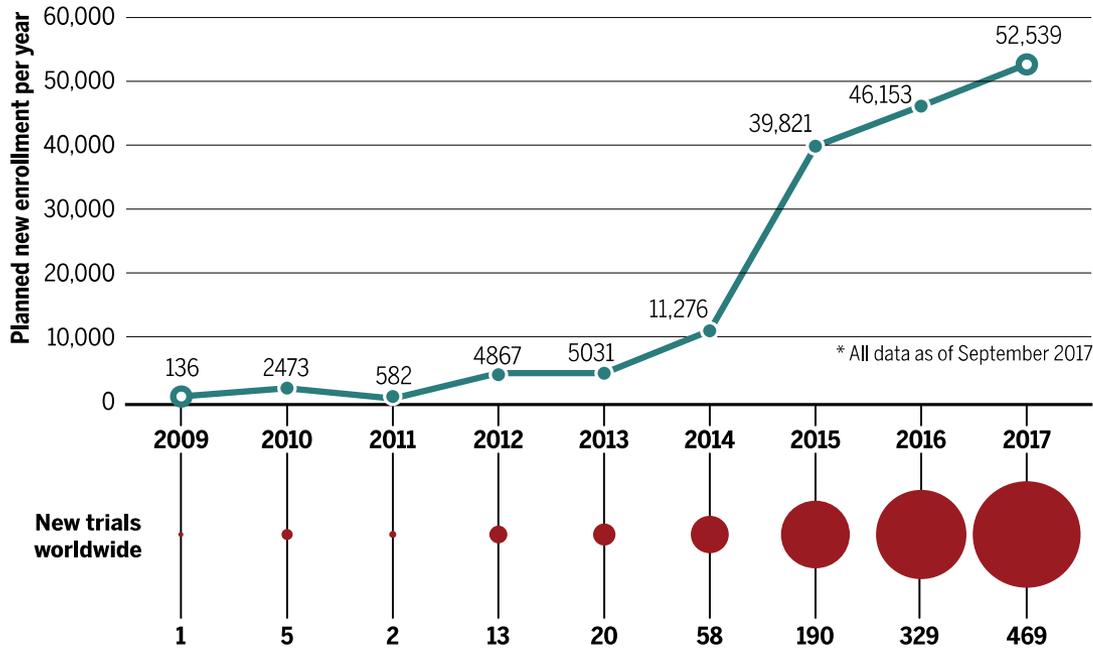


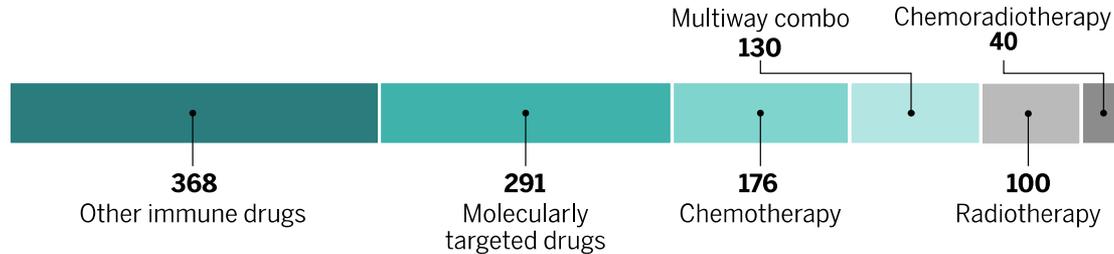
Enabling precision immuno-oncology

Zlatko Trajanoski
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Email: zlatko.trajanoski@i-med.ac.at
<http://icbi.at>

The promise of immuno-oncology



Combination trials with PD-1/PD-L1 inhibitors



December 2017: 3042 trials

Major issues in cancer immunotherapy

- Identify mechanisms of **intrinsic resistance** to checkpoint blockade
 - Predictive biomarkers for response (genetic, immunological, metabolic, microbiome)
- Identify mechanisms of **acquired resistance** to checkpoint blockade
 - Predictive biomarkers for relapse?
- Identify combination therapies with synergistic potential
 - PD-1/PD-L1 and targeted agents (or other drugs)

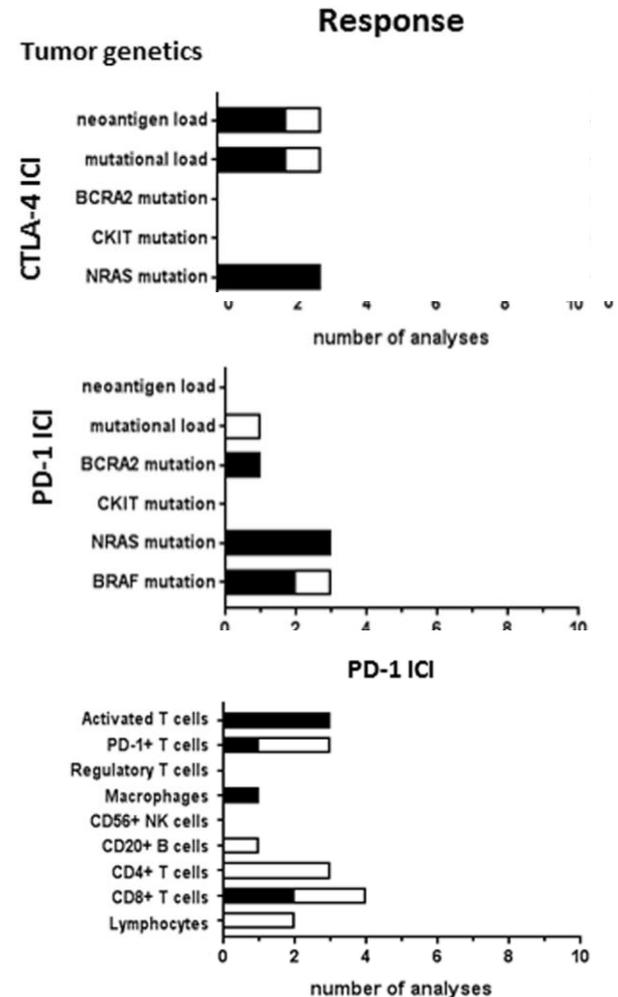
For every complex problem, there is
an answer that is clear, simple, and wrong

H.L. Mencken

Predictive markers for immunotherapy in melanoma

Implication of potential biomarkers in response to immunotherapy.

Biomarker	Checkpoint(s)/Therapies	Effect on response
Molecular		
NRAS mutations	CTLA-4, PD-1	Positive
NF1 mutations	PD-1	Positive
CD8+ T cells at tumor margin	PD-1	Positive
DC infiltration	T cell transfer therapy	Positive
PD-1 expression	PD-1	Positive
CTLA-4 expression	CTLA-4	Positive
PD-L1 expression	CTLA-4, PD-1, PD-L1	Positive
WNT/ β -catenin activation	CTLA-4, PD-L1	Negative
PTEN loss	PD-1	Negative
Mutational burden/ neoantigens	CTLA-4, PD-1, PD-L1	Positive
SERPIN family mutations	CTLA-4	Positive
IPRES transcriptional signature	PD-1	Negative
GNAQ/GNA11 mutations	CTLA-4, PD-1	Negative
MHC I/II expression	PD-1, PD-L1	Positive
JAK1/2 loss	PD-1	Negative
Clonal TCR repertoire	PD-1	Positive
Cytotoxic response	CTLA-4	Positive
IFN- γ response	CTLA-4, PD-L1	Positive
Peripheral Blood		
Elevated LDH	CTLA-4, PD-1	Negative
Low MDSCs	CTLA-4	Positive
Angiopoietin-2	CTLA-4	Negative
sPD-L1	CTLA-4, PD-1	Mixed
Circulating tumor DNA	PD-1	Negative
Clinical		
irAEs	CTLA-4, PD-1, PD-L1	Positive
Prior treatment	PD-1	Negative
Visceral involvement/tumor bulk	PD-1	Negative
Bifidobacterium in microbiome	PD-1	Positive
Uveal subtype	PD-1, PD-L1	Negative

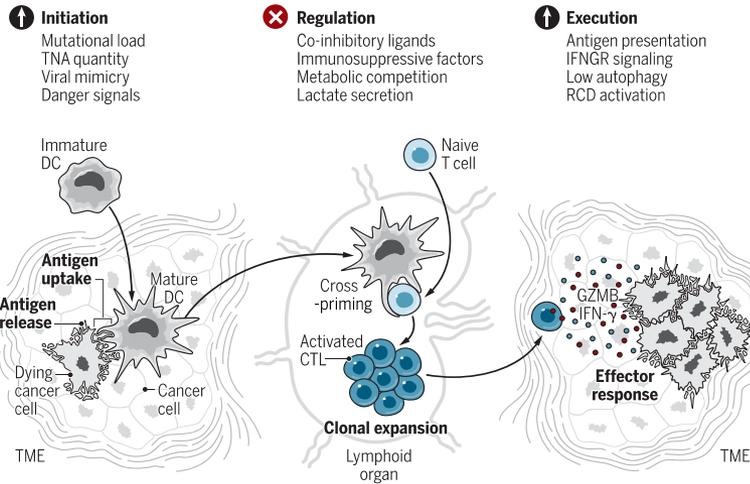


Predictive markers for immunotherapy with anti-PD-1 antibodies in melanoma

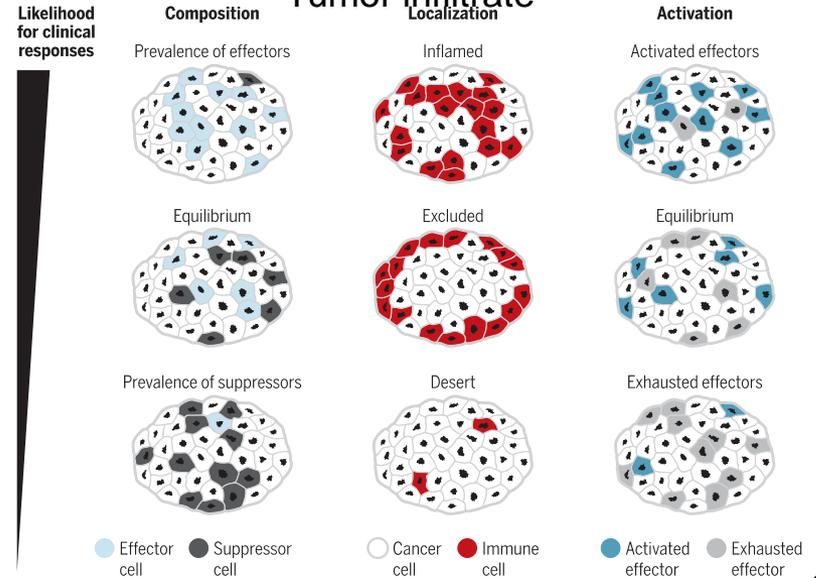
Publication	Marker(s)	Cohort size /p Value
Johnson DB <i>et al.</i> , <i>Cancer Immunol Res</i> 2016, 4:959-967	Mutational load	n=32/33; p=0.003/0.002
Hugo <i>et al.</i> , <i>Cell</i> 2016, 165:35-44	IPRES signature	n=28; p=0.04
Johnson DB <i>et al.</i> , <i>Nat Commun</i> 2016, 7:10582	HLA-DR	n=30/23; p=0.055/0.046
Diem <i>et al.</i> , <i>Br J Cancer</i> 2016, 114:256-261	LDH	n=29; p<0.001 (ANOVA)
Charoentong <i>et al.</i> , <i>Cell Rep</i> 2017, 18:248-262	162 immune genes	n=28; p=0.025
Ayers <i>et al.</i> , <i>J Clin Invest</i> 2017, 127:2930-2939	28 immune genes	n=62; p=0.027

The hallmarks of successful immunotherapy

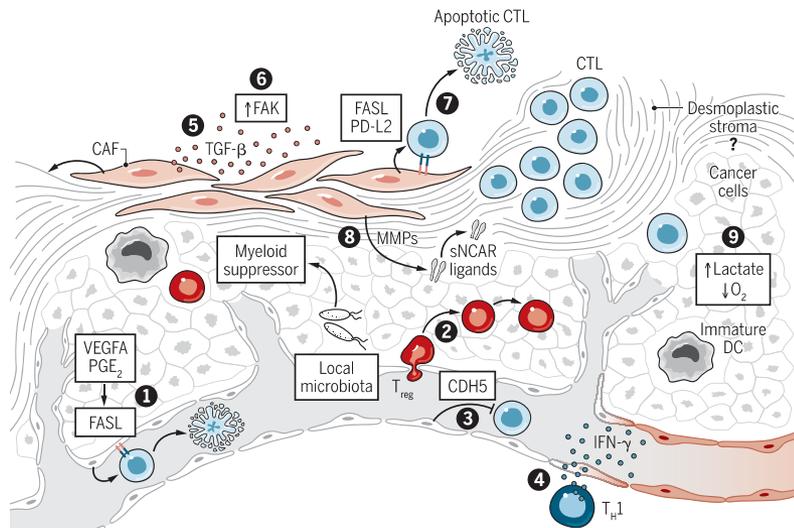
Malignant cells



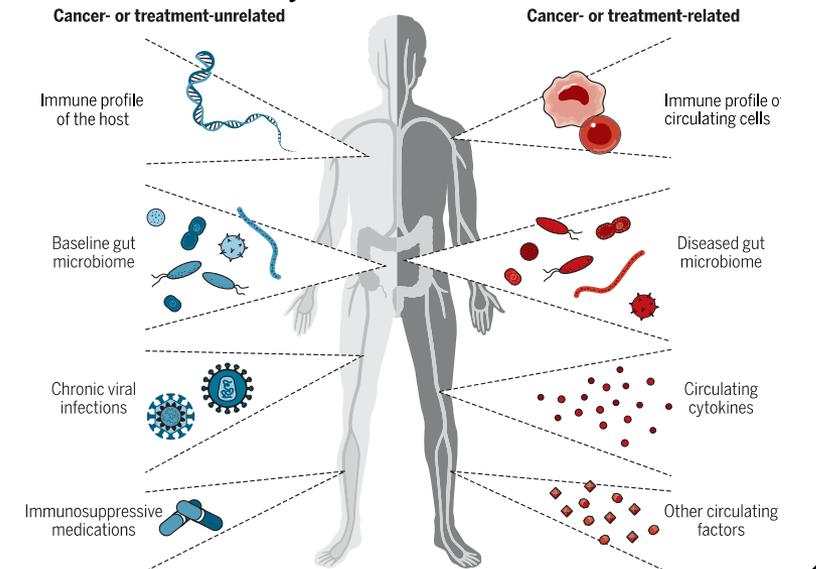
Tumor infiltrate



Tumor stroma and vasculature



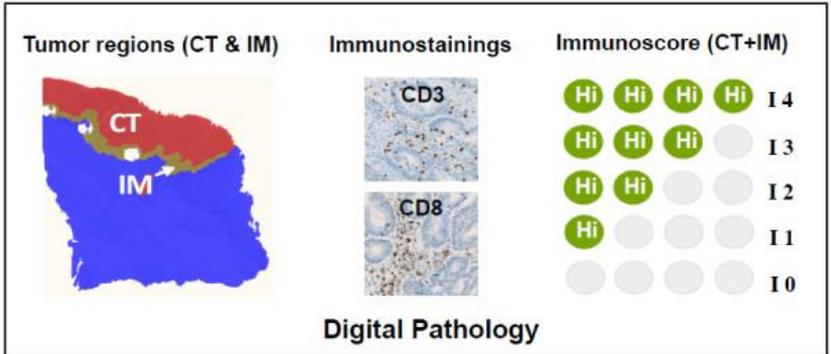
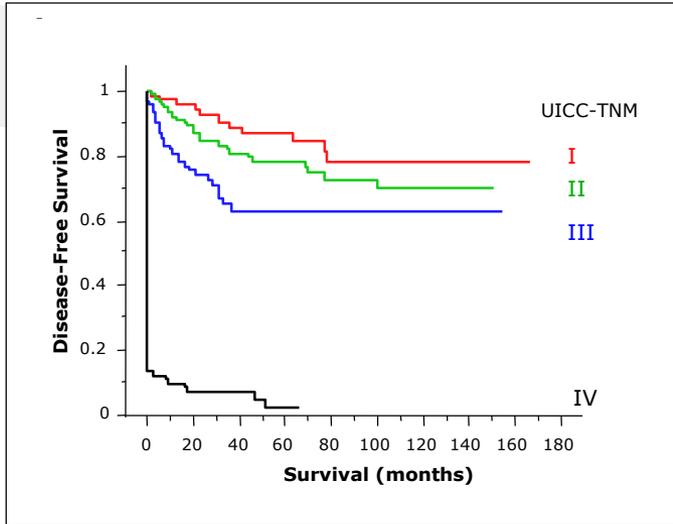
Systemic factors



Immunoscore: prognostic biomarker for tumor recurrence in colorectal cancer

Tumor histopathology

UICC-TNM Staging system

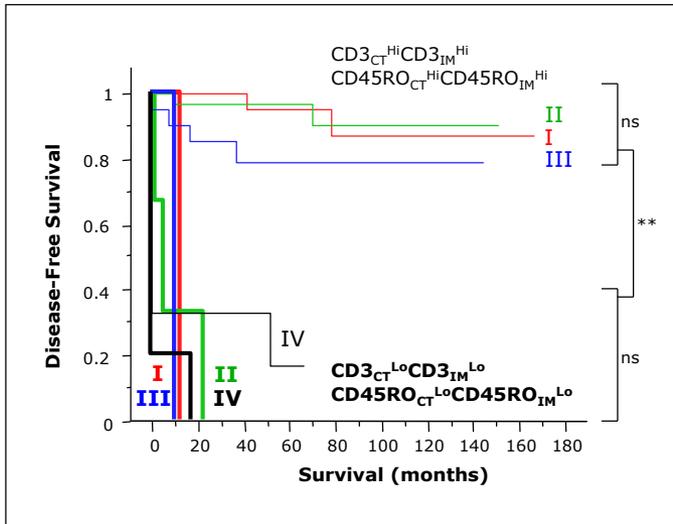


Mlecnik *et al. J Clin Oncol* 2011, 29:610-618

CD3_{CT} CD3_{IM} evaluation

plus

CD45RO_{CT} CD45RO_{IM} evaluation

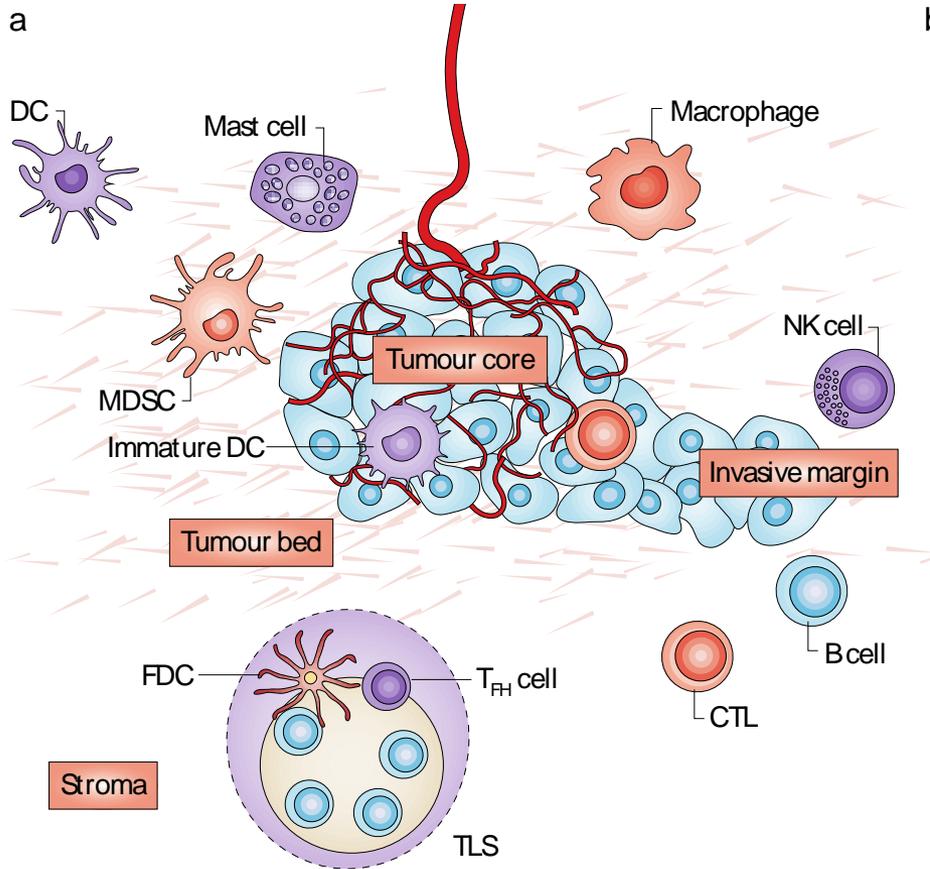


International validation of the consensus Immunoscore for the classification of colon cancer: a prognostic and accuracy study

Franck Pagès, Bernhard Mlecnik, Florence Marliot, Gabriela Bindea, Fang-Shu Ou, Carlo Bifulco, Alessandro Lugli, Inti Zlobec, Tilman T Rau, Martin D Berger, Iris D Nagtegaal, Elisa Vink-Börger, Arndt Hartmann, Carol Geppert, Julie Kolwelter, Susanne Merkel, Robert Grützmann, Marc Van den Eynde, Anne Jouret-Mourin, Alex Kartheuser, Daniel Léonard, Christophe Remue, Julia Y Wang, Prashant Bavi, Michael H A Roehrl, Pamela S Ohashi, Linh T Nguyen, SeongJun Han, Heather L MacGregor, Sara Hafezi-Bakhtiari, Bradly G Wouters, Giuseppe V Masucci, Emilia K Andersson, Eva Zavadova, Michal Vocka, Jan Spacek, Lubos Petruzzelka, Bohuslav Konopasek, Pavel Dundr, Helena Skalova, Kristyna Nemejcova, Gerardo Botti, Fabiana Tatangelo, Paolo Delrio, Gennaro Ciliberto, Michele Maio, Luigi Laghi, Fabio Grizzi, Tessa Fredriksen, Bénédicte Buttard, Mihaela Angelova, Angela Vasaturo, Pauline Maby, Sarah E Church, Helen K Angell, Lucie Lafontaine, Daniela Bruni, Carine El Sissy, Nacilla Haicheur, Amos Kirilovsky, Anne Berger, Christine Lagorce, Jeffrey P Meyers, Christopher Paustian, Zipei Feng, Carmen Ballesteros-Merino, Jeroen Dijkstra, Carlijn van de Water, Shannon van Lent-van Vliet, Nikki Knijn, Ana-Maria Musină, Dragos-Viorel Scripcariu, Boryana Popivanova, Mingli Xu, Tomonobu Fujita, Shoichi Hazama, Nobuaki Suzuki, Hiroaki Nagano, Kiyotaka Okuno, Toshihiko Torigoe, Noriyuki Sato, Tomohisa Furuhashi, Ichiro Takemasa, Kyogo Itoh, Prabhu S Patel, Hemangini H Vora, Birva Shah, Jayadrakumar B Patel, Kruti N Rajvik, Shashank J Pandya, Shilin N Shukla, Yili Wang, GuanJun Zhang, Yutaka Kawakami, Francesco M Marincola, Paolo A Ascierto, Daniel J Sargent*, Bernard A Fox, Jérôme Galon

Lancet 2018; 391: 2128-39

Immune contexture



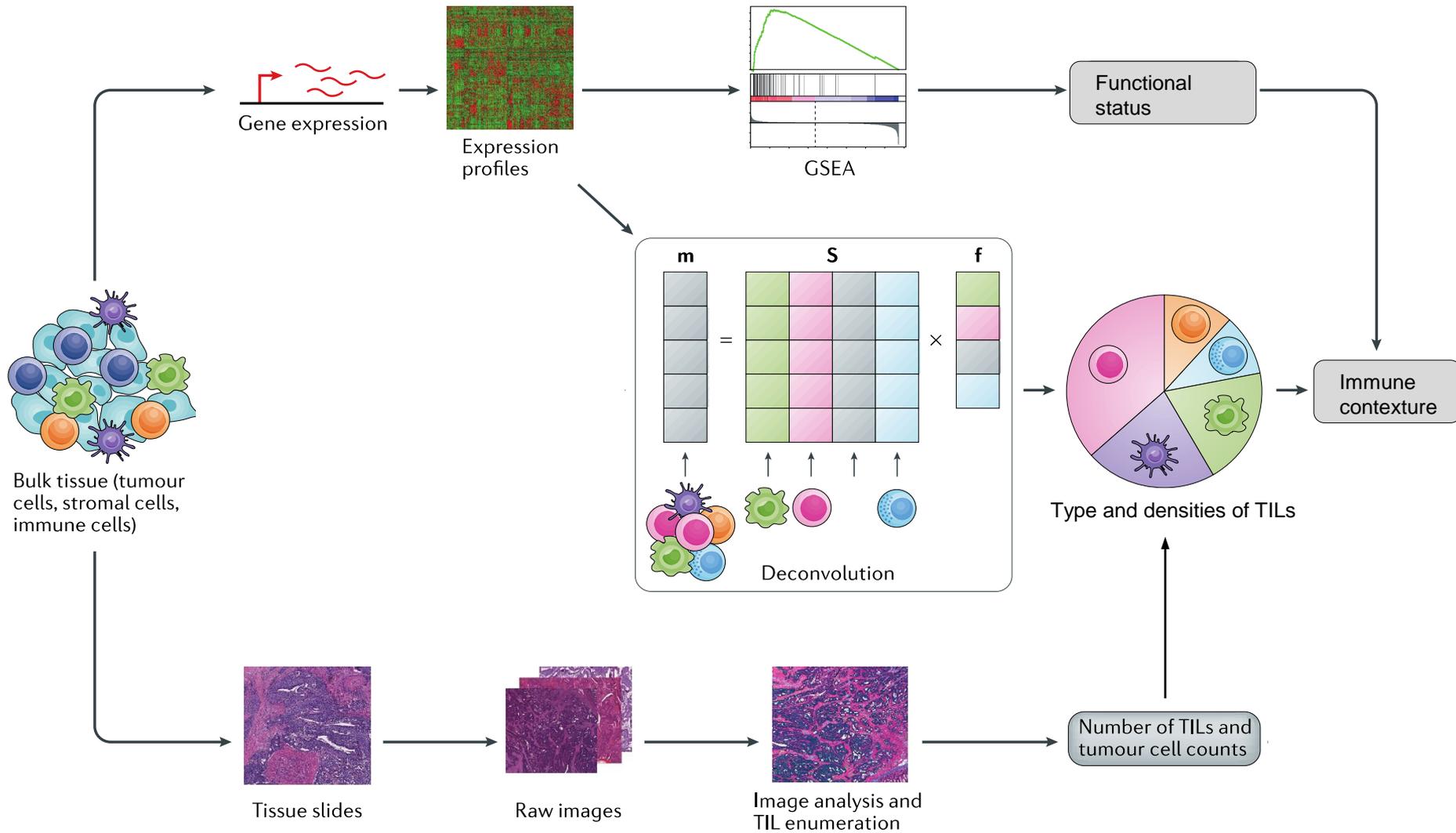
b

Immune contexture	Parameters: positive association with survival
Type	CTLs (CD3 ⁺ CD8 ⁺) Memory T cells (CD45RO ⁺)
Location	Core of the tumour Invasive margin
Density	<p>Number of cells per mm²</p> <p>1 10 100 1,000 10,000</p> <p>CD3⁺_{CT}</p> <p>CD3⁺_{IM}</p> <p>CD8⁺_{CT}</p> <p>CD8⁺_{IM}</p> <p>CD45RO⁺_{CT}</p> <p>CD45RO⁺_{IM}</p>
Functional orientation	<p>T_H1 cell-associated factors (IFNγ, IL-12, T-bet and IRF1)</p> <p>Cytotoxic factors (granzymes, perforin and granulysin)</p> <p>Chemokines (CX3CL1, CXCL9, CXCL10, CCL5 and CCL2)</p> <p>T_H17 cells, T_{Reg} cells and T_H2 cells have a variable</p>
TLS	Presence and quality

Additional obstacles

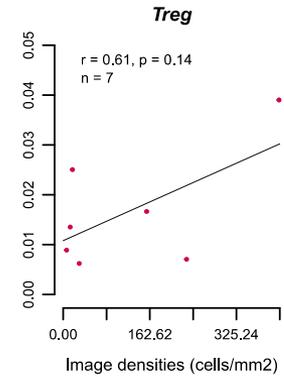
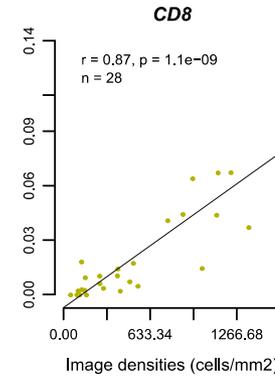
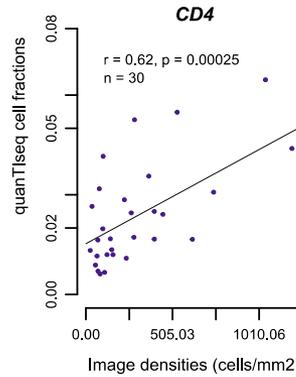
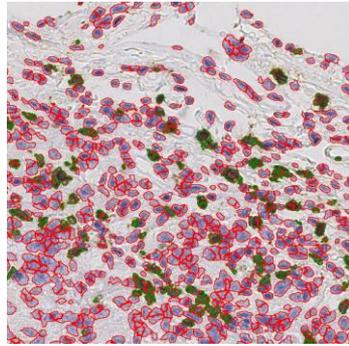
1. Epigenetic, genetic and immunologic heterogeneity; tumor evolution
 - Multi-region sampling
 - Multiple time points sampling
2. Cancer-cell extrinsic factors
 - Metabolism
 - Cellular senescence
3. Interactions between other treatment forms and immunotherapy
 - Chemotherapy, radiotherapy
 - Targeted therapy
4. Missing techniques to assay the entire complexity of TME
 - RNA-seq
 - Immunohistochemistry

Quantifying immune contexture using NGS and imaging data

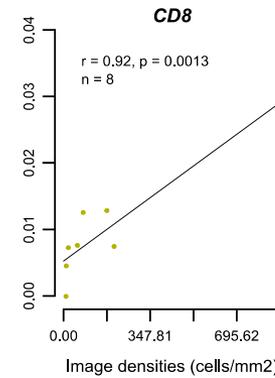
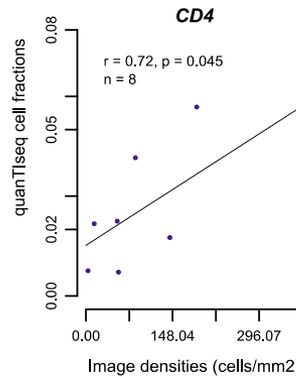
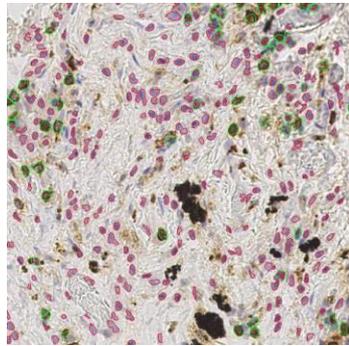


quanTiseq validation

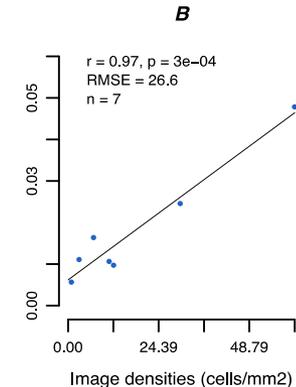
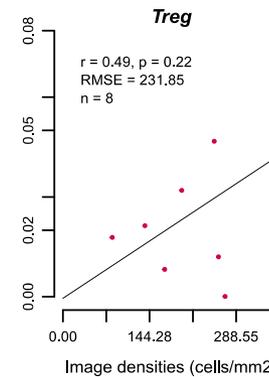
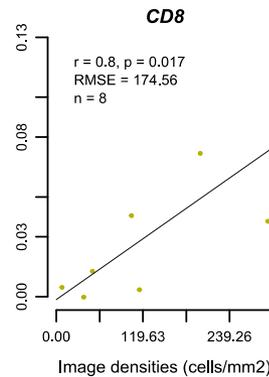
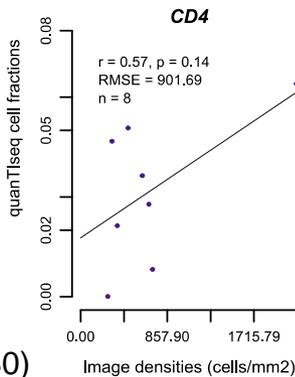
Melanoma
n=30
J. Balko
Vanderbilt University



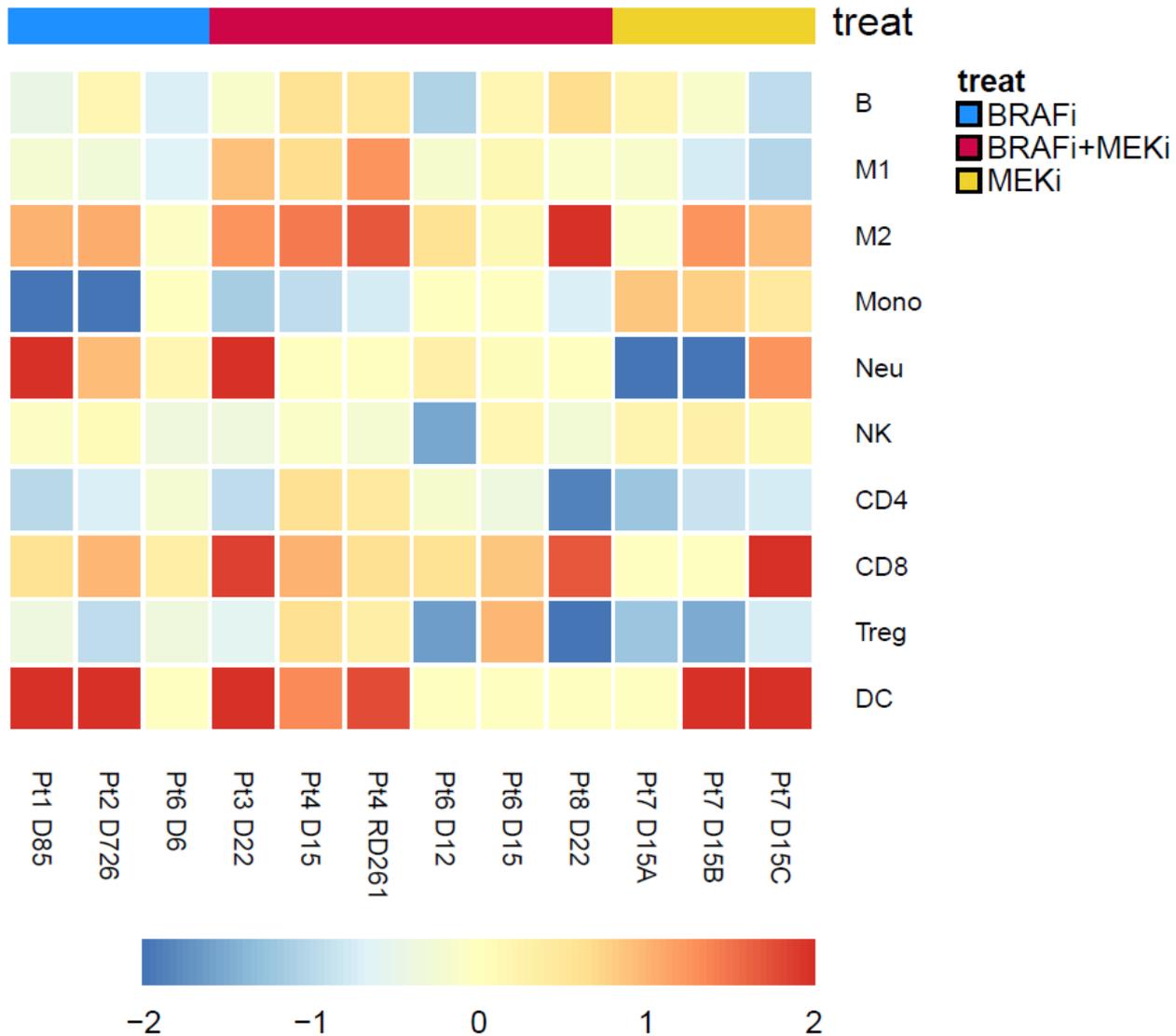
Lung cancer
n=8
J. Balko
Vanderbilt University



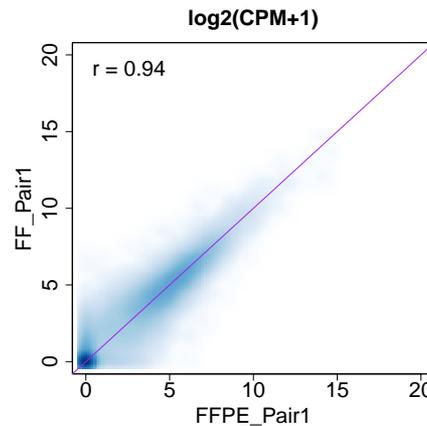
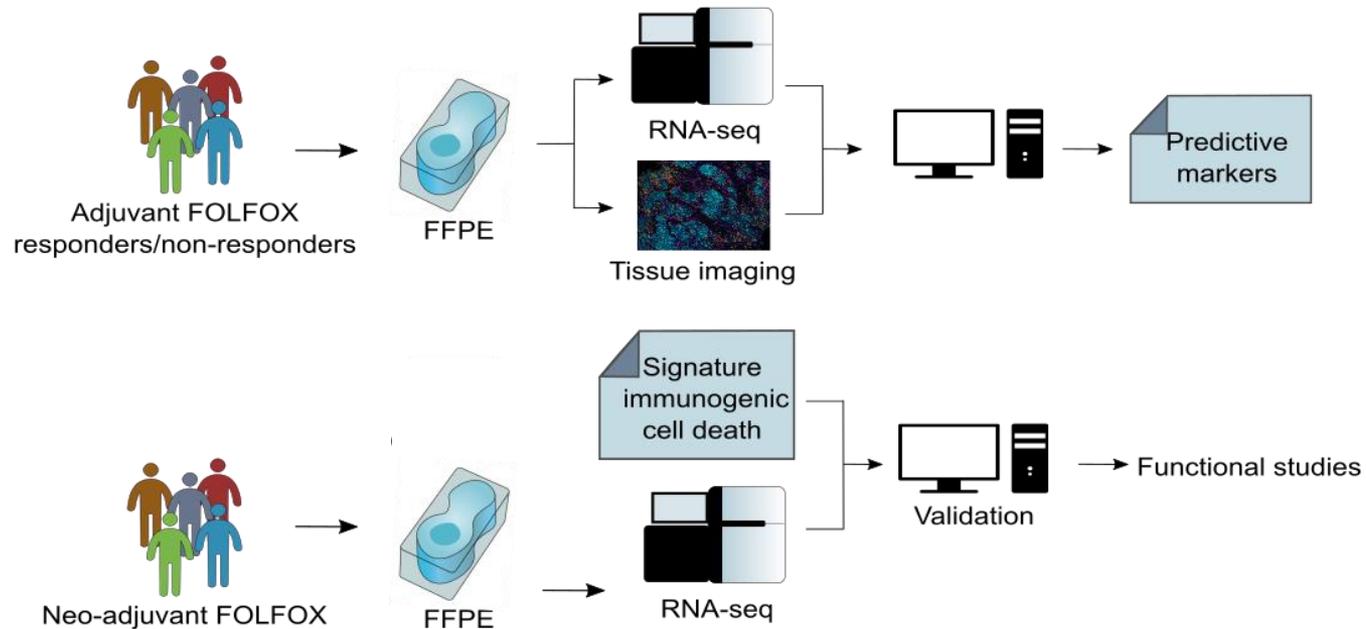
CRC
n=8
N. de Miranda
Leiden University



Immunogenic effects of BRAFi/MEKi in melanoma



Immunogenic effects of chemotherapy

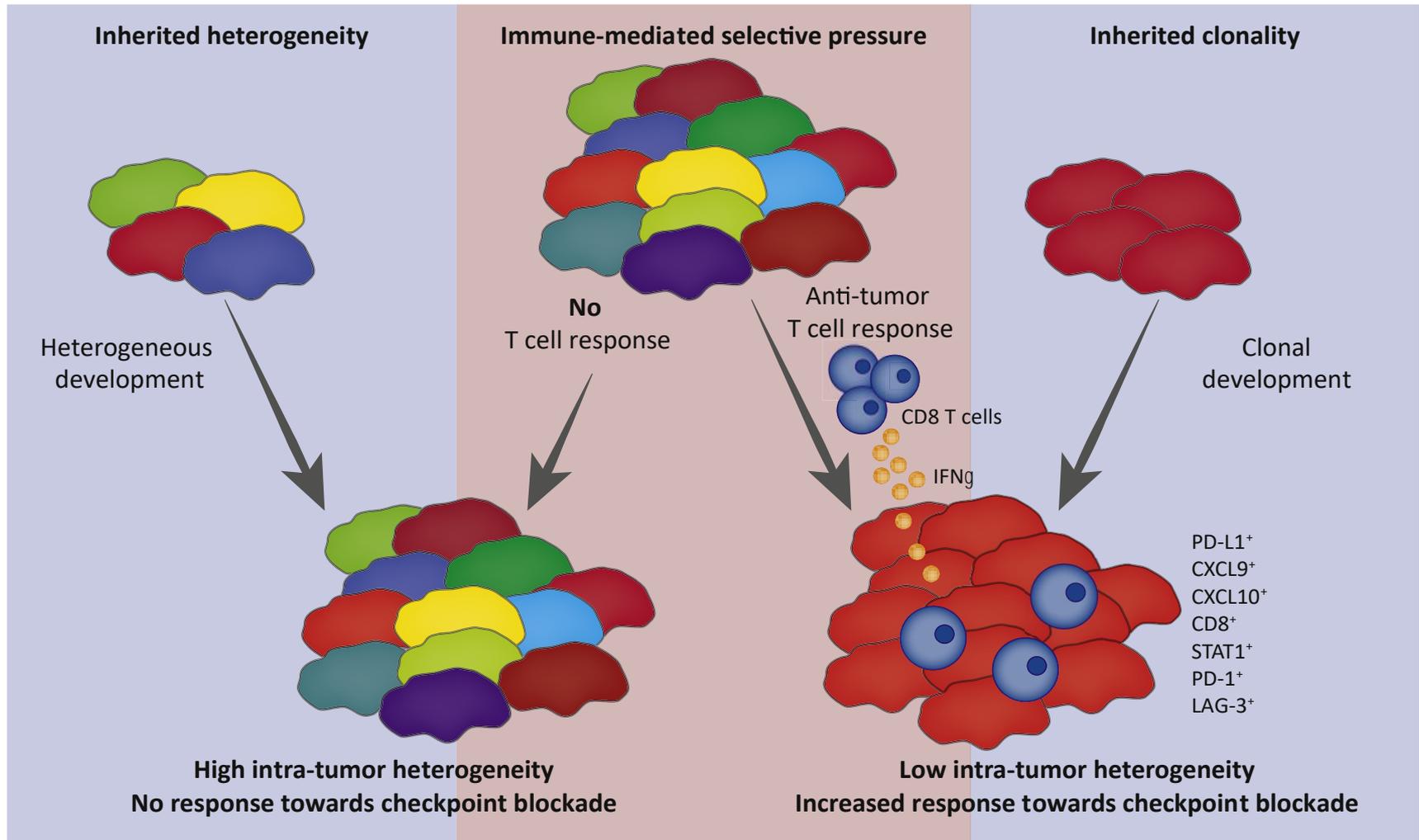


Collaboration with H. Fiegl, C. Marth
Medical University Innsbruck

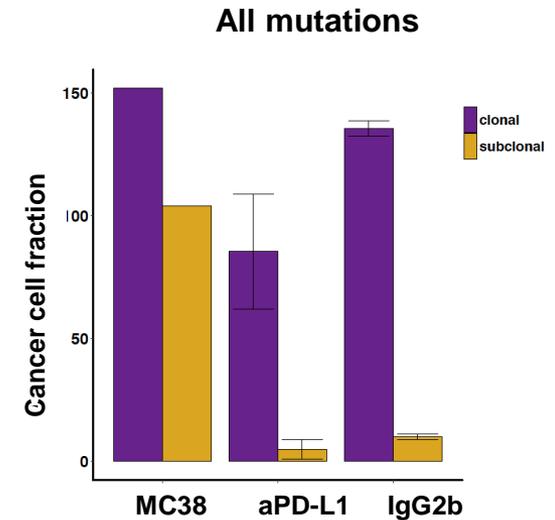
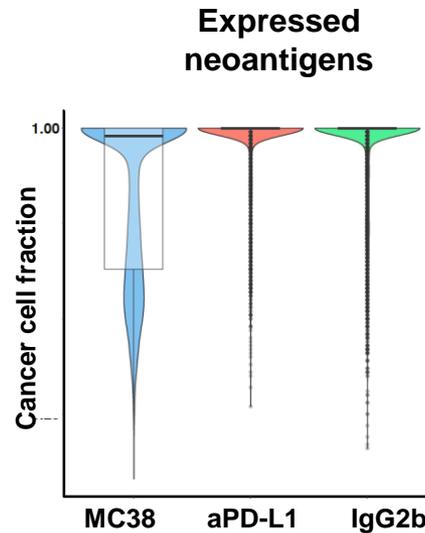
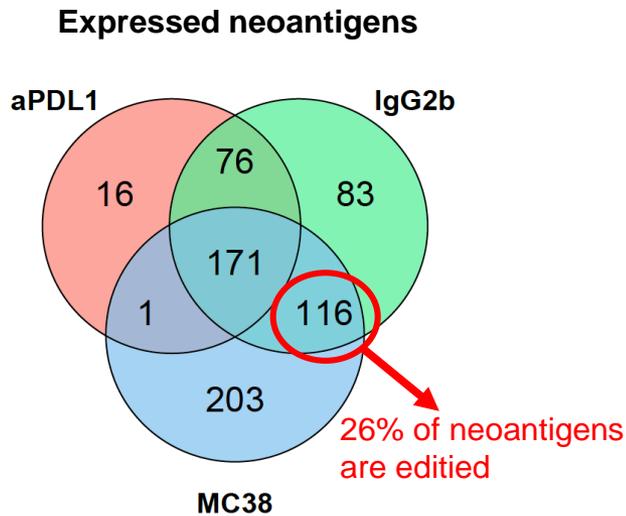
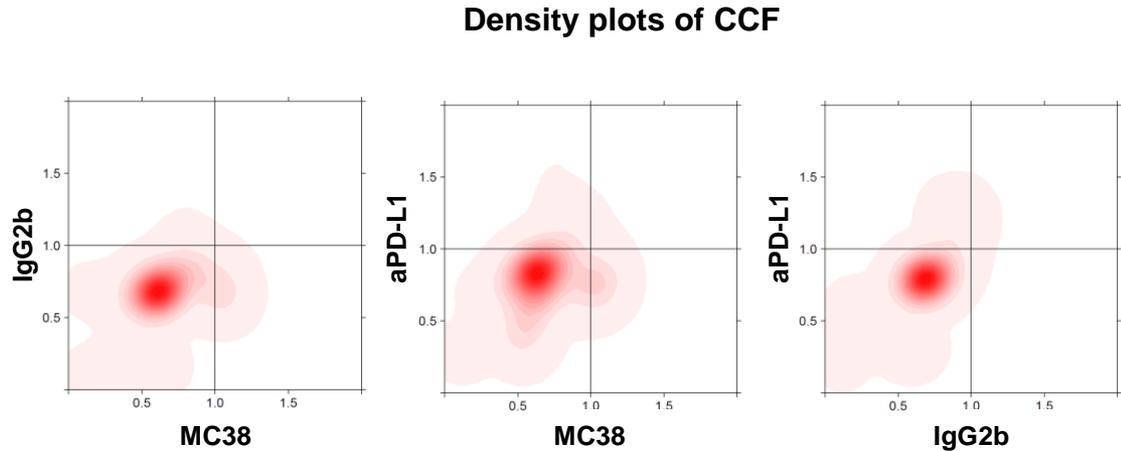
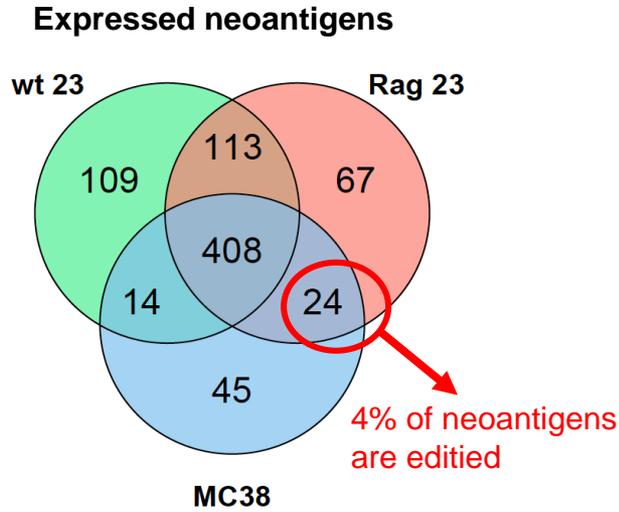
Major issues in cancer immunotherapy

- Identify mechanisms of intrinsic resistance to checkpoint blockade
 - Predictive biomarkers for response (genetic, immunological, metabolic, microbiome)
- Identify mechanisms of **acquired resistance** to checkpoint blockade
 - Predictive biomarkers for relapse?
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 - PD-1/PD-L1 and targeted agents (or other drugs)

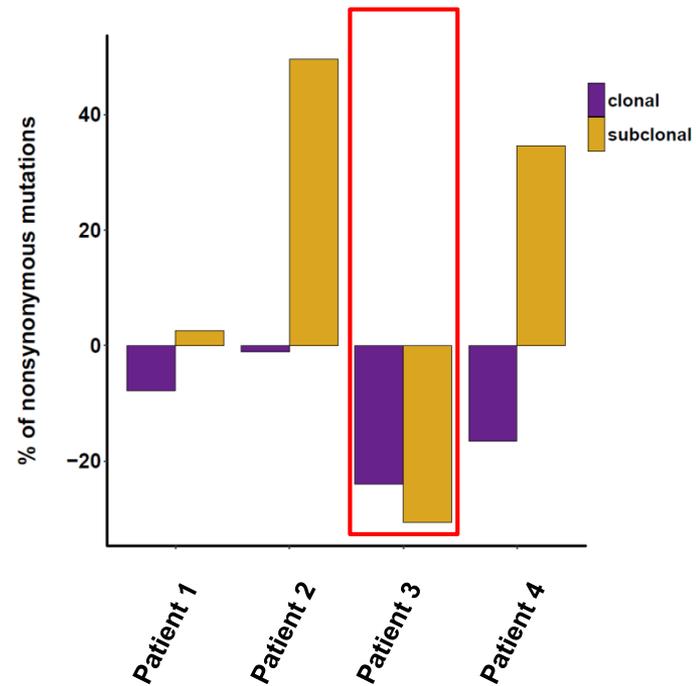
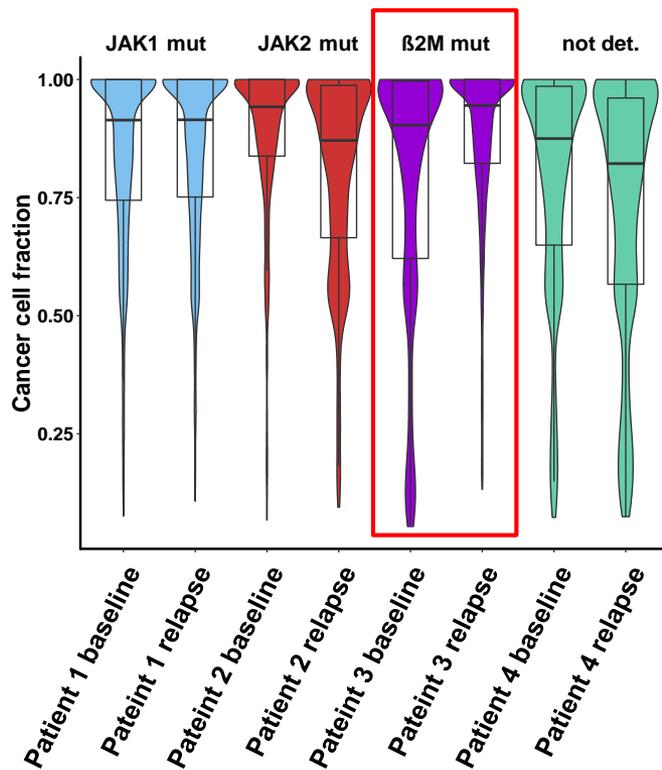
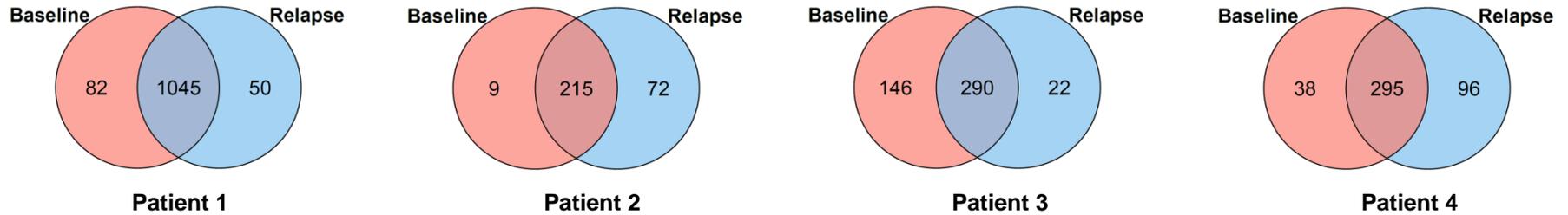
Immunoediting and tumor heterogeneity



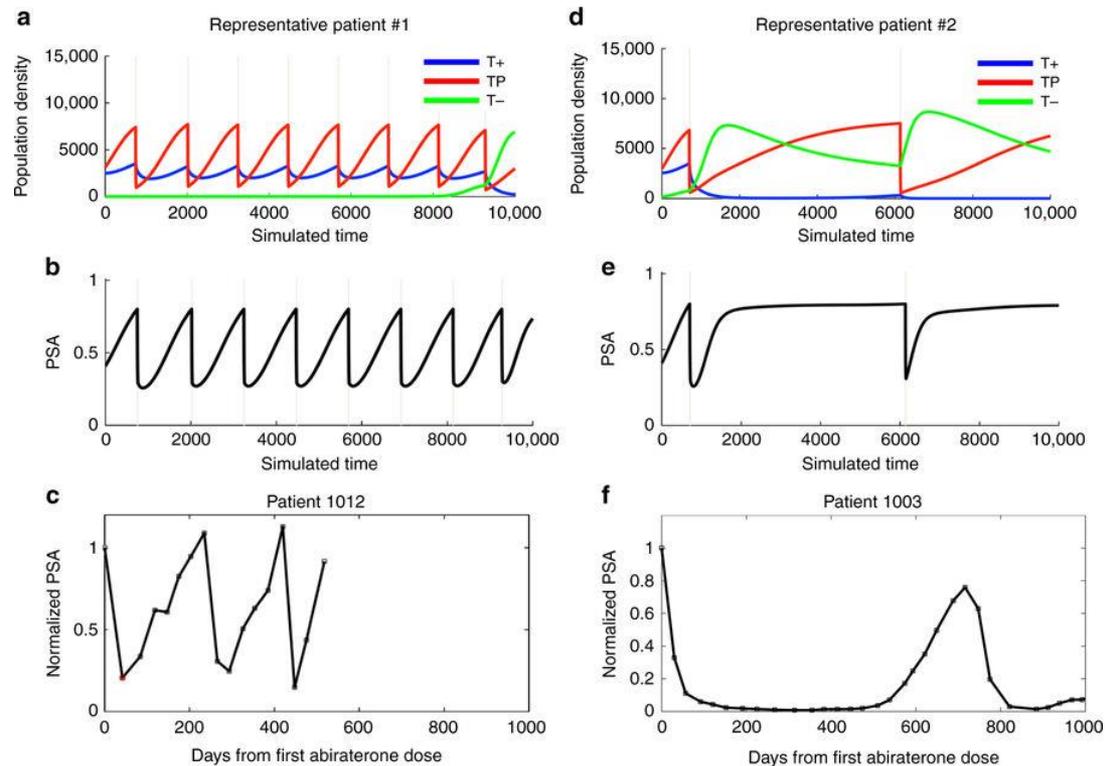
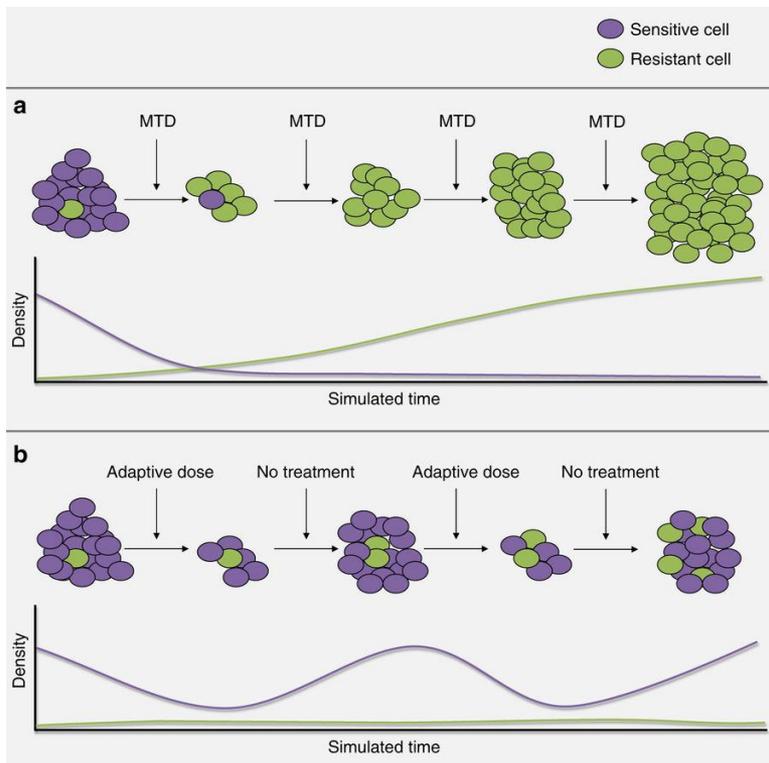
Targeting the PD-1/PD-L1 pathway broadens the T cell repertoire and renders the tumors more homogeneous



Acquired resistance to PD-1 blockade in melanoma



Adaptive therapy in metastatic prostate cancer using patient-specific evolutionary dynamics



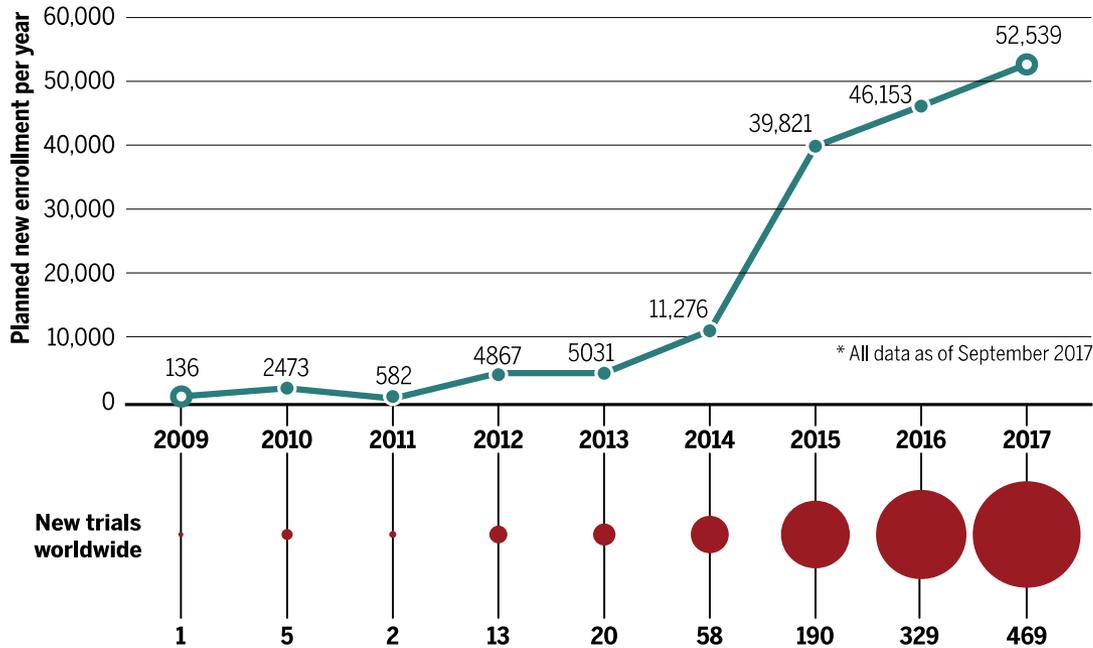
Development of treatment algorithms for cancer immunotherapy:

- Monitoring tumor evolution (liquid biopsies, radiomics, TCR repertoire)
- Evolutionary dynamic models of immunoediting

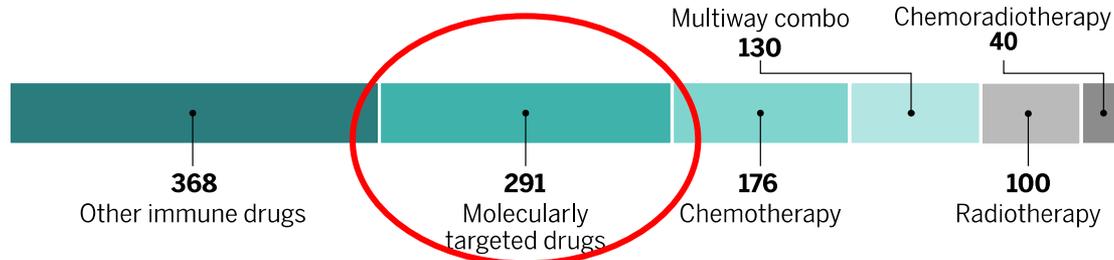
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Cancer immunotherapy



Combination trials with PD-1/PD-L1 inhibitors



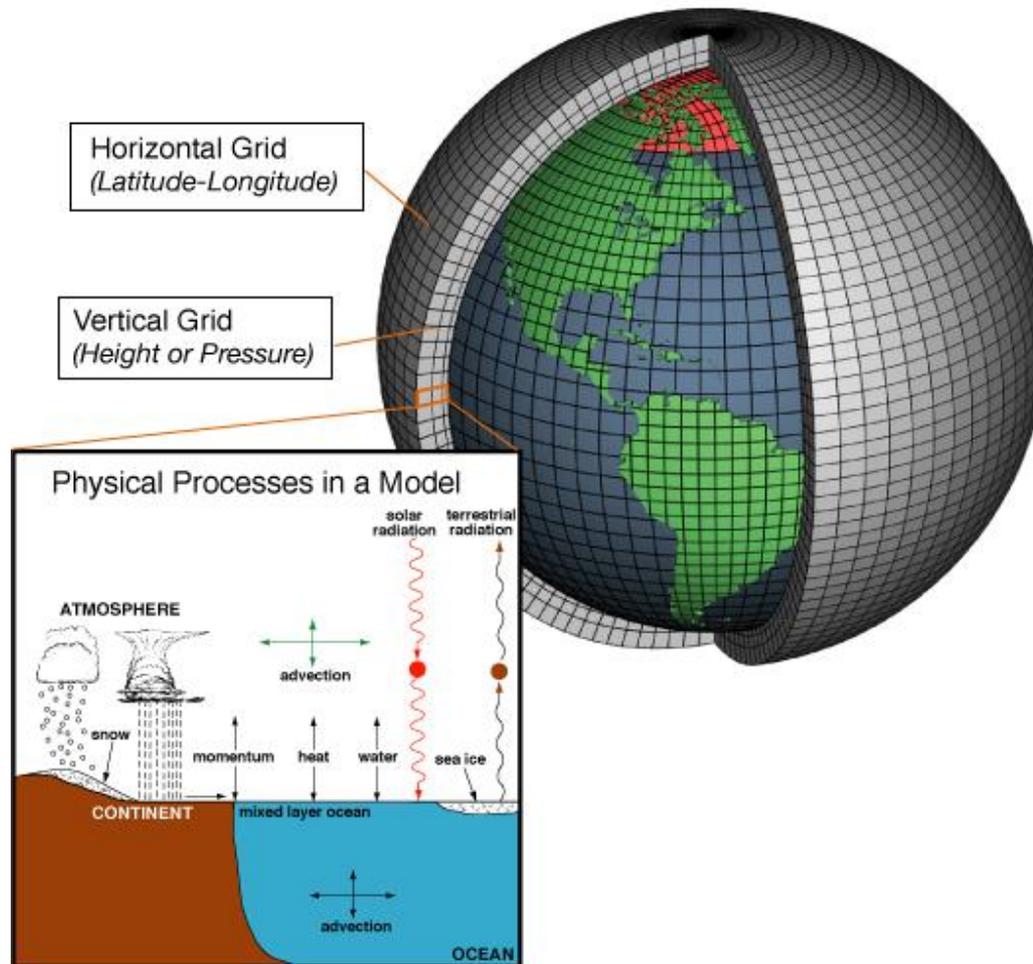
December 2017: 3042 trials

Enabling precision immuno-oncology?

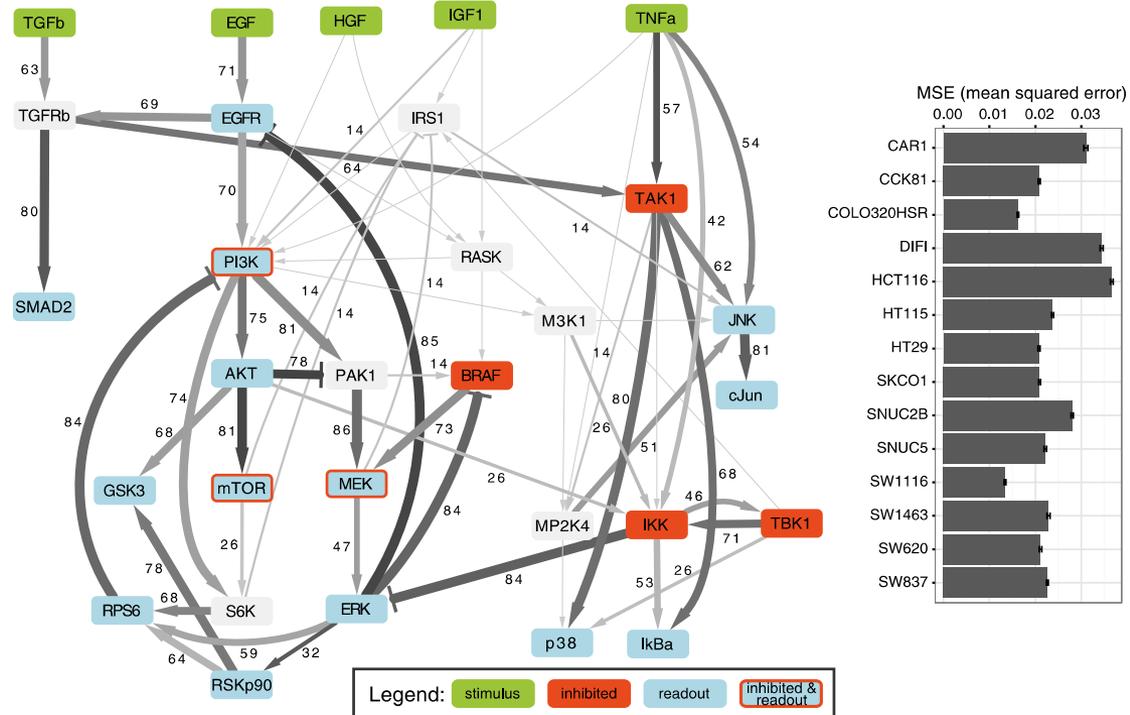
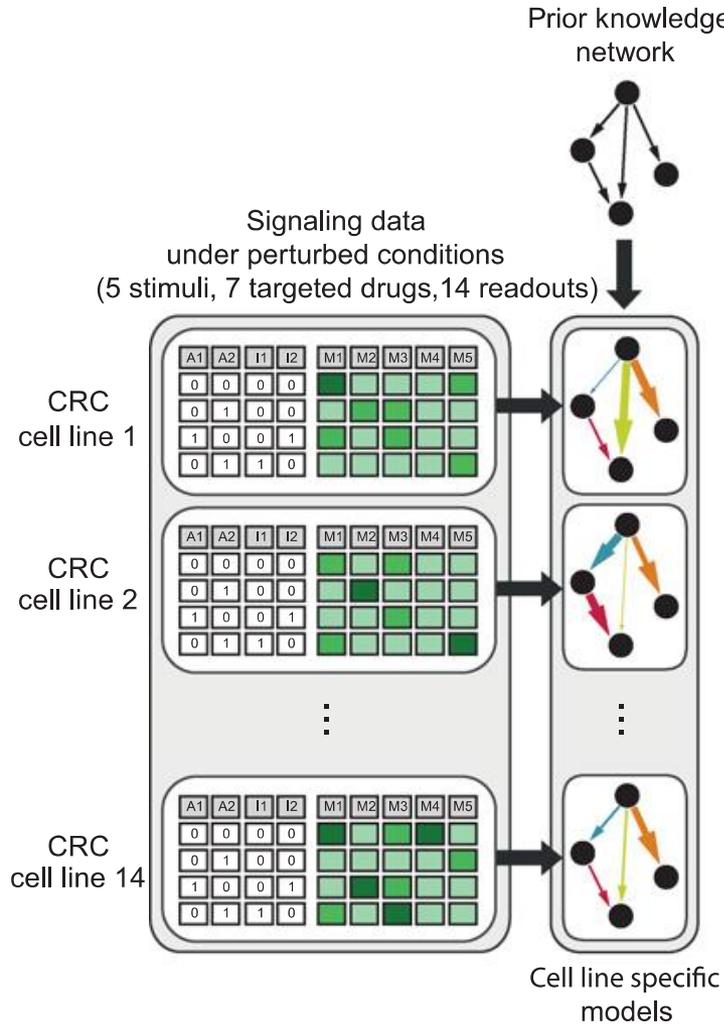
- Mutational and neoantigen landscapes are highly individual
 - Colorectal cancer (Angelova *et al.*, *Genome Biology*, 2015, 16:64)
 - Solid cancers (Charoentong *et al.*, *Cell Rep*, 2017, 18:248-262)
- Oncogenic signaling is cell context-specific
 - BRAF mutant CRC is resistant to BRAF inhibitors (Pralhad *et al.*, *Nature* 2012, 483:100-103)
- Tumor evolution and immune responses are dynamic and interwoven systems
 - In general molecular data only from single time points is available
- Mouse models are not suitable for testing precision immuno-oncology
 - Patient-derived xenografts (PDX) mouse models are immunocompromised
 - Current humanized mouse models cannot mimic the entire complexity of the tumor microenvironment

Prediction of dynamic systems

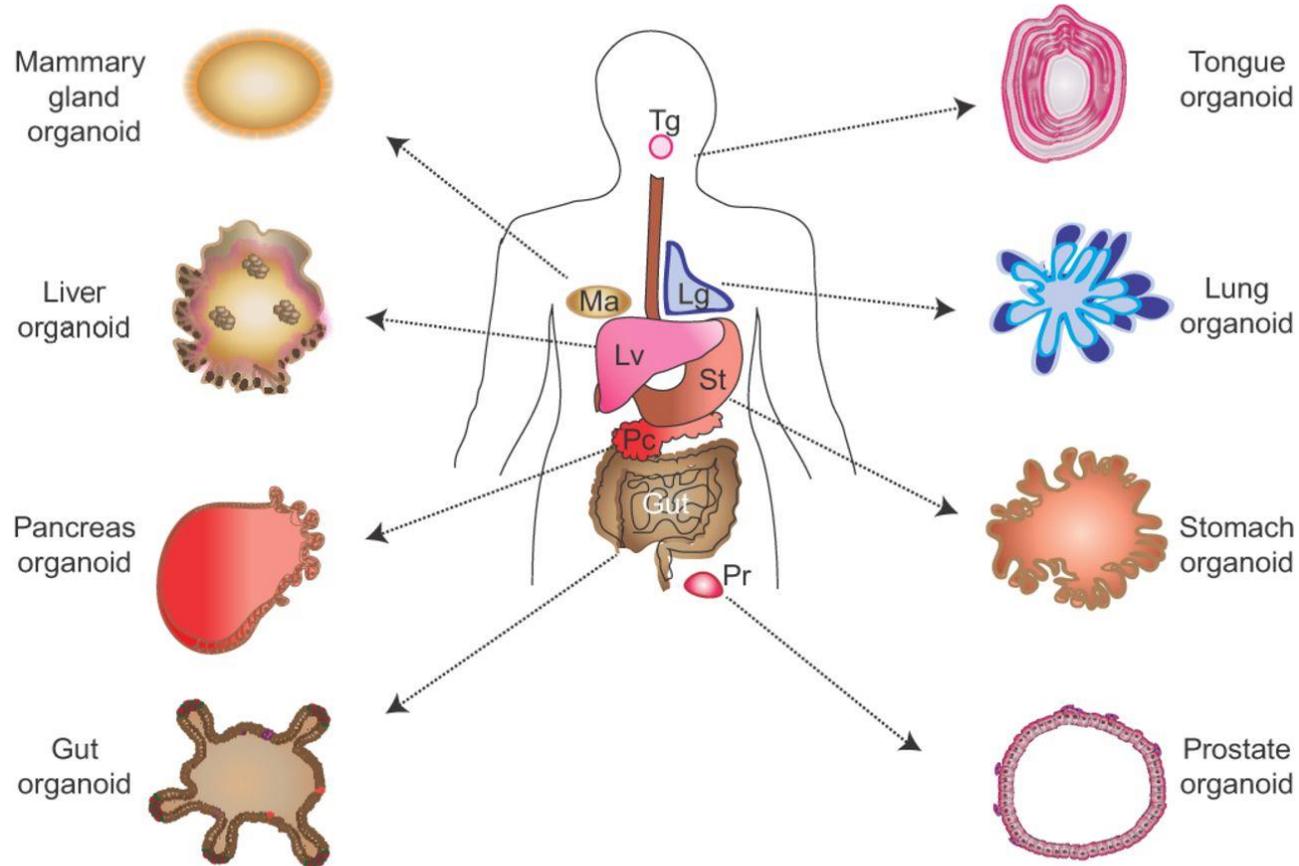
Weather forecasting



Perturbation biology

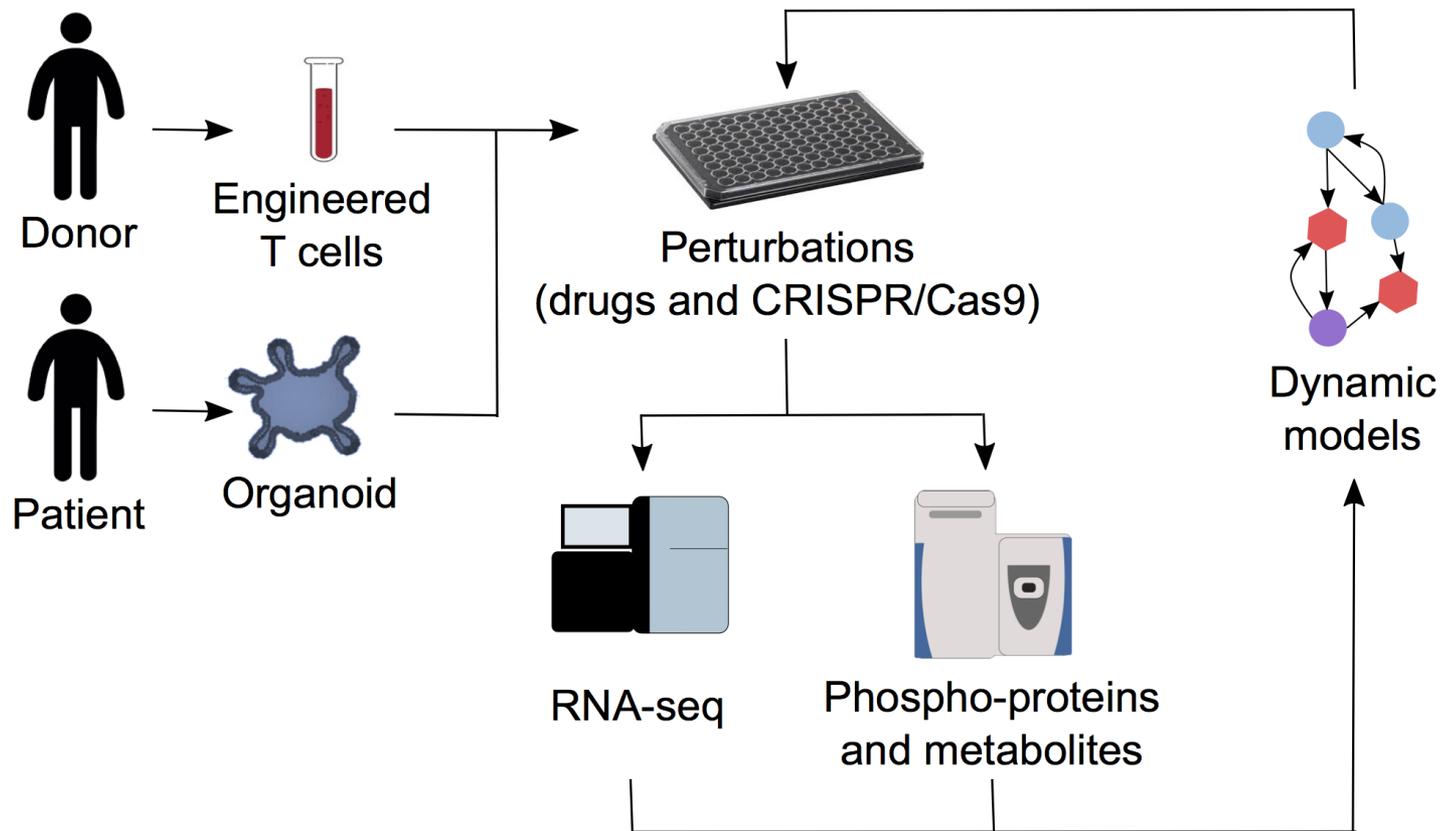


In vitro models: Organoids



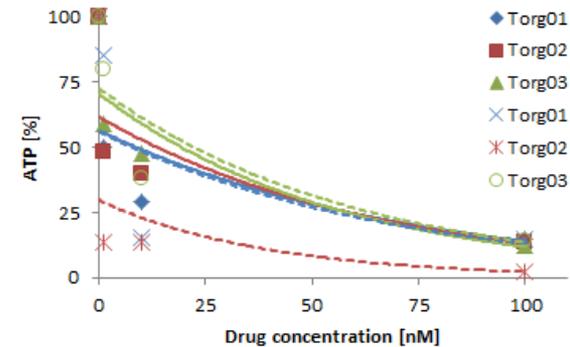
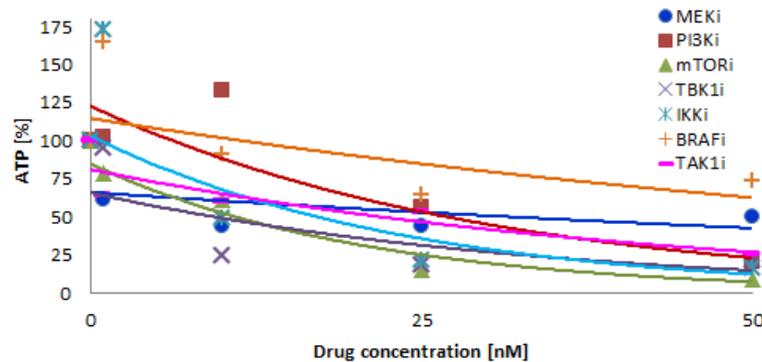
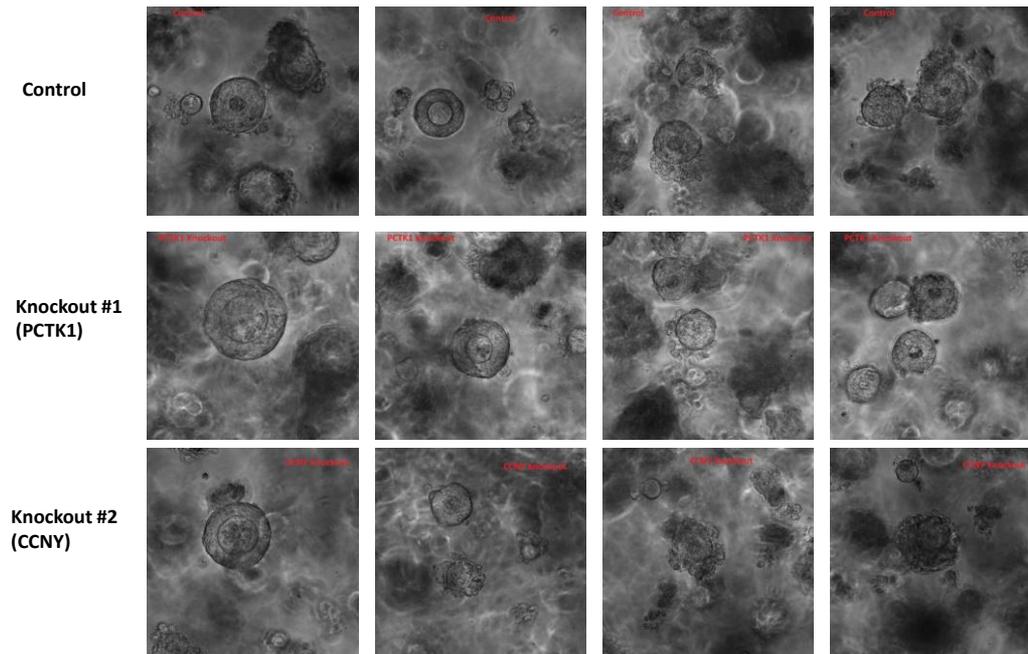
Hybrid avatars using *in vitro* and *in silico* models

Perturbation biology to derive mechanistic rationale



Hybrid avatars using *in vitro* and *in silico* models

Perturbation biology to derive mechanistic rationale



Summary

- Identification of predictive biomarkers for response to immune checkpoint blockade:
 - Novel comprehensive assays/assay combinations are required
- Identification of predictive biomarkers for relapse:
 - Non-invasive assays for tumor monitoring are required
- Enabling precision immuno-oncology:
 - Need for avatars and perturbation data

Enabling precision immuno-oncology



Grossvenediger 3666 m



Stilfser Joch

Zlatko Trajanoski

Division for Bioinformatics, Medical University of Innsbruck, Austria

Email: zlatko.trajanoski@i-med.ac.at <http://icbi.at>