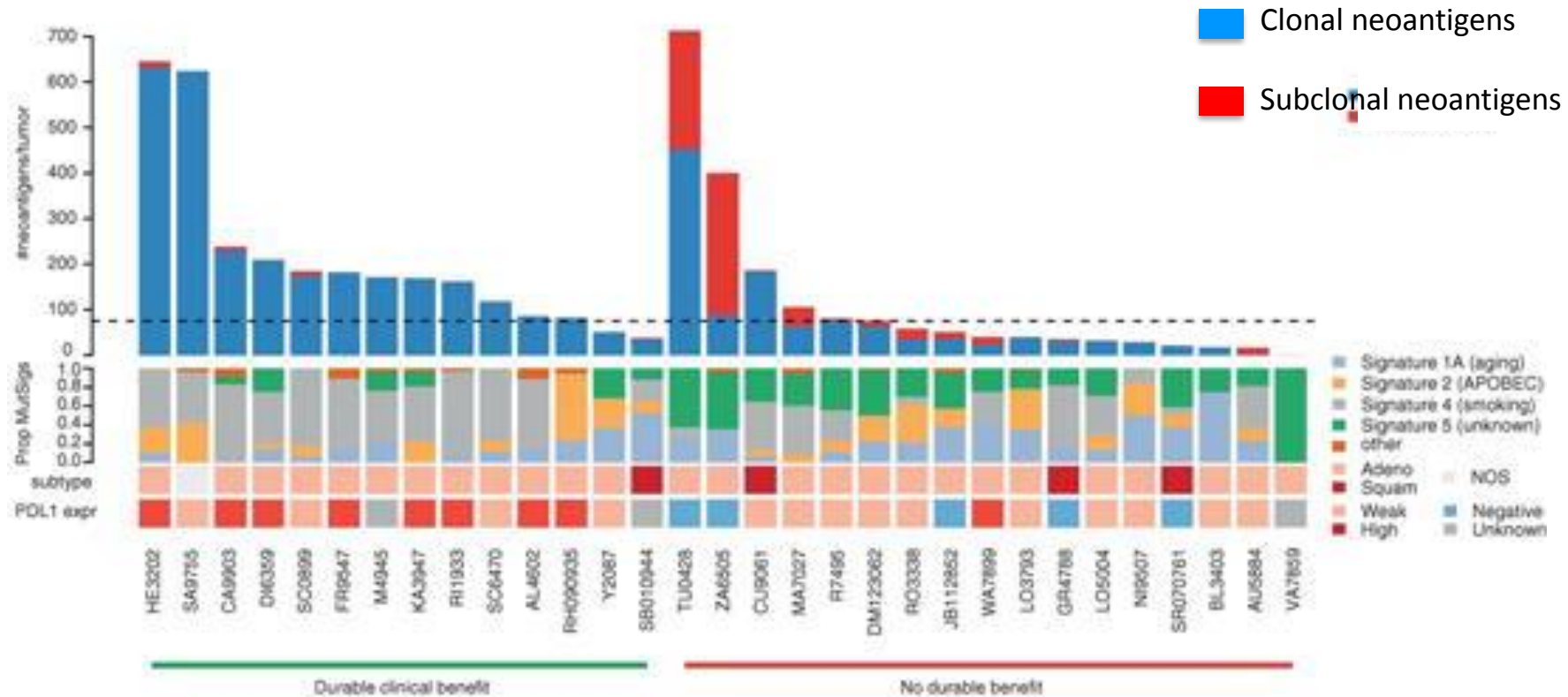
Resistance

Single gene alterations – *STK11* mut, *PTEN* loss, *CDK4/CCND1* ampl

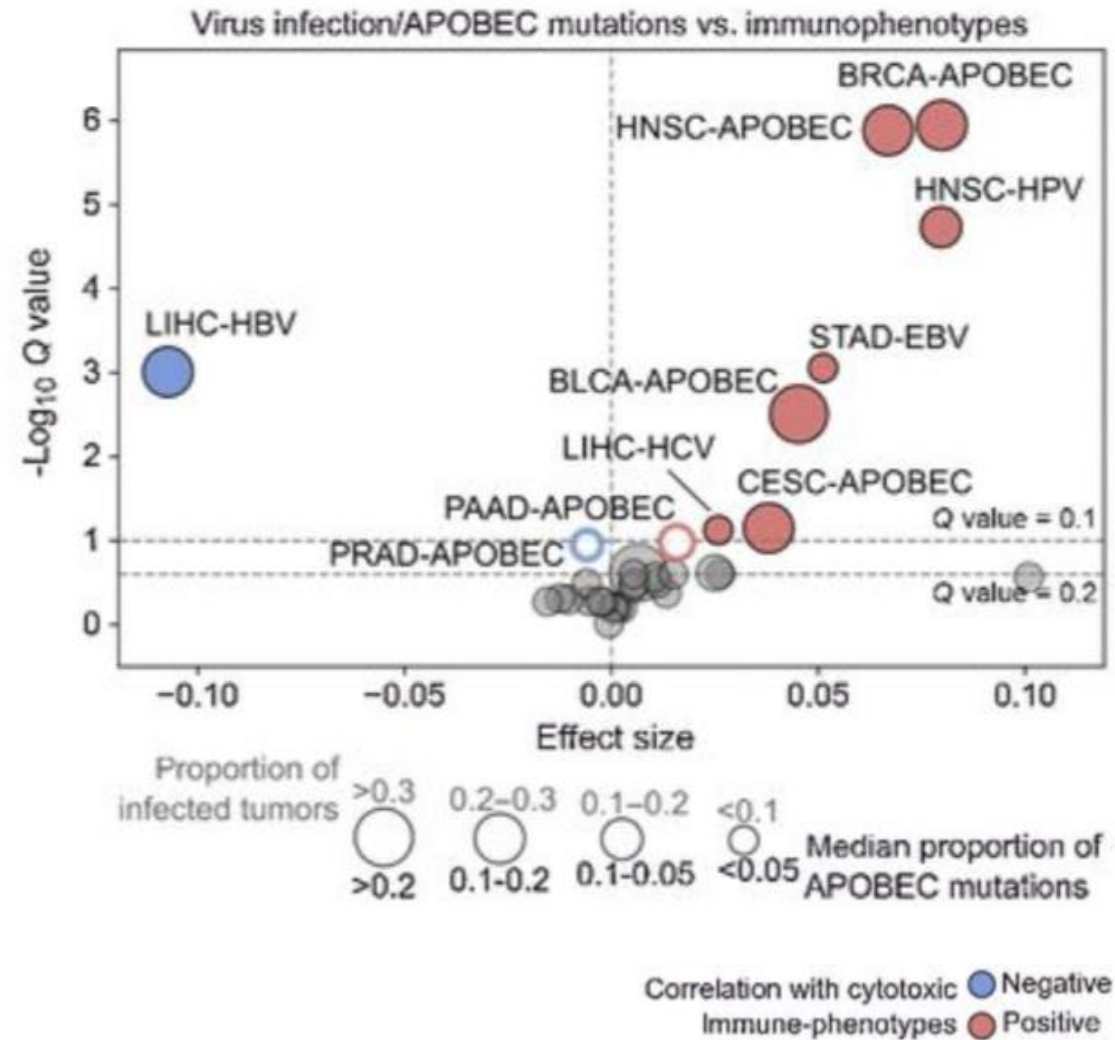
High levels of copy number loss

Alterations in Interferon and antigen presentation signaling

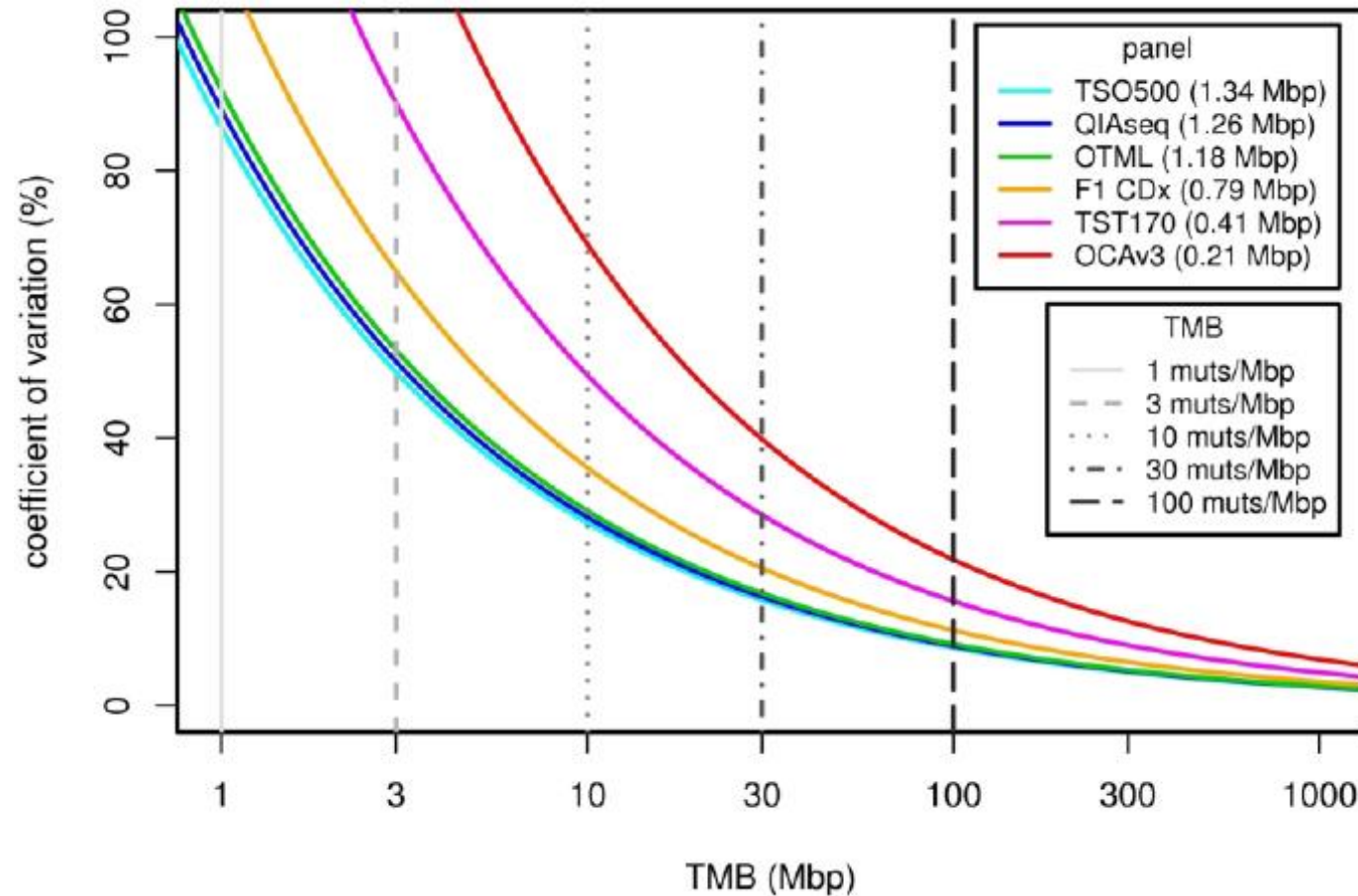
CLONAL NEOANTIGENS IN LUNG CANCER



VIRAL NEOANTIGENS AND APOBEC MUTATION SIGNATURE



OPTIMIZING PANEL-BASED TMB MEASUREMENT



~ 33% misclassification rate
across commercial panels

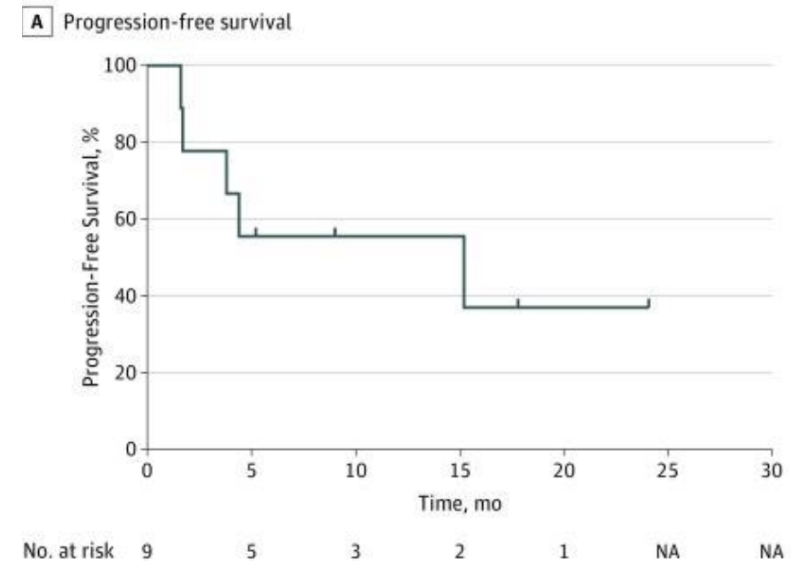
Important aspects:

- Cut-offs (tumor type/ setting)
- Source (tissue, ctDNA)
- Harmonization
- Prognostic vs. predictive (OS gain)

***PDL1* amplification in solid tumors**

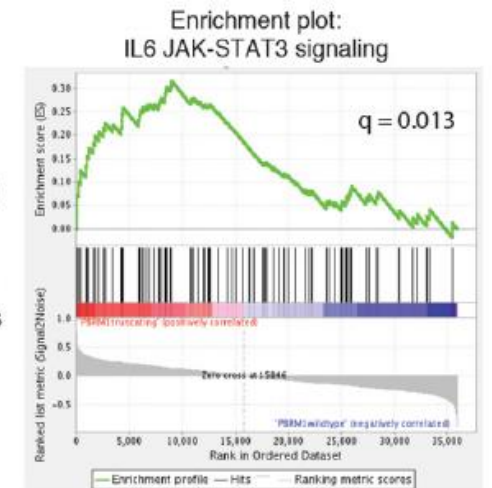
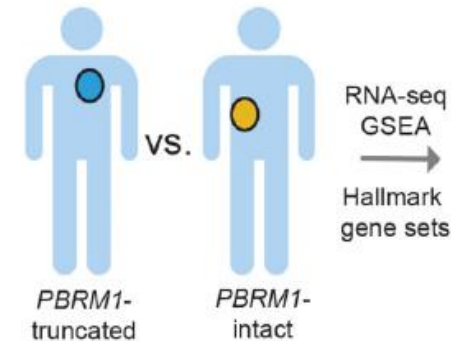
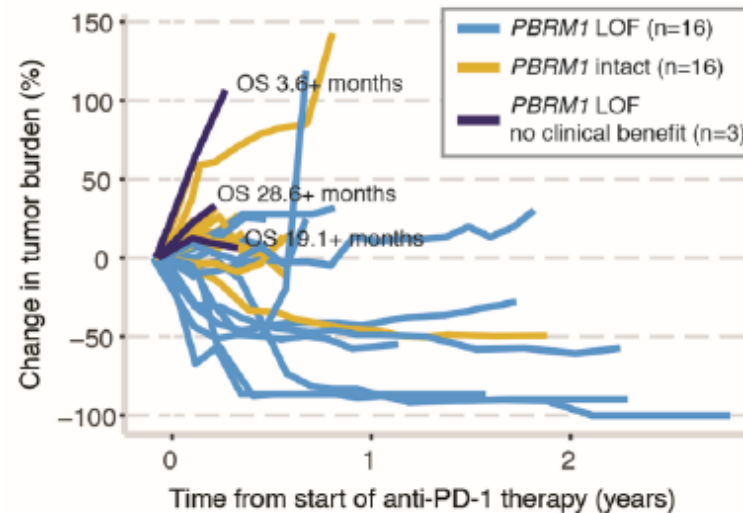
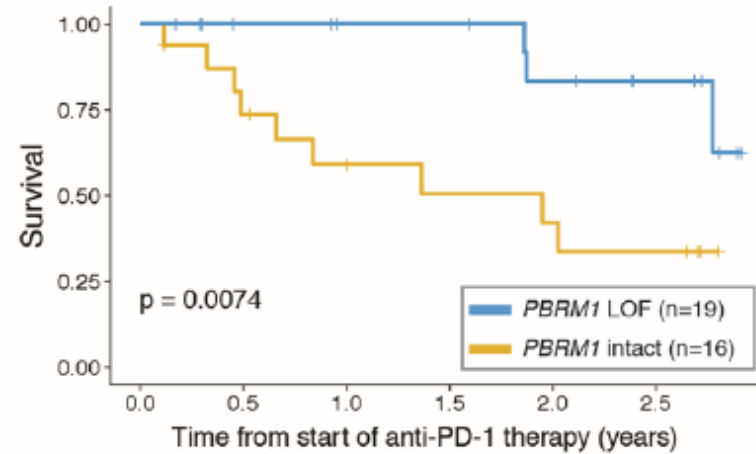
ORR 67%

Goodman et al, JAMA Oncol 2018



EPIGENETIC ALTERATIONS IN RENAL CANCER

SWI/SNF alterations:
PBRM1 mut



EPIGENETIC ALTERATIONS ACROSS TUMORS



ARID1A deficiency promotes mutability and potentiates therapeutic antitumor immunity unleashed by immune checkpoint blockade

ARID1A mut

Nature Medicine 2018

A major chromatin regulator determines resistance of tumor cells to T cell-mediated killing

PBMR1 mut, *ARID2* mut

Science 2018

Immune-Active Microenvironment in Small Cell Carcinoma of the Ovary, Hypercalcemic Type: Rationale for Immune Checkpoint Blockade

SMARCA4 mut

JNCI 2018

GENOMIC MARKERS



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Clonal mutations, viral neoantigens

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Upregulation of *PDL1* (ampl)

Epigenetic events (*PBMR1*, *SMARCA4*, *ARID1A*, *ARID2*...)

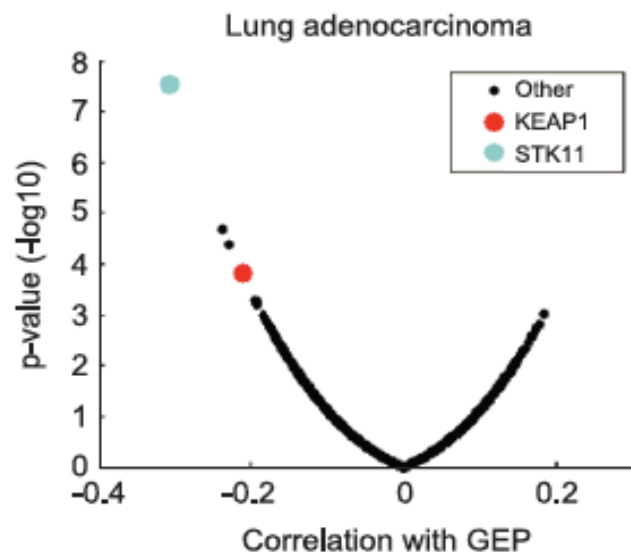
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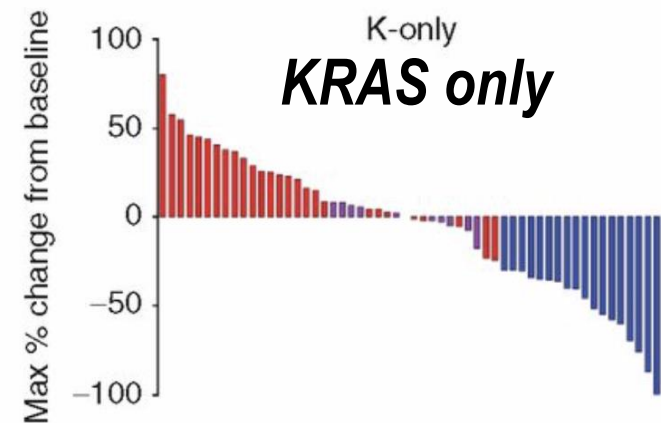
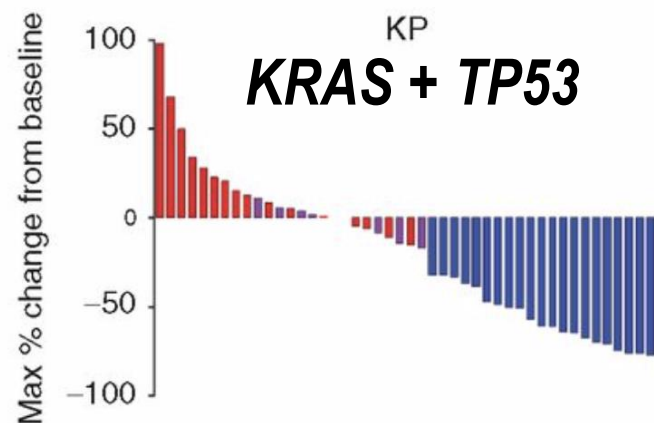
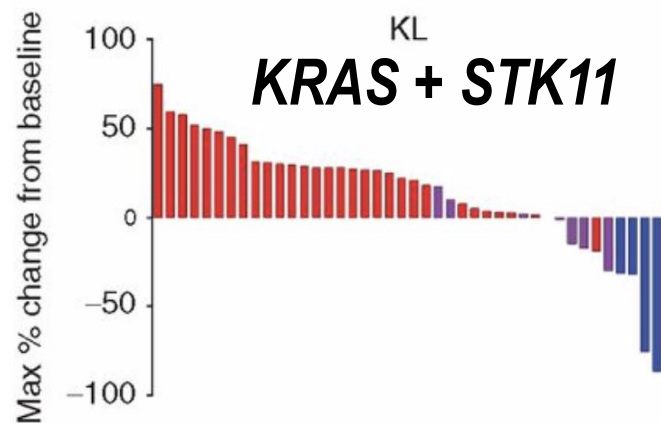
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STK11 MUTATIONS IN LUNG CANCER



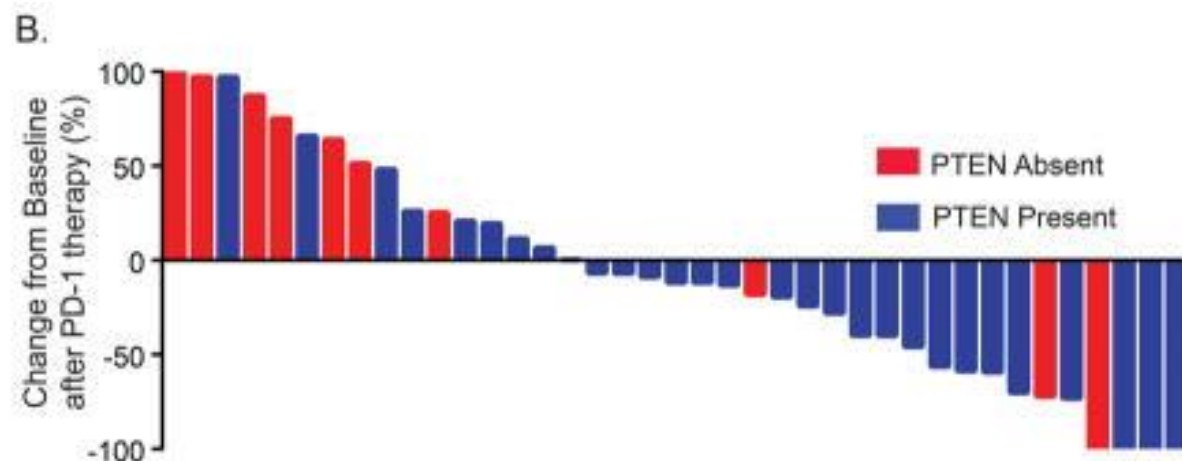
Cristescu et al, Science 2018



Skoulidis et al, Cancer Discov 2018

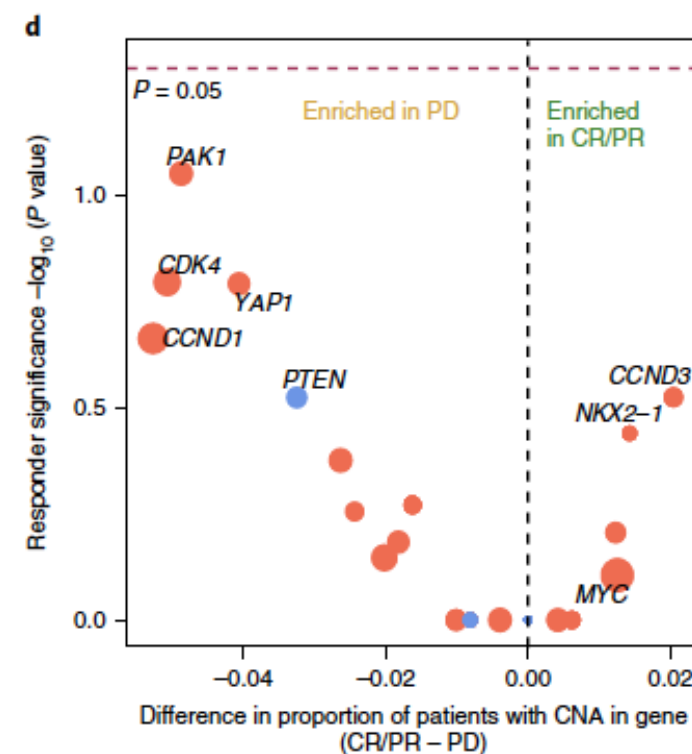
PTEN LOSS AND CDK4/CCND1 AMPL

Melanoma



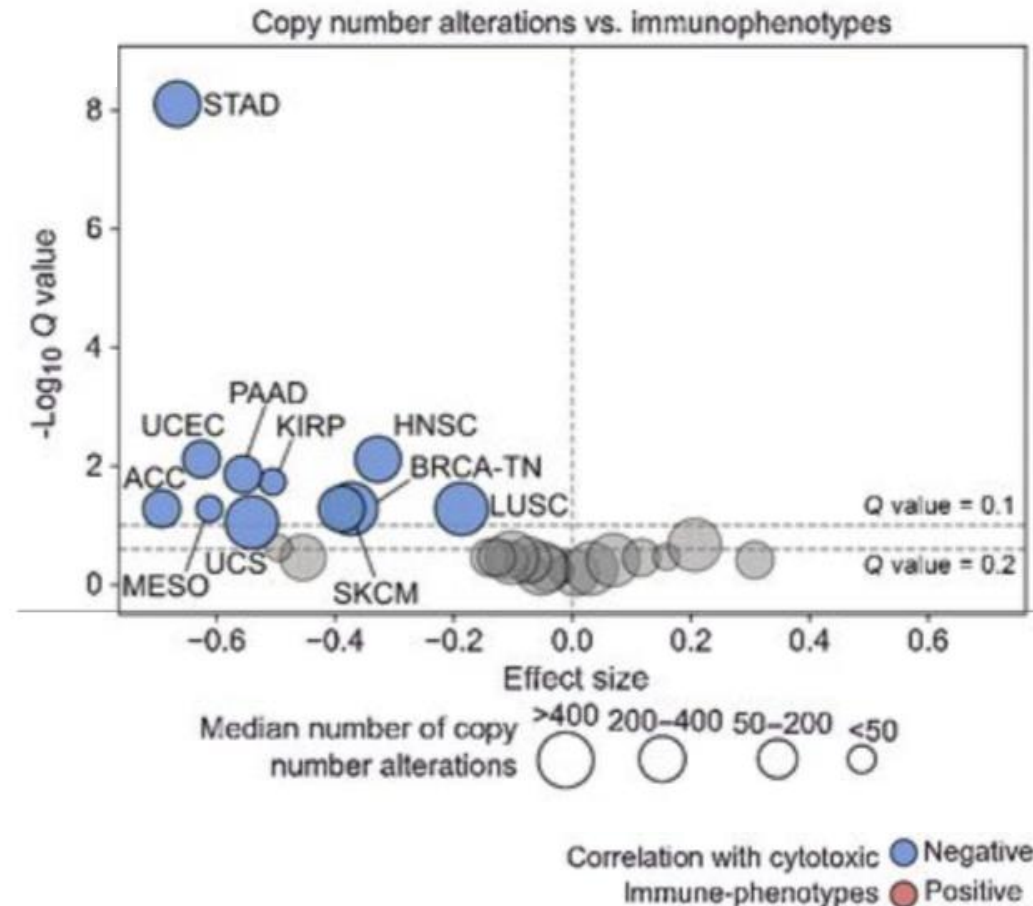
Peng et al, Cancer Discov 2016

Pan-tumor

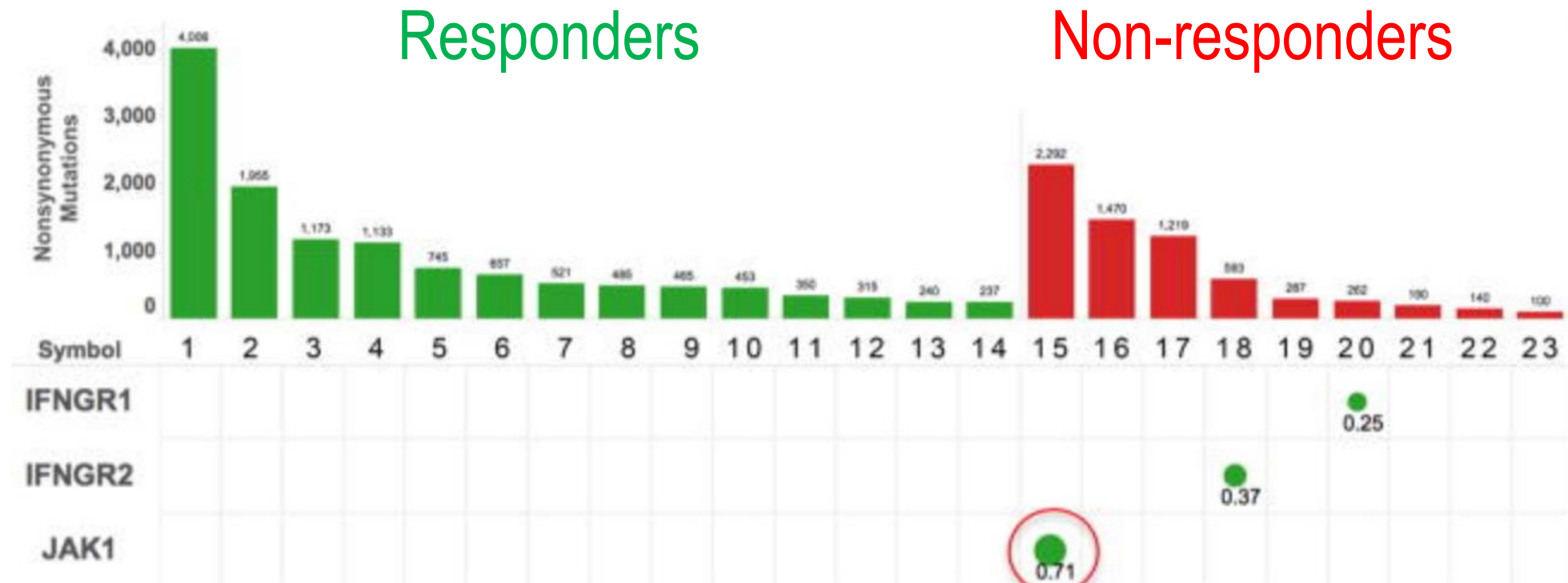


Miao et al, Nat Genet 2018

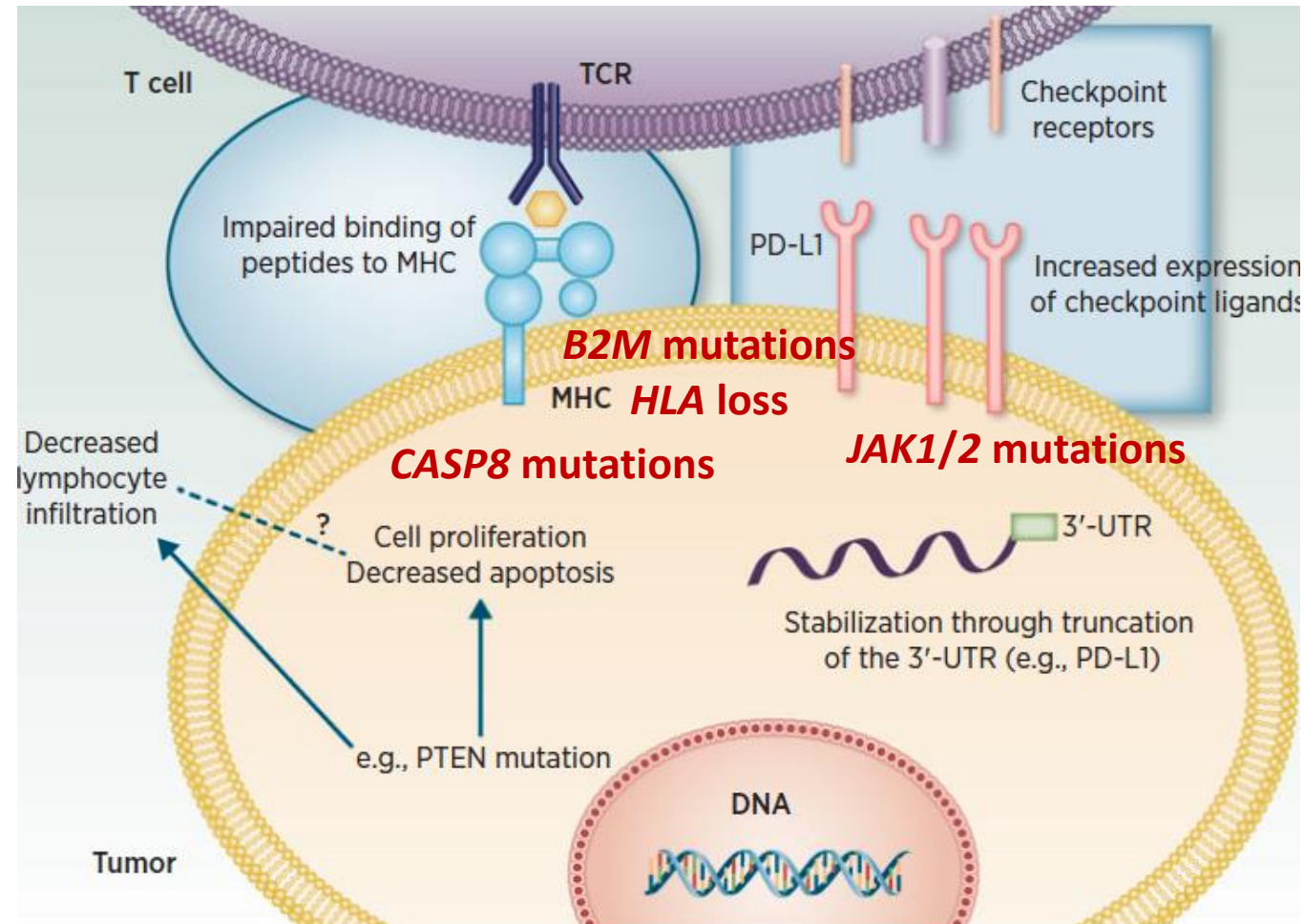
HIGH COPY NUMBER ALTERATIONS ACROSS TUMORS



INTERFERON PATHWAY MUTATIONS IN MELANOMA AND MSI COLON CANCER



ACQUIRED ANTIGEN PRESENTATION PATHWAY ALTERATIONS



TRANSCRIPTOMIC MARKERS



Response

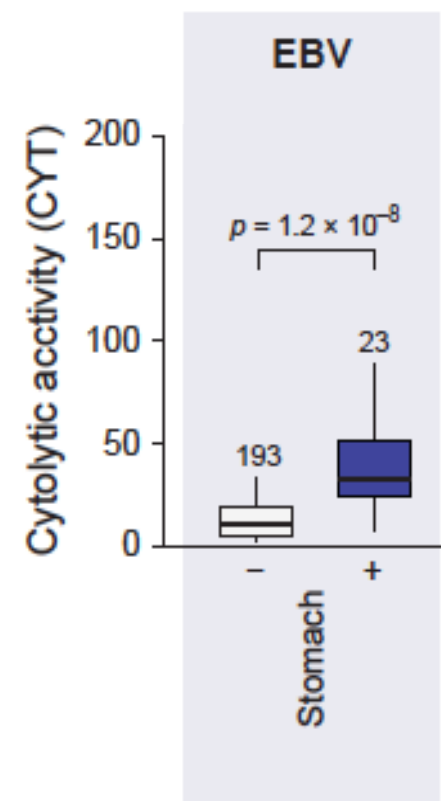
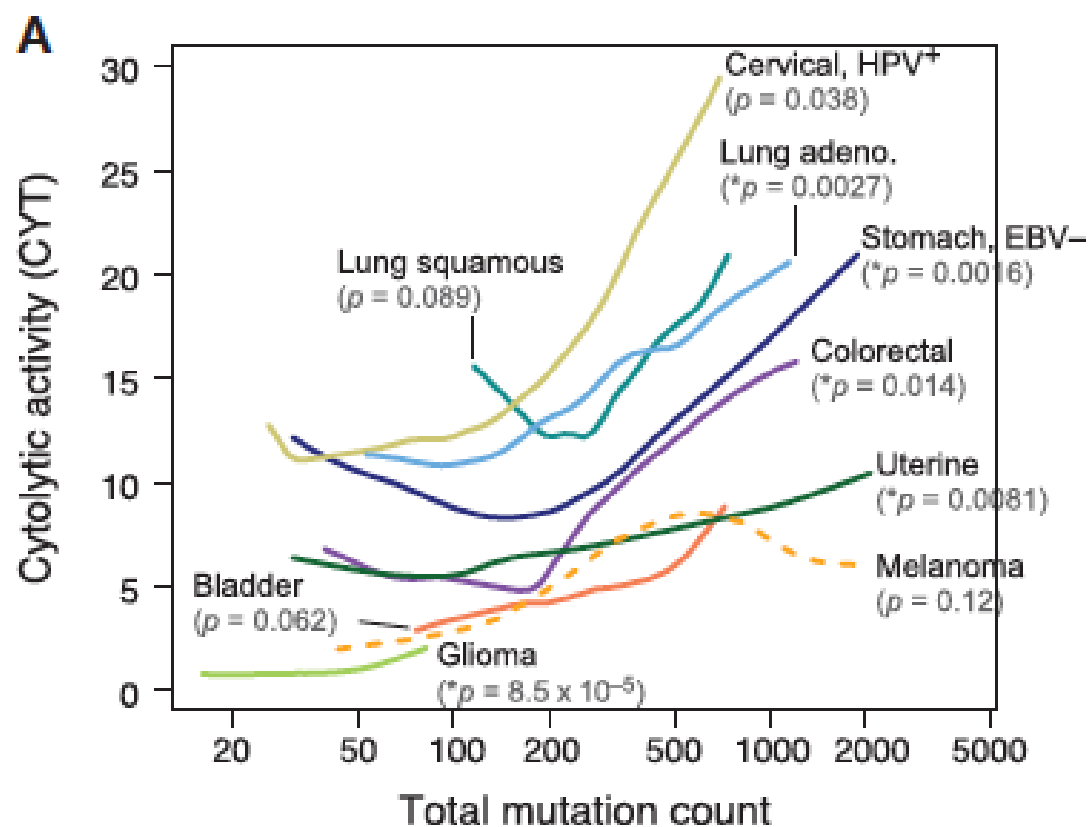
Cytotoxic (CD8 T cell inflammation) microenvironment signatures

Resistance

Pathway alterations – WNT high, TGFB high

CYTOTOXIC SIGNATURES

granzyme A (GZMA) and perforin (PRF1)

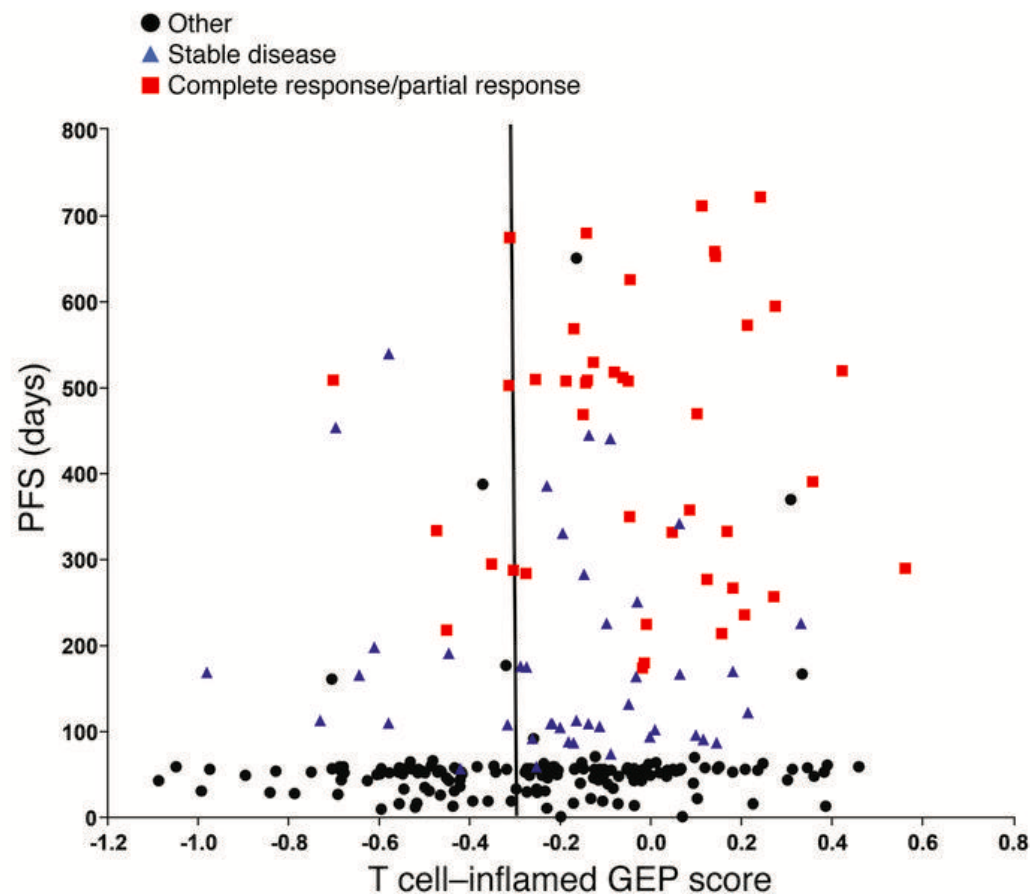


CYTOTOXIC SIGNATURES

Metanalysis

Melanoma, HNSCC, gastric
Pembrolizumab

GEP cytotoxicity signature (10 genes)
Nanostring nCounter

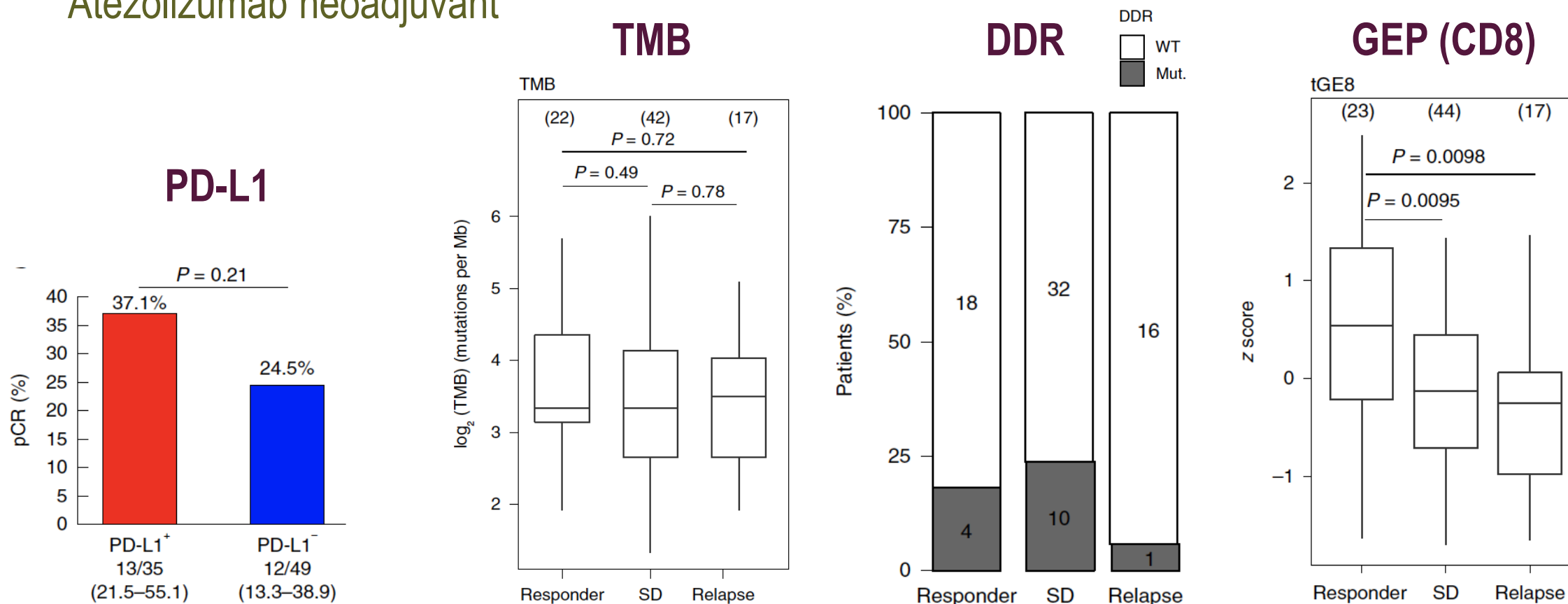


CYTOTOXIC SIGNATURES



ABACUS trial Bladder Cancer

Atezolizumab neoadjuvant



TRANSCRIPTOMIC MARKERS



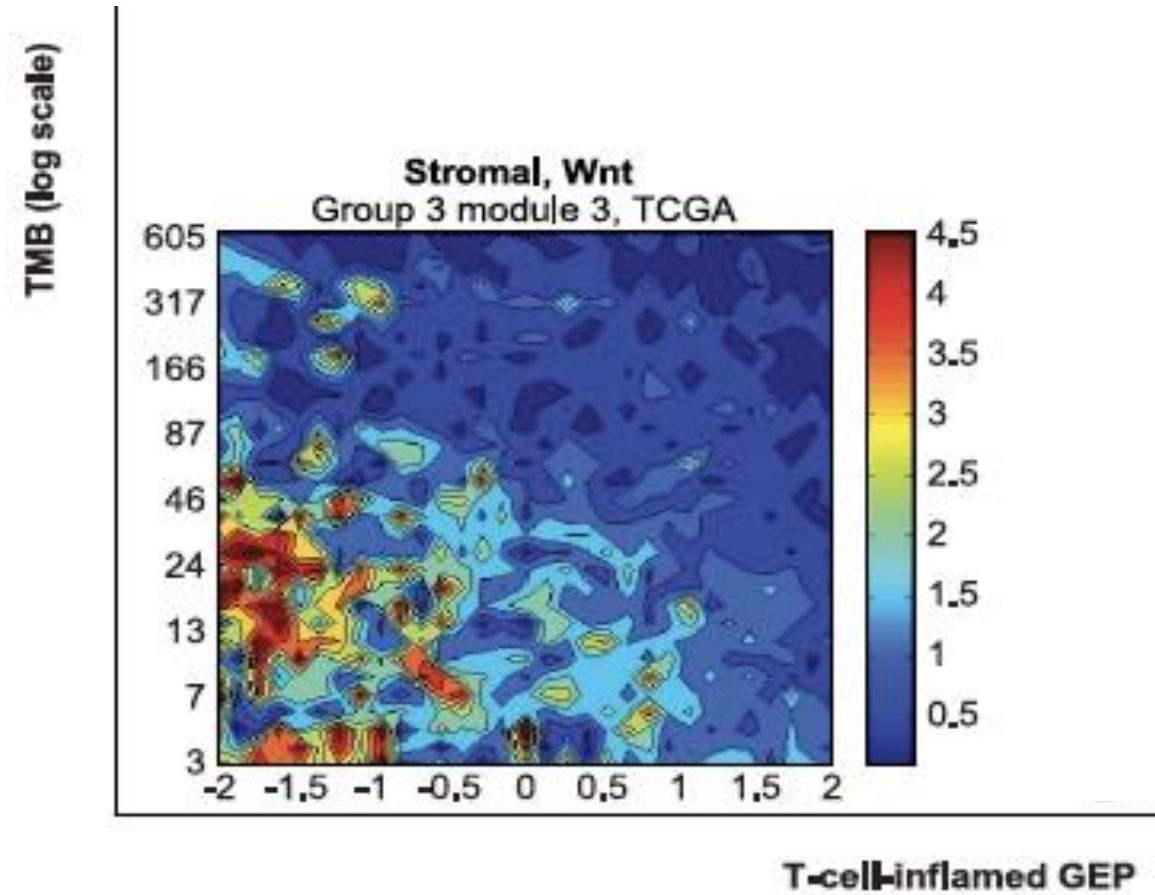
Response

Cytotoxic (CD8 T cell inflammation) microenvironment signatures

Resistance

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WNT AND TGFB HIGH SIGNATURES

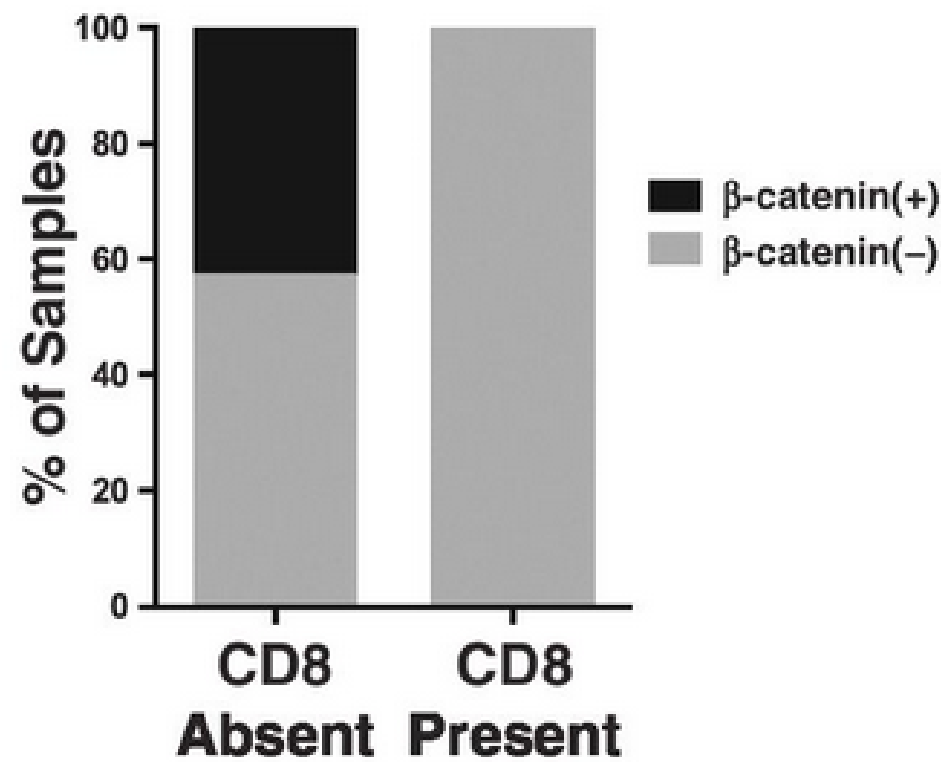


Cristescu et al, Science 2018

B CATENIN POSITIVE/HIGH

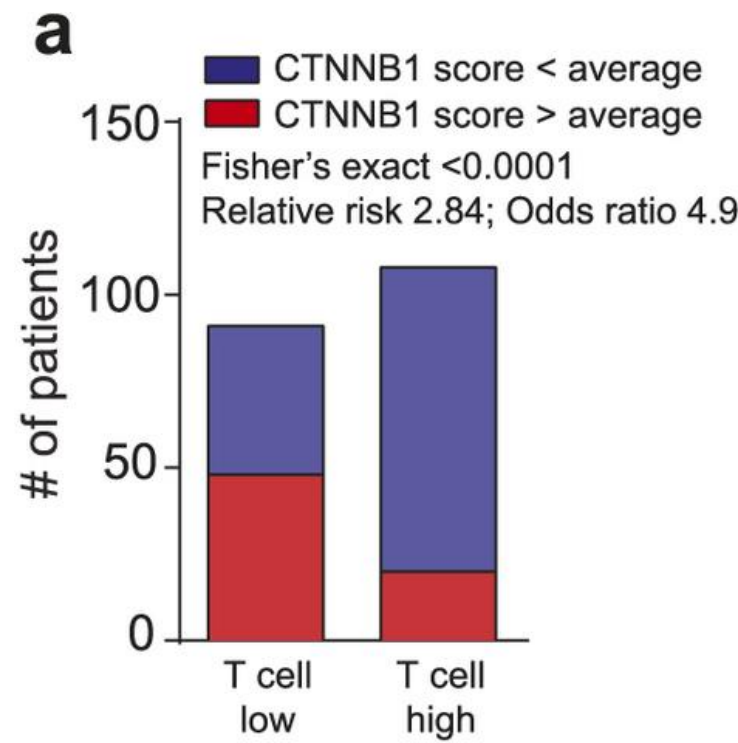


Bladder



Sweis et al, Cancer Immunol Res. 2016

Melanoma



Spranger et al, Nature 2015

CONCLUSIONS



- Response to immune checkpoint inhibitors is multi-factorial
- PD-L1, TMB and CD8 signatures (and combinations) have highest potential
 - Tumor type and disease setting matter
- There is room for further biomarker development (digital pathology, radiomics, patient genetics/phenomics)
- Design clinical trials with power for retrospective correlative analysis (not only “exploratory analysis”) in homogeneous populations.

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TIME TO MOVE TO NEGATIVE PREDICTIVE MARKERS?
ARE THERE UNIVERSAL RESISTANCE MARKERS?