

Data and Specimen sharing- Industry perspective

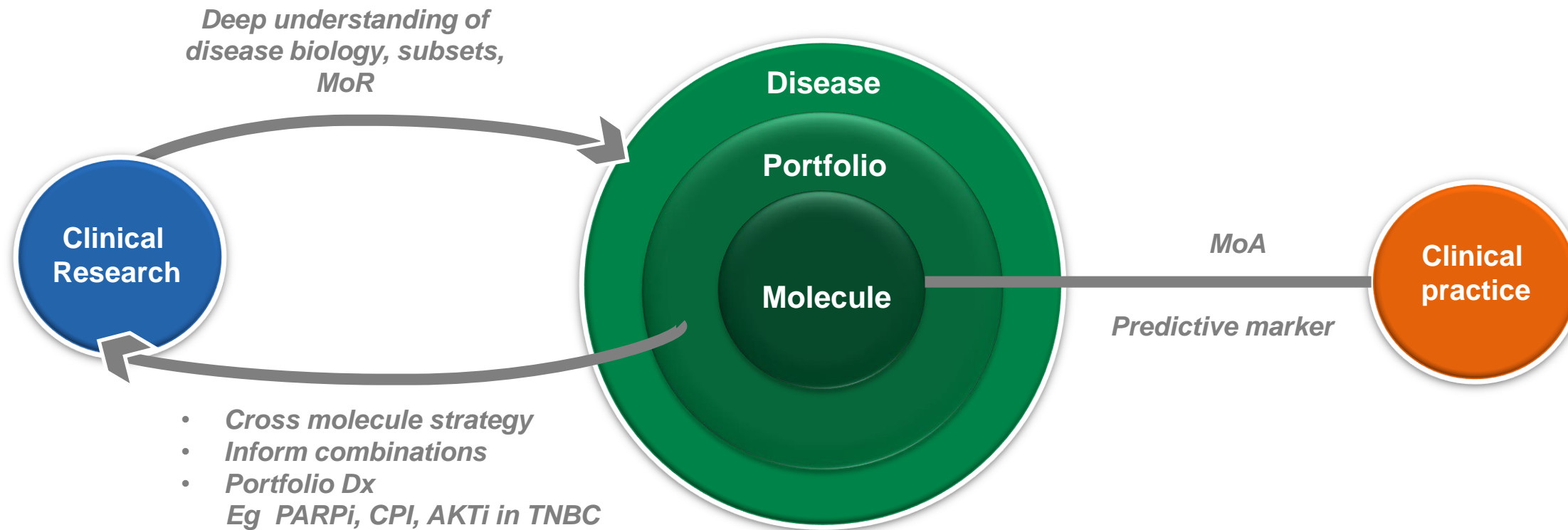
Genentech

Priti Hegde, PhD

**CIT Franchise Biomarker Lead
Genentech**

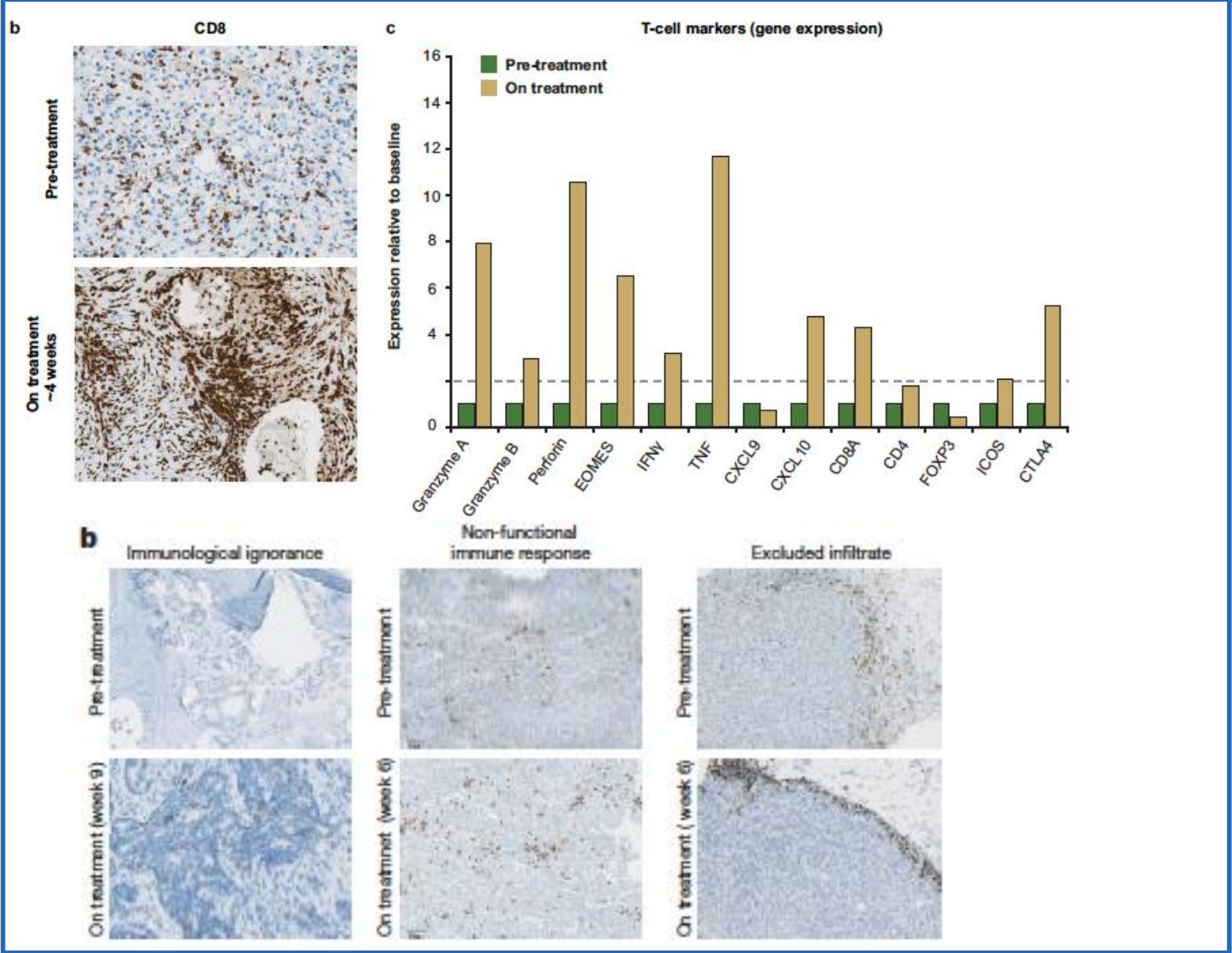
**SITC BMW
San Francisco, May 2018**

Genentech Biomarker Program

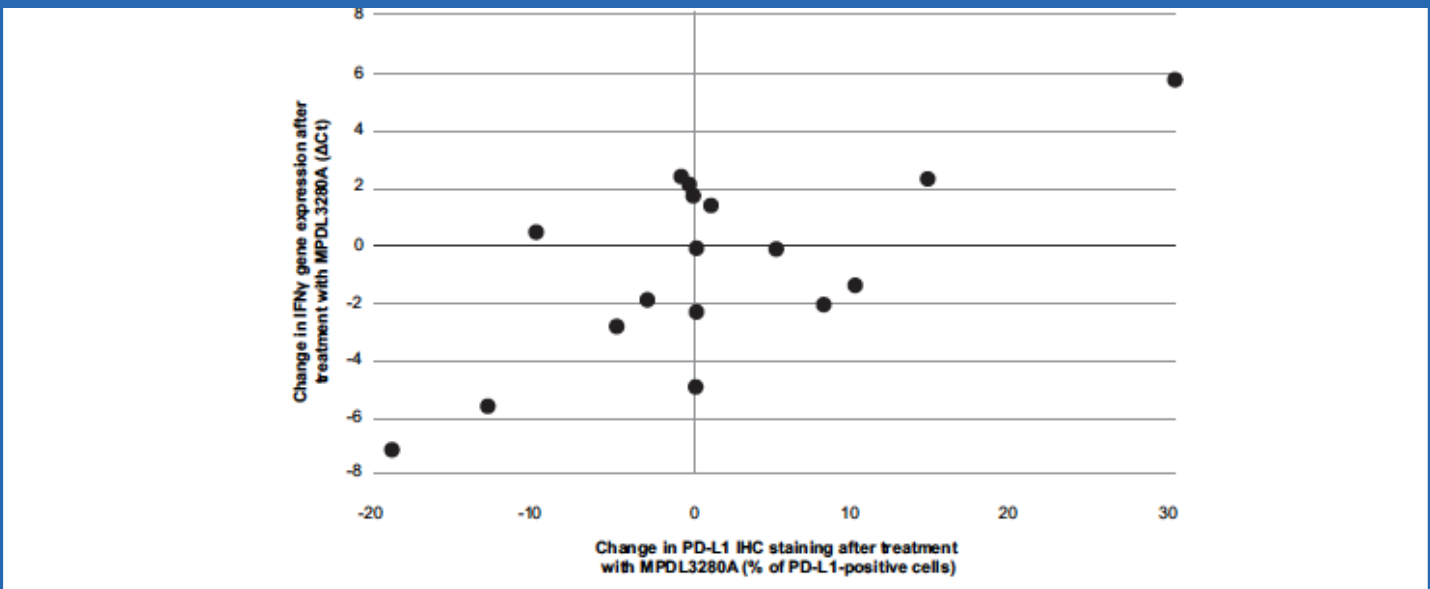


Molecule MoA- the science

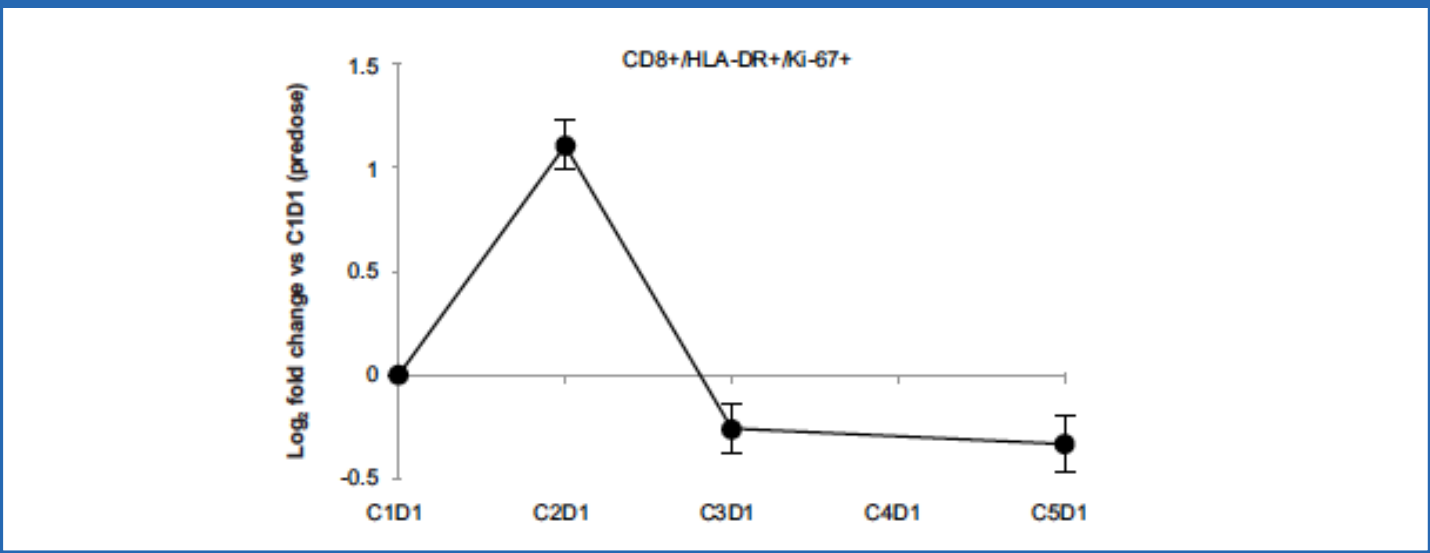
Spatial and temporal changes in infiltrating CD8+ T-cells upon atezolizumab tx



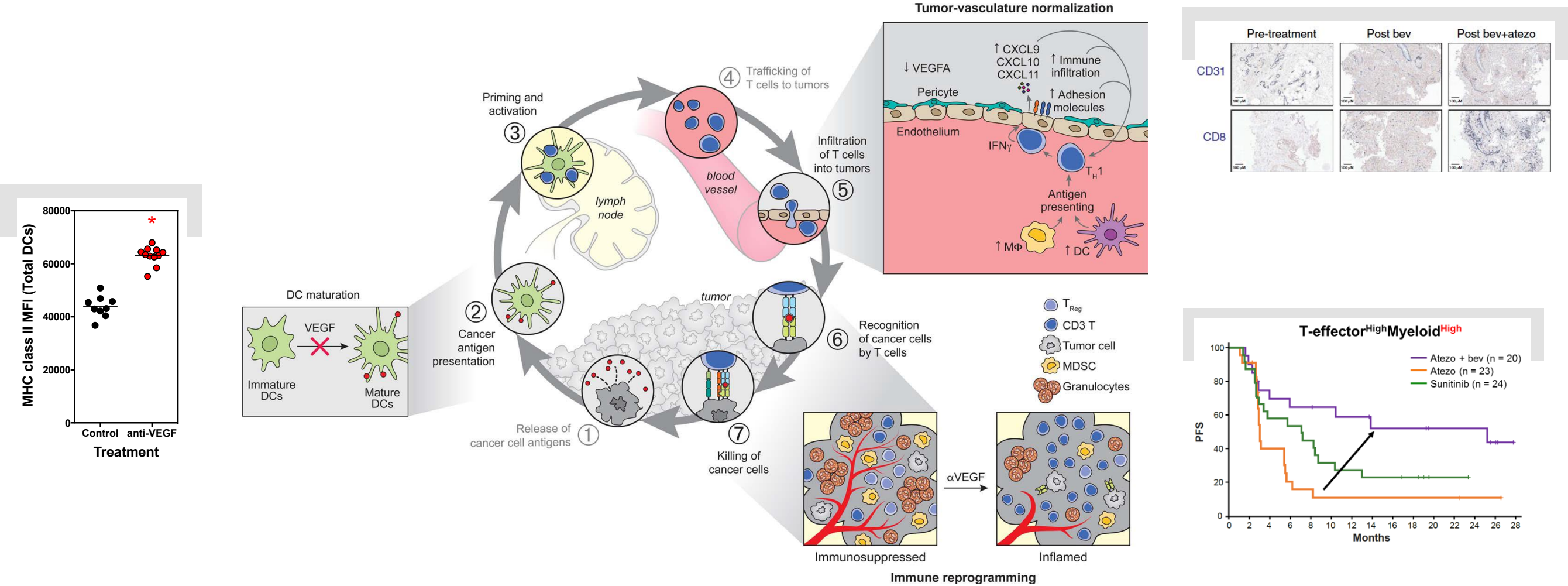
IFN γ mediated adaptive regulation of tumor PD-L1 upon atezolizumab tx



Acute rise in proliferating T-cells and associated cytokines upon atezolizumab tx



Combinations MoA- Portfolio perspective



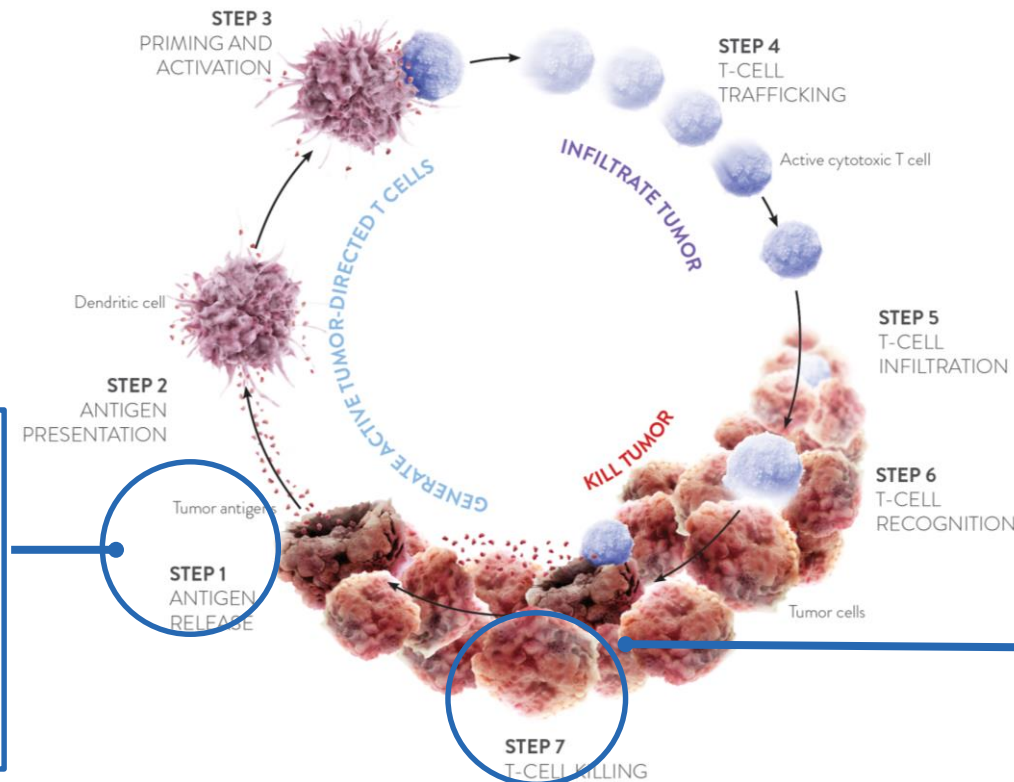
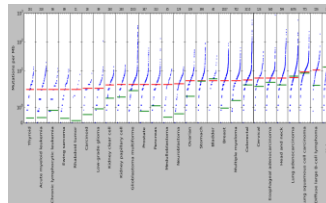
Understanding the immune modulatory properties of a-VEGF have guided the regimens for IMpower150, IMmotion151 and IMbrave151

Predictors of response- shape clinical practice

SIGNAL 1

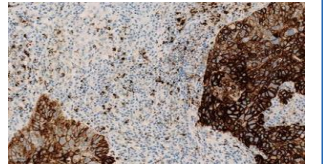
Tumor Mutation Burden

MSI¹, tMB²



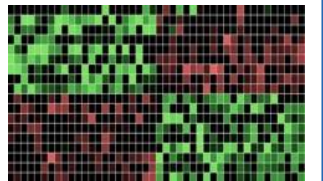
PD-L1 expression

IHC



Tumor Gene Expression
IFN γ signature

NGS



SIGNAL 2

Chen and Mellman, *Immunity*, 2013

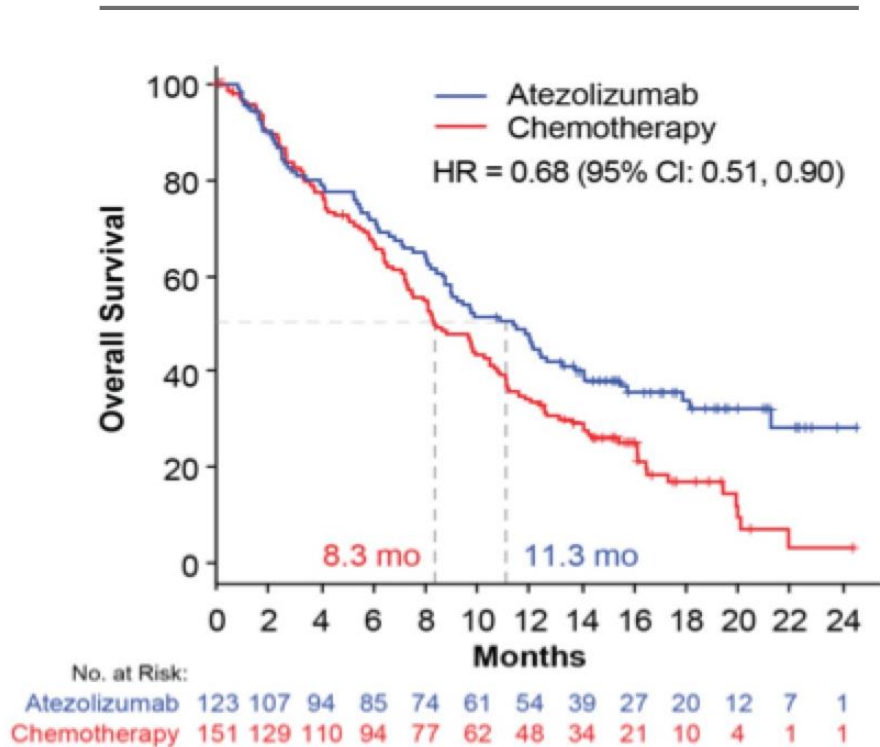
¹Le et al., *NEJM* 2015

² Powles T et al., *Lancet* 2017

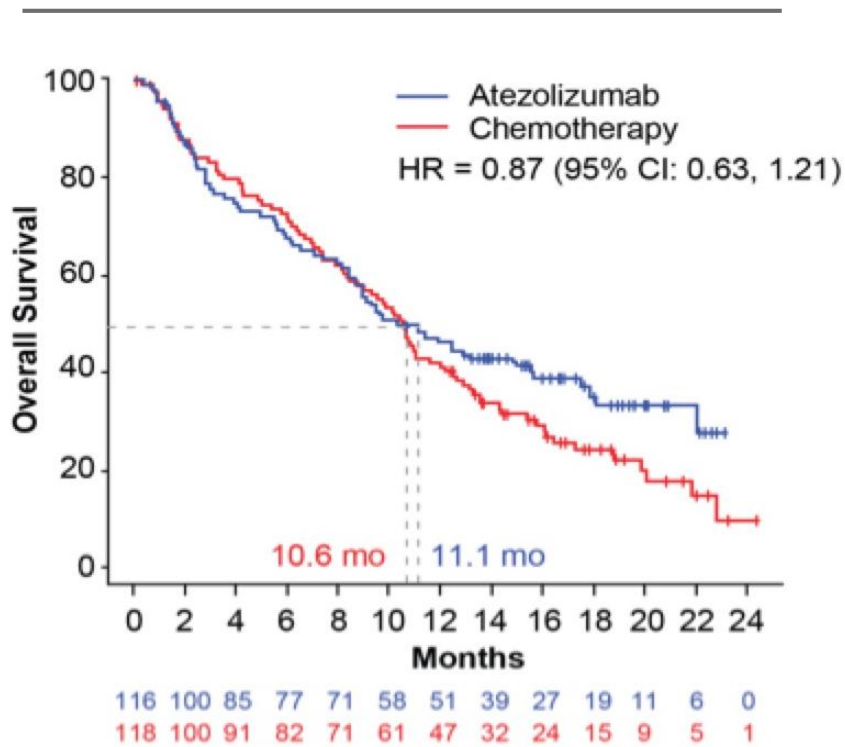
PD-L1 IHC and TMB are predictors of clinical benefit

Example: Ph3 IMvigor211 study in mUC

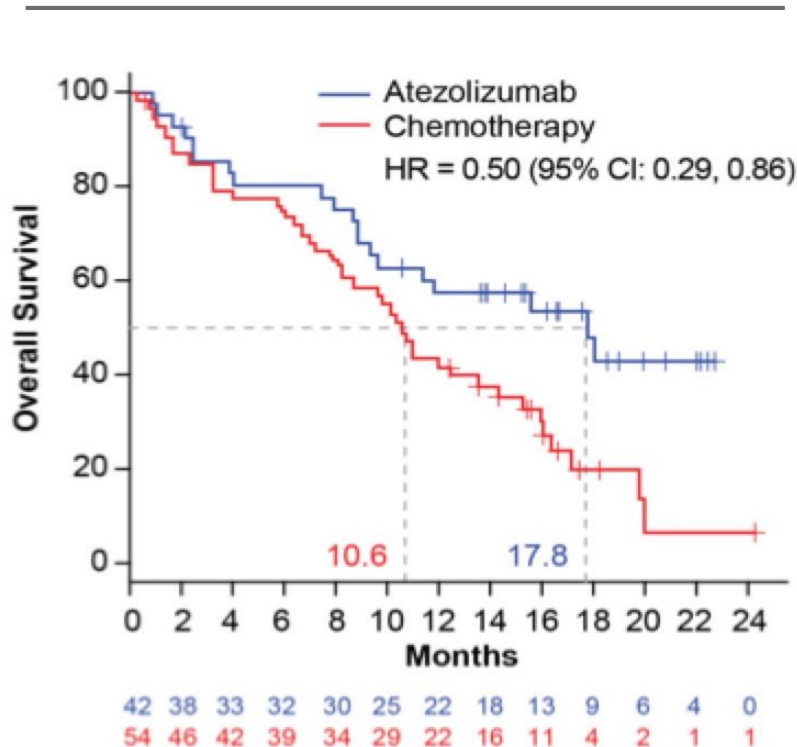
Signal 1
TMB high^a



Signal 2
PD-L1 IC2/3



Signals 1 + 2
TMB high and PD-L1 IC2/3



Improved OS benefit was observed in patients with high TMB as well as high PD-L1 IC scores

ASCO GU 2018, Powles T
Unstratified HRs are displayed. ^aMedian scores were used to define assessment cutoffs: TMB-high (≥ median) or TMB-low (< median). Median TMB in the biomarker-evaluable population was 9.65 mutations/Mb.

Disease tumor immune biology

inflamed

non-inflamed

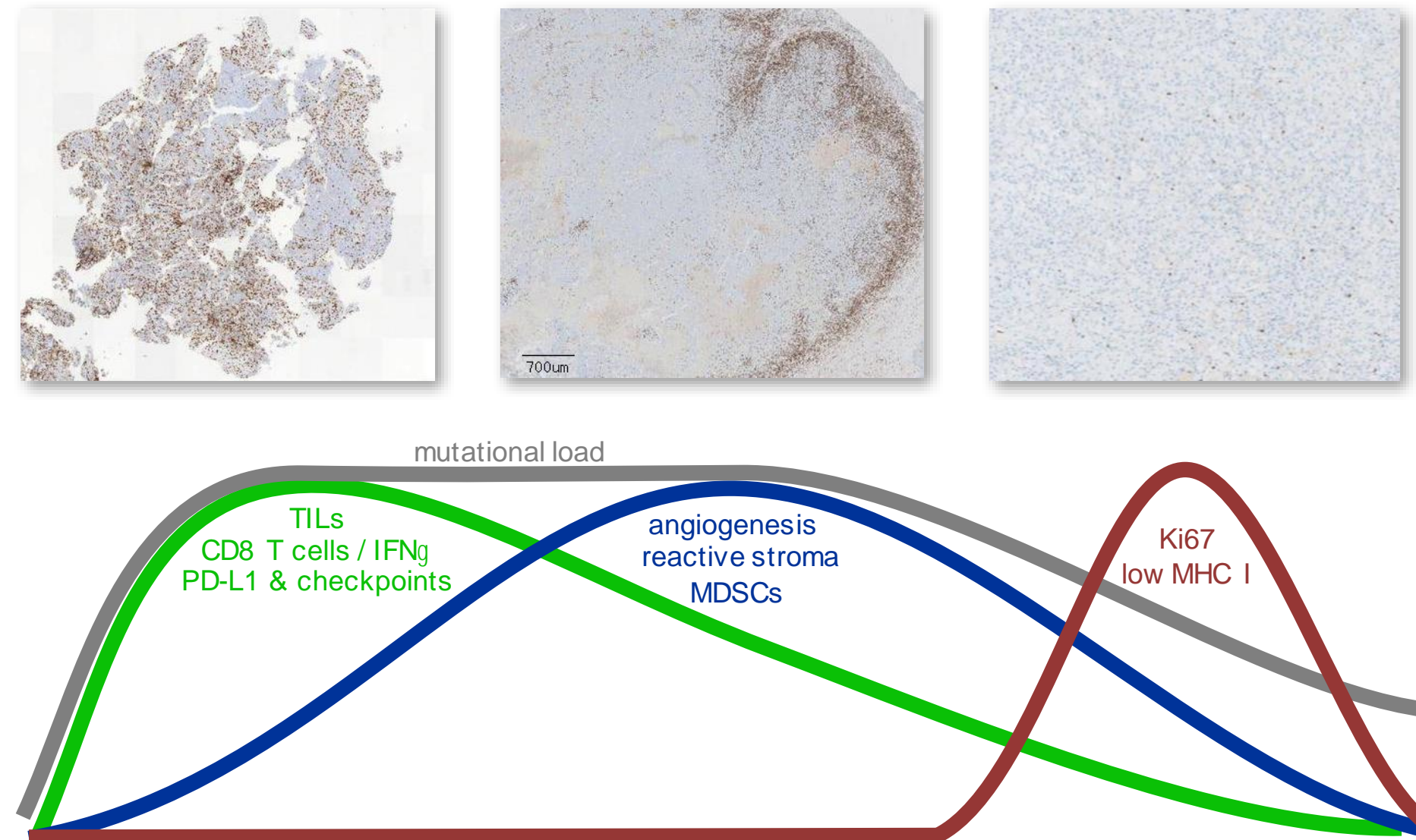
infiltrated

excluded

desert

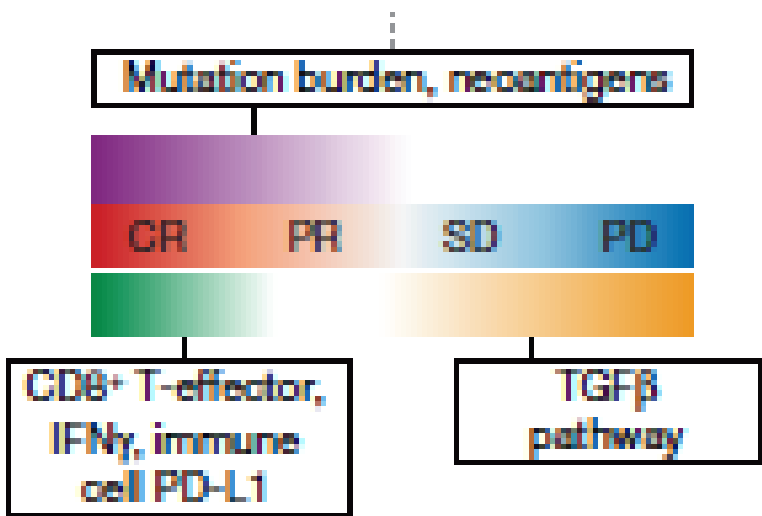
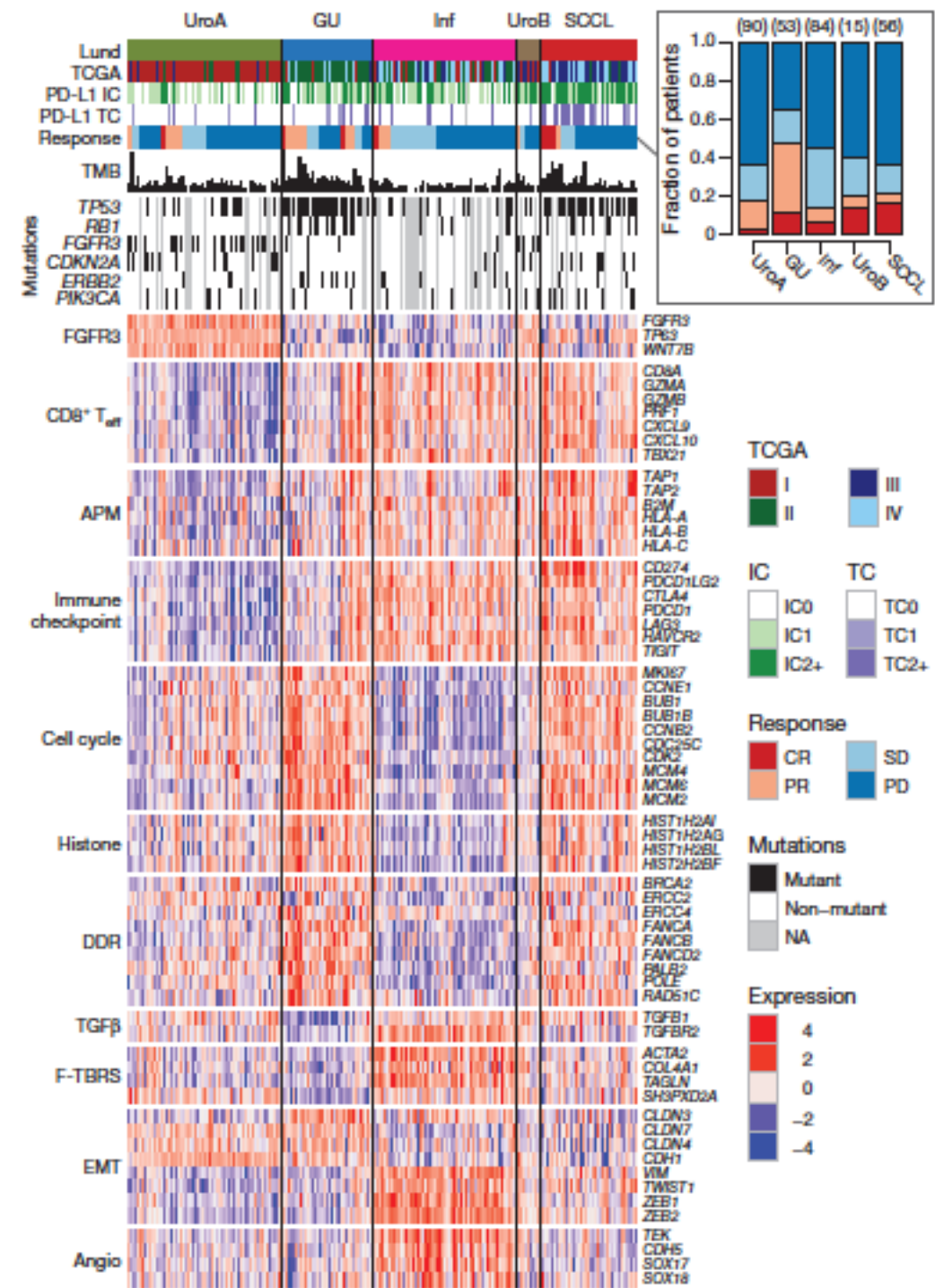
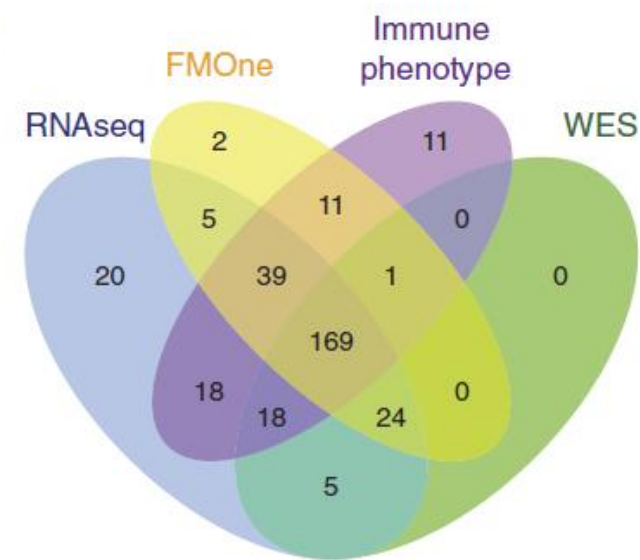
Leveraged archival tissue RNA and CD8 IHC data from legacy Phase III trials (distinct diseases) and clinical outcomes data to develop a deeper understanding of disease immune biology and prognostic significance

- Disease strategy
- Clinical strategy
- Science



Impact of molecule on disease understanding

IMvigor210
(Atezo single agent Phase 2 trial): n = 326



Clinical trial biomarker research

- What is the hypothesis being tested?
- Is the study designed to address the question?
- Can the biomarker question be addressed in a multi-institution study?
 - eg fresh tissue shipments
- Is the biomarker going to enable a clinical decision?
- Is the assay qualified (non primary ep) or validated (primary/co-pri ep)?
- Preferable to use consistent tools and platforms to address questions within and across trials
- Pre-analytical variables may have a greater impact eg RNA/DNA isolation, PBMC prep
- Preferable to execute a biomarker analysis plan (BAP) signed off by biostats prior to data analysis (adds statistical rigor to hypothesis testing)
 - How is the biomarker defined (eg IFNg signature constitutes x,y,z)
 - What is the cutoff used?
 - What is the expected outcome?
- To enable decisions, independent validation required

Clinical trial specimen collections

Global trials

- Multiple sites (typically 10-100 per trial)
- Real time sample shipment (eg blood FACS)
- Real time data turnaround (eg PD-L1, TMB, MSI)

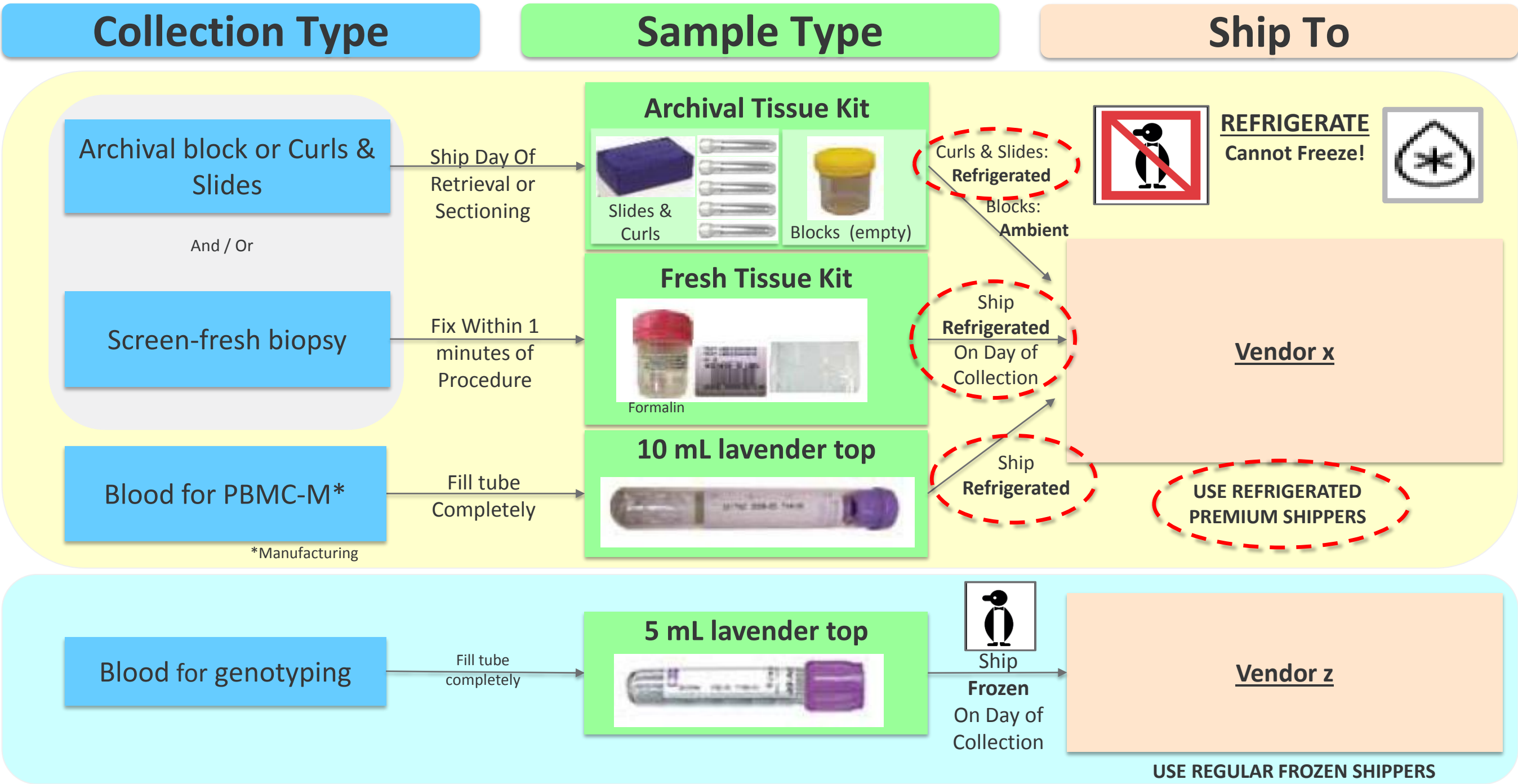


Centralized testing

- Vendor capability
- Assay performance
- Data quality monitoring



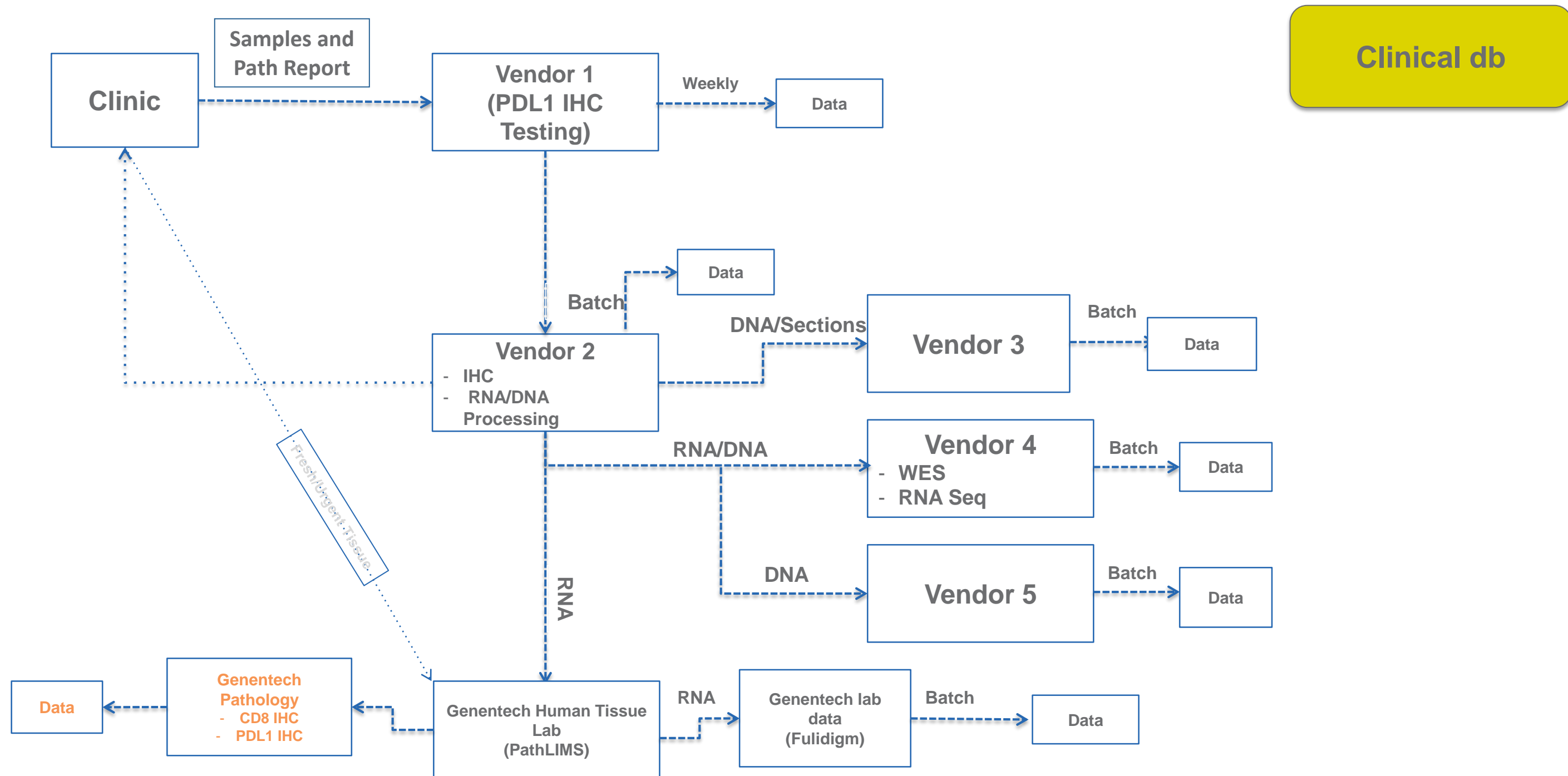
Proper Sample Handling is crucial to executing on biomarker strategy



Specimens from trials

- Informed consent from the patient to conduct specific research
- Sections usually cut and block sent back to patient within 1 month of receipt
- Managed by trial steering committee
- Not up to sponsor to disseminate to third parties

Example of Single Tissue Flow Lifecycle



Data Management

Tissue tracker

- Mechanism to match IDs (pre-screen, screen, different vendors IDs for data)
- Real-time assessment of tissue quality, how much plasma left, how much tissue

Raw data tracker

- Omics
- Digital path
- FACS
- Plasma cytokines

Analyzed data tracker

- Ppts
- PDFs
- Spotfire files, jmp, codes for analysis etc

Data sharing

Phase I trials

At time of publication, data contributing to publication is shared as per journal requirements

Phase II learning trials

GNE policy is to share data to enable learning on certain trials [IMvigor 210 (mUC), IMmotion 150 (RCC)]

Eg Mariathasan S et al., Nature 2018

- RNAseq, WES, CD8 IHC, PD-L1 data made publicly available with outcomes data for all 326 patients enrolled in the trial. Easily downloadable.
- Allows TCGA-like learning with clean outcomes data to atezolizumab

Phase III trials

Typically not data rich

Sharing of data contributing to publication as per journal requirements

Investigator-initiated trials- All about authorship!



- Publication philosophy
- Biomarker statement of work developed and signed between institutions
- Clarity of who does what and timelines for data generation
- Data sharing philosophy within group of investigators
- Generally select CRO for routine data (eg CD8 IHC, PD-L1, RNAseq, DNAseq)
 - Inst. contract with CRO
- Analysis plan depends on expertise within the consortium

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