

# CDDF Multi-stakeholder Workshop

## Measurable Residual Disease (MRD) and Circulating Tumour Nucleotides (ctDNA)

Axel Glasmacher  
(CDDF, DE)

# Disclaimer

- I am a board member and the treasurer of the CDDF
- I have no financial conflicts related to the workshop's topics

# CDDF Multi-Stakeholder-Workshops on MRD

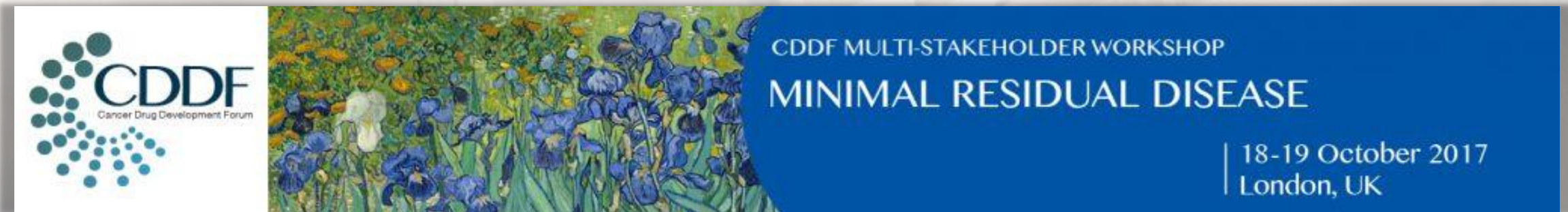



**BDA**  
WORKSHOP

MINIMAL RESIDUAL DISEASE:  
END-POINTS IN CLINICAL TRIALS

13-14 MAY 2014 | LONDON, UNITED KINGDOM

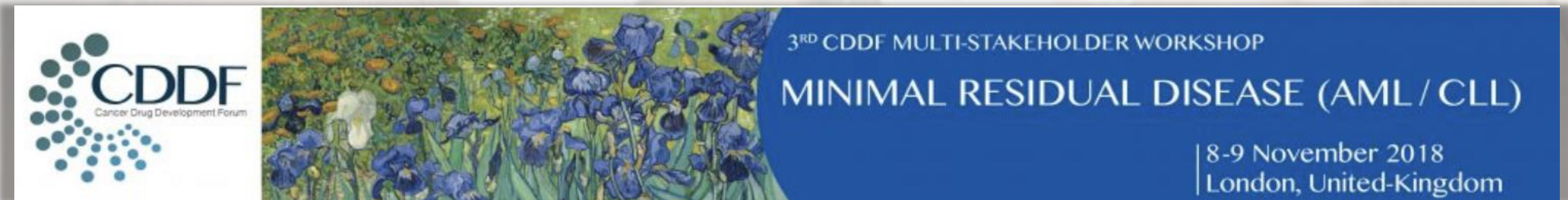
**BDA**  
BIOTHERAPY DEVELOPMENT ASSOCIATION




 CDDF  
Cancer Drug Development Forum

CDDF MULTI-STAKEHOLDER WORKSHOP  
MINIMAL RESIDUAL DISEASE

18-19 October 2017  
London, UK



 CDDF  
Cancer Drug Development Forum

3<sup>RD</sup> CDDF MULTI-STAKEHOLDER WORKSHOP  
MINIMAL RESIDUAL DISEASE (AML / CLL)

8-9 November 2018  
London, United-Kingdom



## Scientific Programme Committee



**Axel Glasmacher**  
CDDF, DE



**John Smyth**  
CDDF, UK



**Veerendra  
Munugalavadla**  
AstraZeneca, US



**Hans Scheurer**  
Myeloma Patient Europe, NL



**Natalie Dimier**  
Roche, UK



**Reshma Patel**  
Johnson & Johnson, UK

# Speakers, Moderators and Panelists

In Order of Appearance

**Bruno Paiva (Univ. de Navarra, ES)**  
**Dominic Rothwell (Manchester, UK)**  
**Jürgen Gschwend (TU Munich, DE)**  
**Marie Morfouace (EORTC, BE)**  
**Veerendra Munugalavadla (AstraZeneca, US)**  
**Pierre Demolis (ANSM, FR)**  
**Natalie Dimier (Roche, CH)**  
**Reshma Patel (Johnson & Johnson, UK)**  
**Christopher Hourigan (NIH, USA)**  
**Jeff Allen (Friends of Cancer Research, USA)**

**Nicole Gormley (FDA, USA)**  
**Natalie Dimier (Roche);**  
**Paula van Hennik (MEB, NL)**  
**Shirley Hopper (MHRA, UK)**  
**Donna Roscoe (FDA, USA)**  
**Claudia Popp (Roche, CH)**  
**Hans Scheurer (MPE, NL)**  
**Carole Longson (NICE, UK)**  
**Darren Hodgson (AstraZeneca, UK)**

# Definitions

- **MRD**

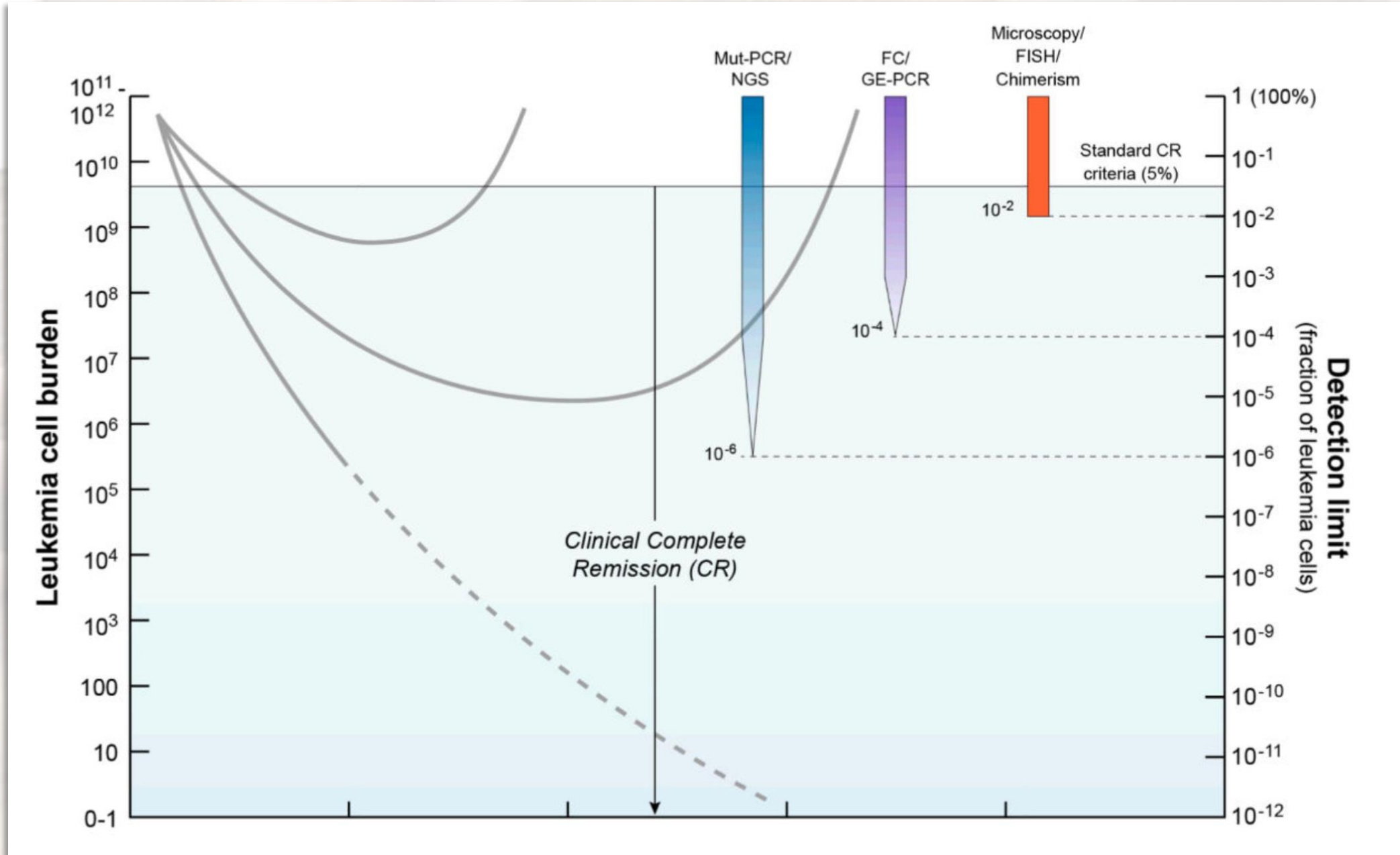


- Measurable (previously termed minimal) residual disease
- Usually analysed in bone marrow samples
- Detection of malignant cells down to levels of  $1:10^4$  to  $1:10^6$  white blood cells, compared with 1:20 in morphology-based assessments

- **ctDNA**

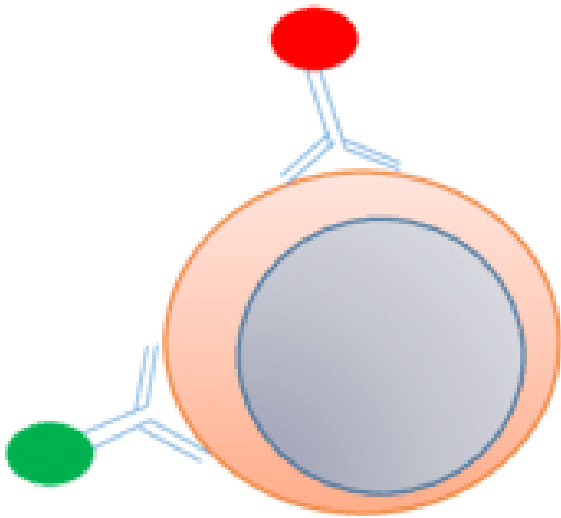


- Circulating (cell-free) tumour DNA (nucleotides)
- Usually analysed in peripheral blood samples
- Allows to detect/track tumour-specific nucleotides ('liquid biopsy') to determine e.g. response or relapse of a tumour

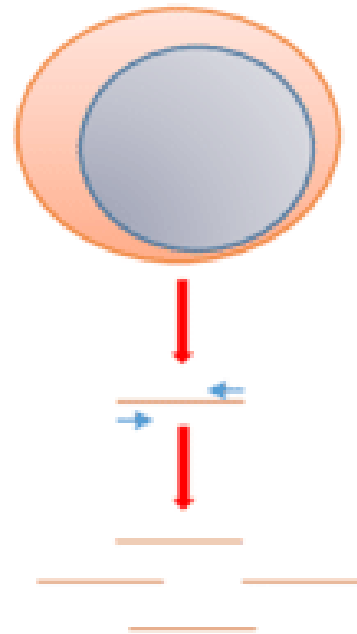


# MRD Modalities

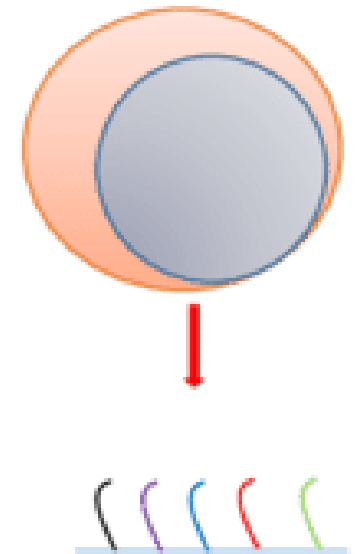
Multi-parameter  
flow cytometry



Quantitative  
PCR



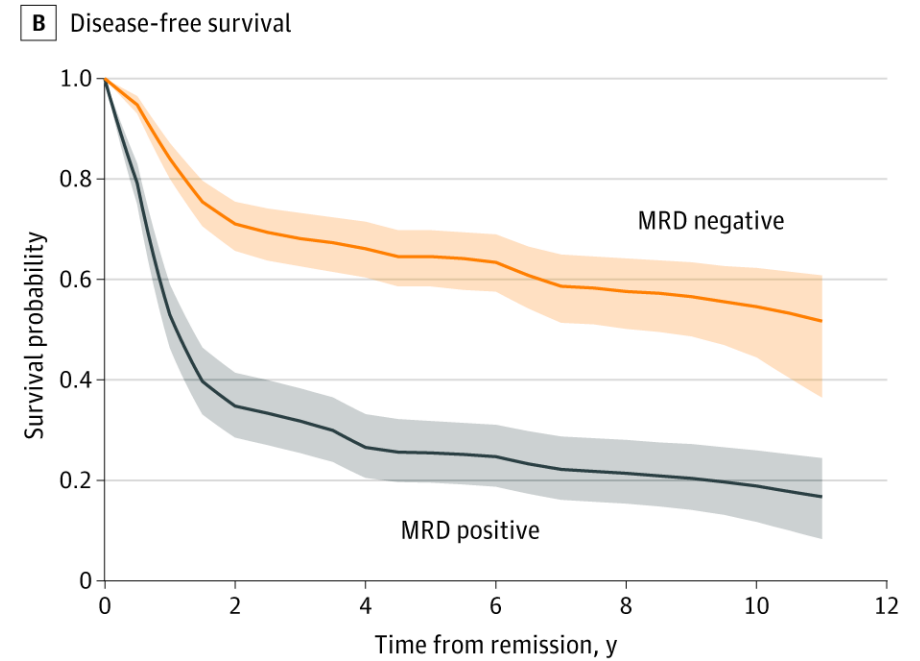
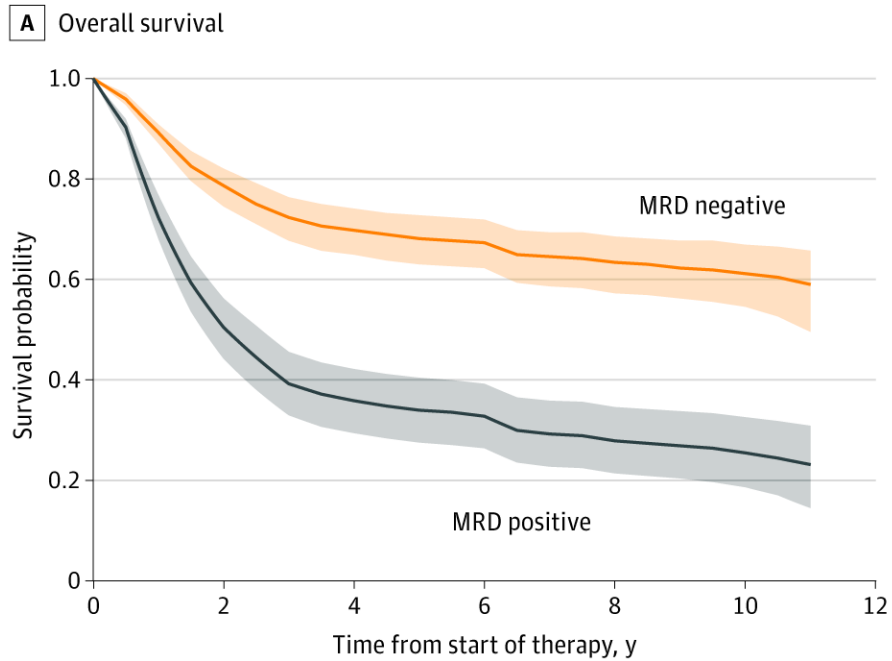
Next generation  
sequencing





# Association of MRD with survival outcomes in AML

## A Systematic Review and Meta-analysis in 11,151 patients



Short NJ, et al. JAMA Oncol. 2020;6(12):1890-1899.

# MRD assessment across some hematological malignancies

Application depending on clinical need and methodological robustness

	ALL	AML	CML	CLL	MM
Complexity	Intermediate	High	Low	Intermediate	Intermediate
Standardization	High	Low	High	Low	Intermediate
Clinical trials	Yes	Yes	Yes	Yes	Yes
Routine practice	Yes	Yes	Yes	Infrequent	Intermediate
Treatment decisions	Yes	Yes	Yes	Infrequent	Infrequent

# Definitions

- **MRD**



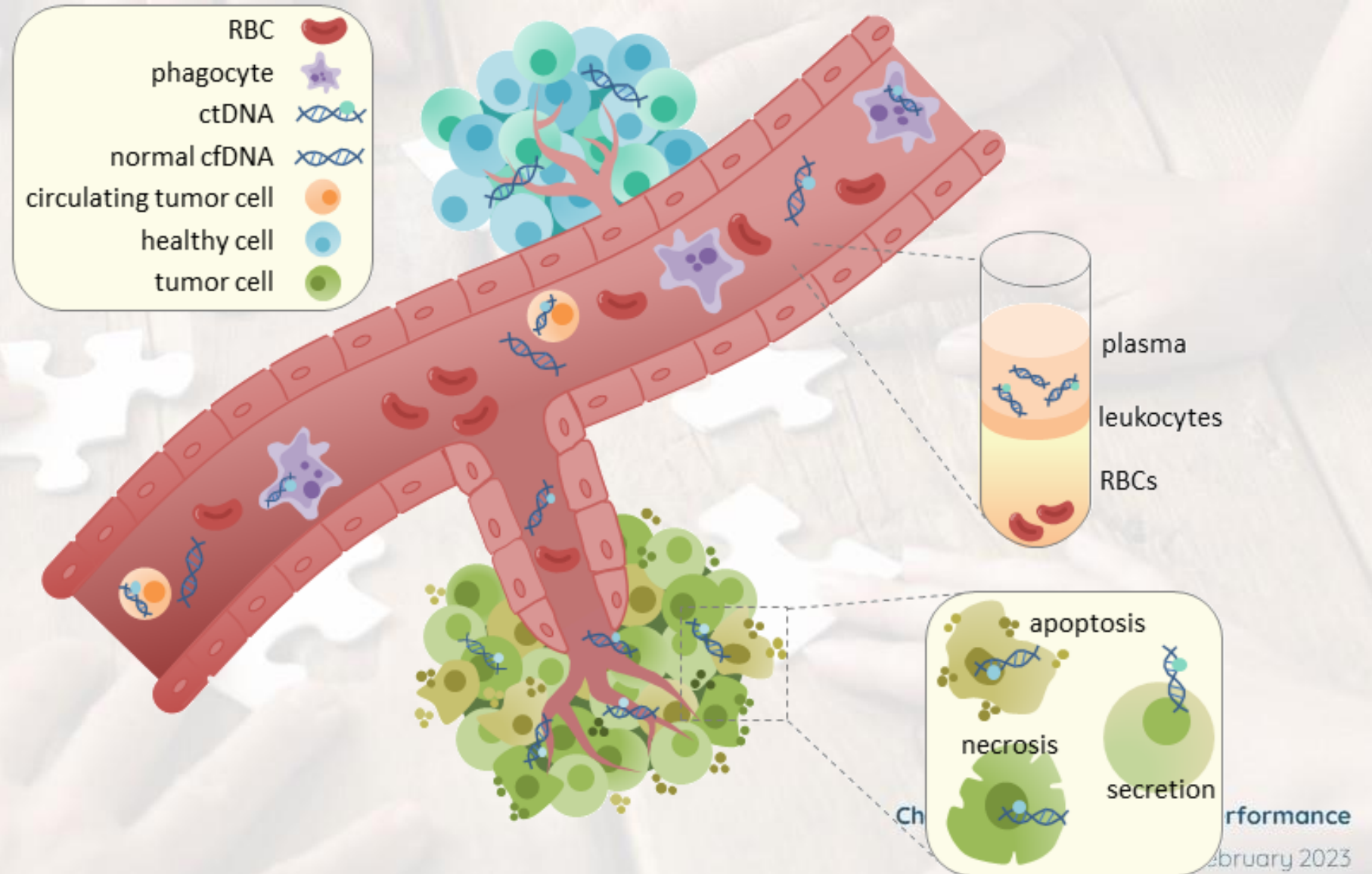
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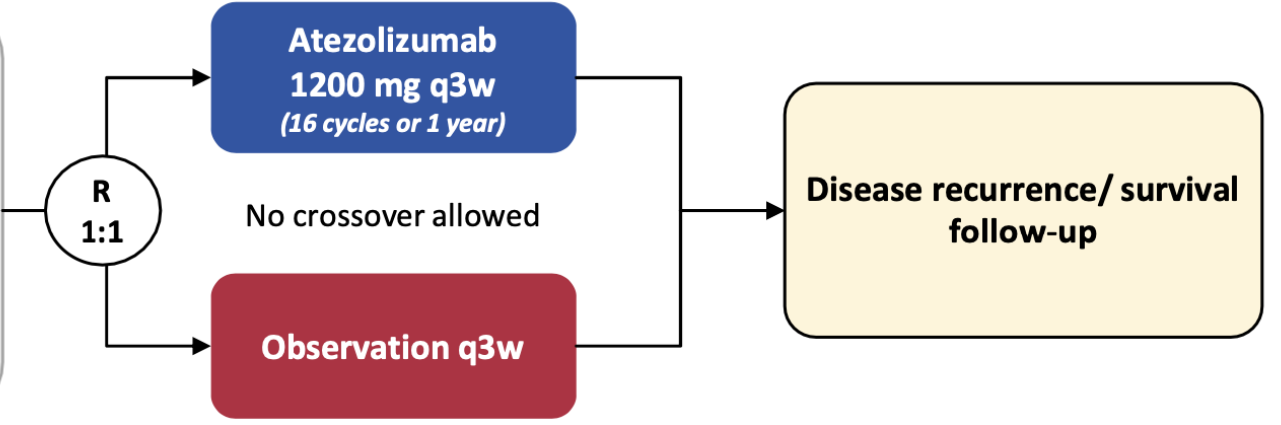
# circulating tumor DNA (ctDNA)



# Phase 3 IMvigor010 adjuvant study in MIUC

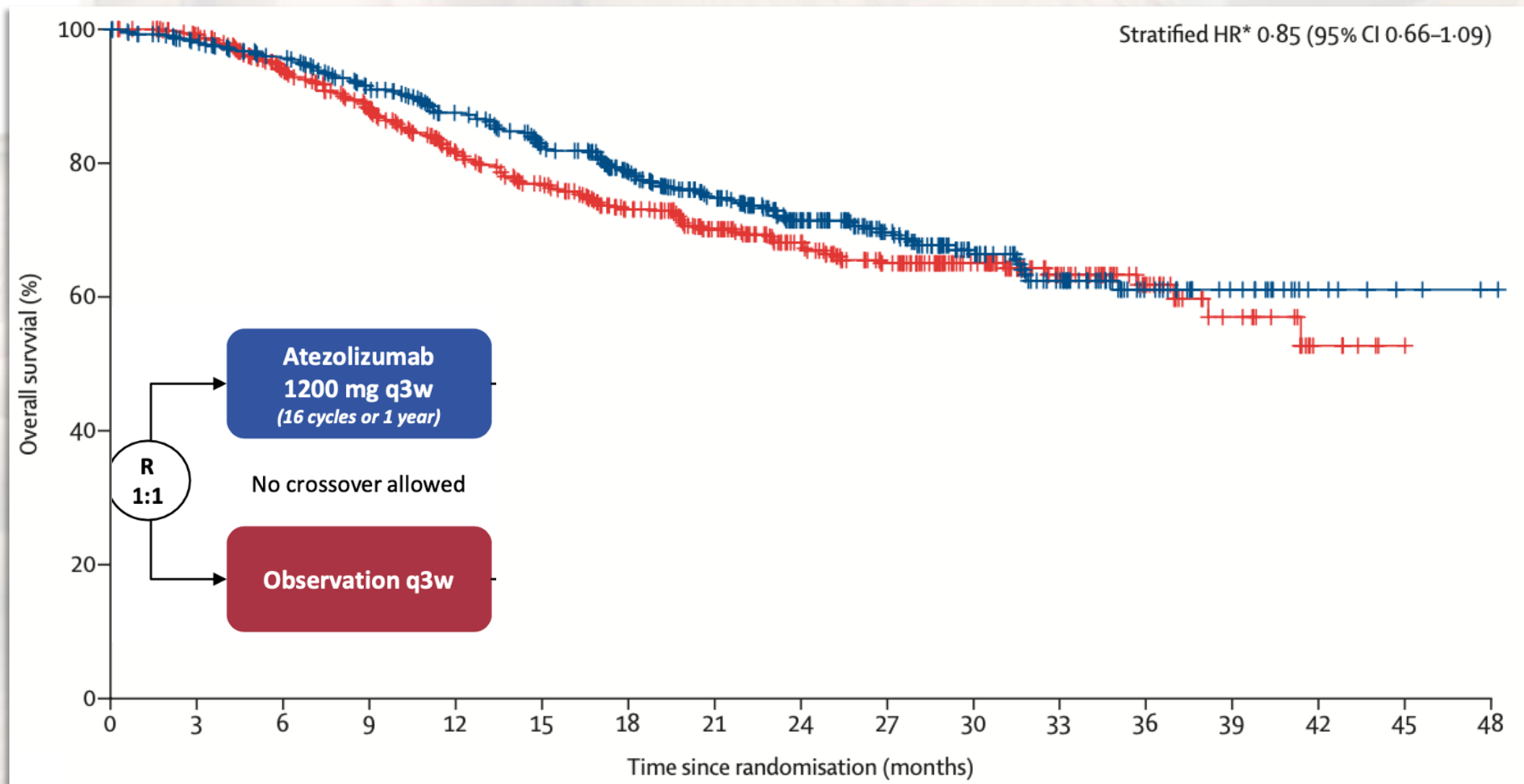
- Key eligibility**
- High-risk MIUC (bladder, renal pelvis, ureter)
  - Radical surgery with lymph node dissection within  $\leq 14$  weeks<sup>a</sup>
  - No postsurgical radiation or AC
  - ECOG PS 0-2
  - Tissue sample for PD-L1 testing

- Stratification factors**
- |                             |                                  |
|-----------------------------|----------------------------------|
| • PD-L1 status <sup>b</sup> | • # of lymph nodes resected      |
| • Nodal status              | • Prior neoadjuvant chemotherapy |
| • Tumor stage               |                                  |

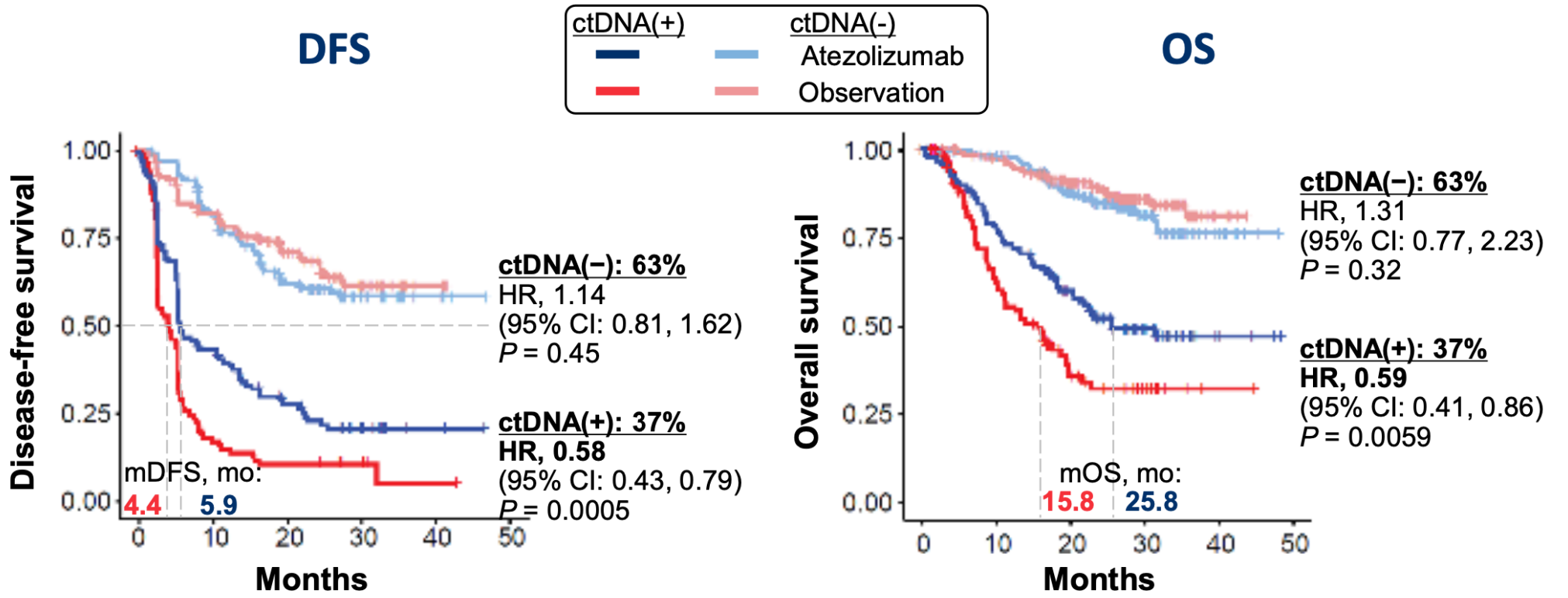


- **Primary endpoint:** DFS (ITT population)
- **Key secondary endpoint:** OS (ITT population)
- **Safety**
- **Exploratory analyses:** Predictive, prognostic and pharmaco-dynamic biomarkers in tumor tissue and blood and their association with disease recurrence<sup>c</sup>

J. Geschwend: Presentation at CDDF Workshop, 2022  
Bellmunt J et al., Lancet Oncology 2021; 22: 525

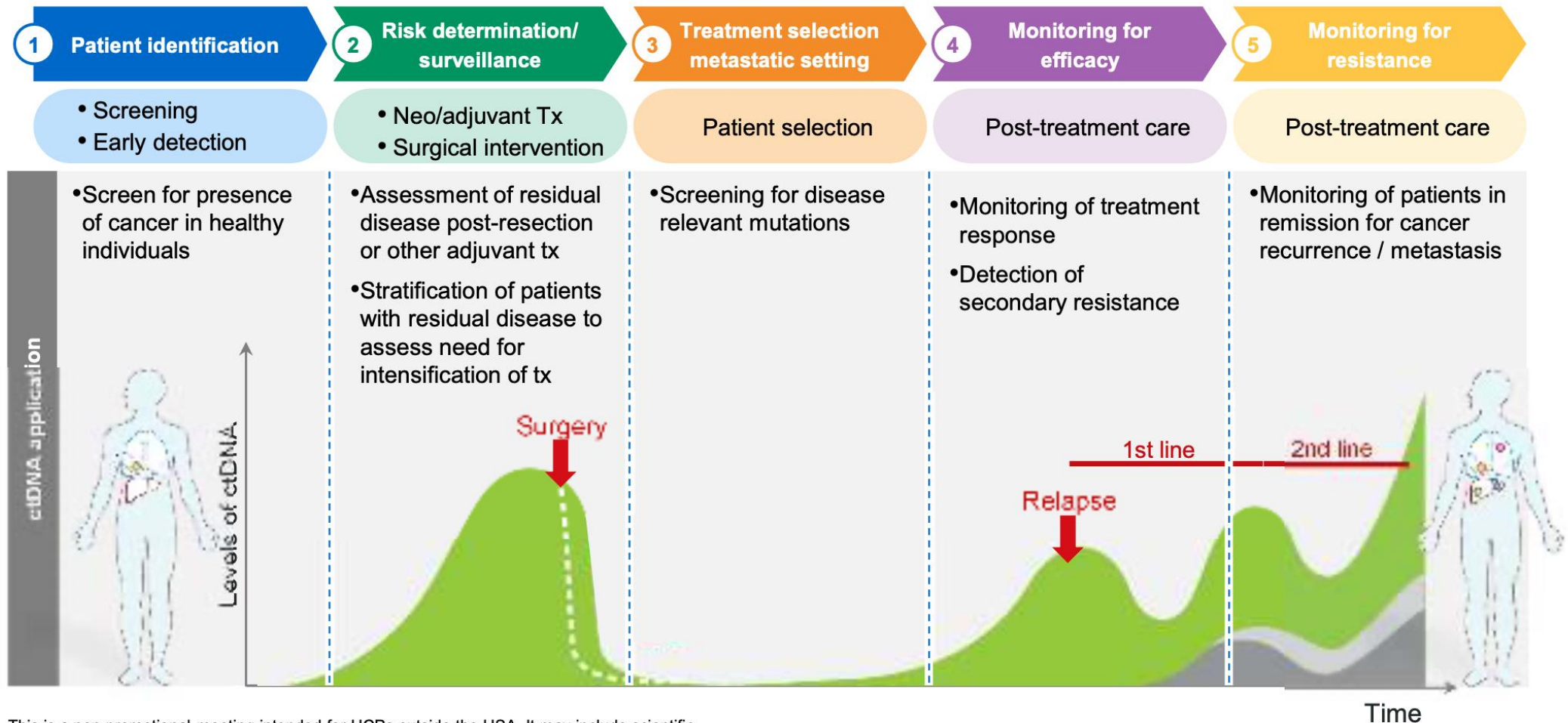


# ctDNA(+) patients had improved DFS and OS with atezolizumab vs observation



- IMvigor010 confirmed the prognostic value of ctDNA status

# ctDNA has substantial potential to be used throughout the patient journey



This is a non-promotional meeting intended for HCPs outside the USA. It may include scientific information about investigational compounds that may not be approved or valid in your jurisdiction

Adapted from Wan et al. Nat Rev Cancer 2017



# FDA: Potential uses of MRD and ctDNA

- Prognostic Biomarker
- **Clinical Uses**
  - Screening/Early Detection
  - Monitor for relapse
  - Guide therapeutic decisions
- **Regulatory Uses**
  - Patient Stratification
  - Patient Selection/Enrichment
  - Risk-based treatment assignment (Escalation / De-Escalation)
  - Intermediate Endpoint or Surrogate Endpoint

# FDA: Potential uses of MRD and ctDNA

- Accelerated Approval
  - is based on an effect on a surrogate endpoint or an intermediate clinical endpoint that is reasonably likely to predict clinical benefit
- **Meta-analytical methods - Patient-level data**
  - Individual Level Surrogacy
    - Correlation between candidate surrogate and true clinical endpoint on an individual level
  - Trial Level Surrogacy
    - Correlation between effect of treatment on the candidate surrogate and the effect of treatment on the true clinical endpoint
  - Surrogate Threshold Effect
    - Minimum treatment effect on the surrogate necessary to predict an effect on the true clinical endpoint

# FDA: Potential uses of MRD and cDNA

- **Meta-analysis Considerations**

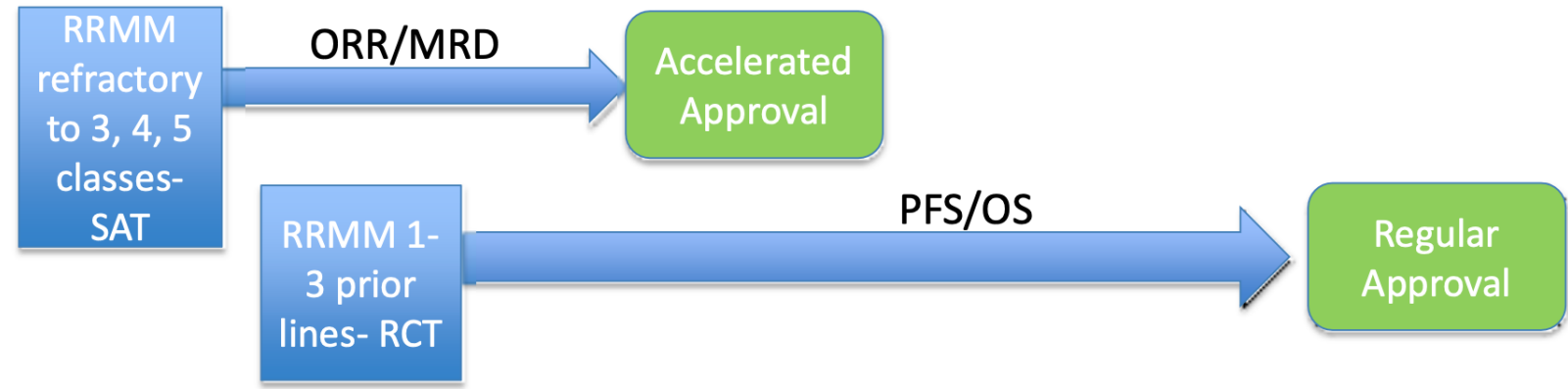
- Inclusion of more trials increases the statistical rigor of the analysis and may allow for more interrogation of the data to address uncertainties.
- Inclusion of trials with a range of treatment effects (positive and negative trials) increases the accuracy and precision of trial level surrogacy assessment.

- **Caveats regarding use of surrogate endpoint**

- Use of surrogate may not be appropriate for subpopulations or future trial populations if there are significant differences between the population in the meta-analysis and the trial population.
- Use of surrogate may not be appropriate for therapeutic modalities that have substantially different MOA (e.g., cytotoxic vs. immunotherapies).

# Drug Development Approach

- Multiple Trial Model



- Single Trial Model



N. Gormley: Presentation at CDDF Workshop, 2022

# Conclusions

- ctDNA and MRD are prognostic biomarkers, but not validated surrogate endpoints
- Existing uncertainty and remaining questions regarding these endpoints for regulatory purposes
- MRD assessments in clinical trials should be discussed with the Agency
- FDA is committed to working with the community on the development of MRD and ctDNA.

# Key Milestones to Date

**2016**



**Duke-Margolis/FDA Public Meeting on MRD**

Janssen invited to present plans to evaluate MRD as an endpoint in AML; other companies invited to create a collaborative program

**2018**



**Alliance formed with two goals:**  
 - Establish MRD as surrogate endpoint for AML  
 - Harmonise MRD testing



**2019**



**Engagement with Health Authorities to design surrogacy analysis and discuss assay methodology**

Initial discussions with the Agencies provided important feedback to help MPAACT design a meta-analysis for surrogacy

**2020-2021**



**Substantial progress made towards MPAACT goals**

- Validation of a flow MRD assay
- Advisory boards held to discuss pediatric and market access considerations
- Third-party statistical Vendor, Mayo Clinic, selected to start data collection

**2022 (so far)**

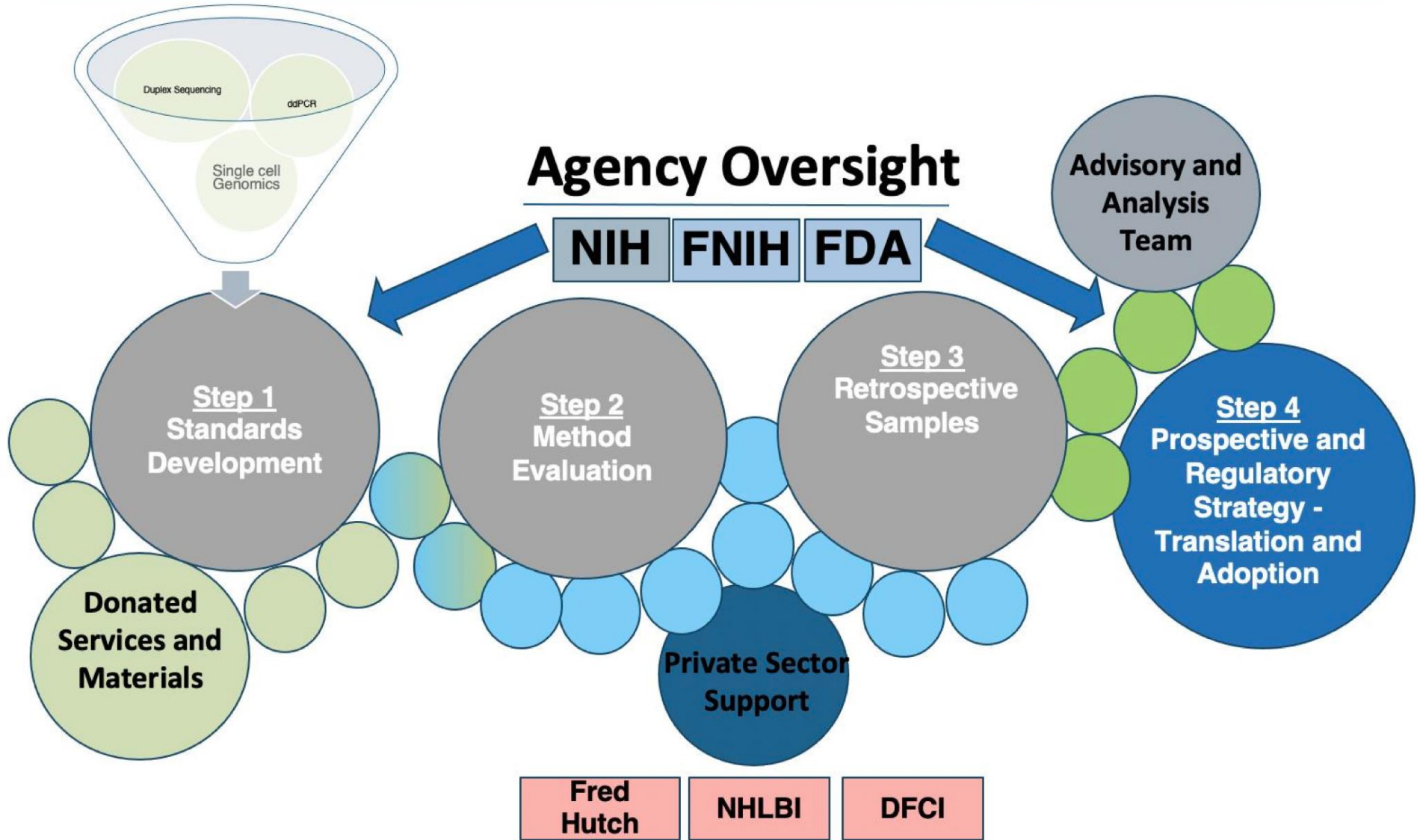


**Inclusion of three new companies to the MPAACT Alliance**

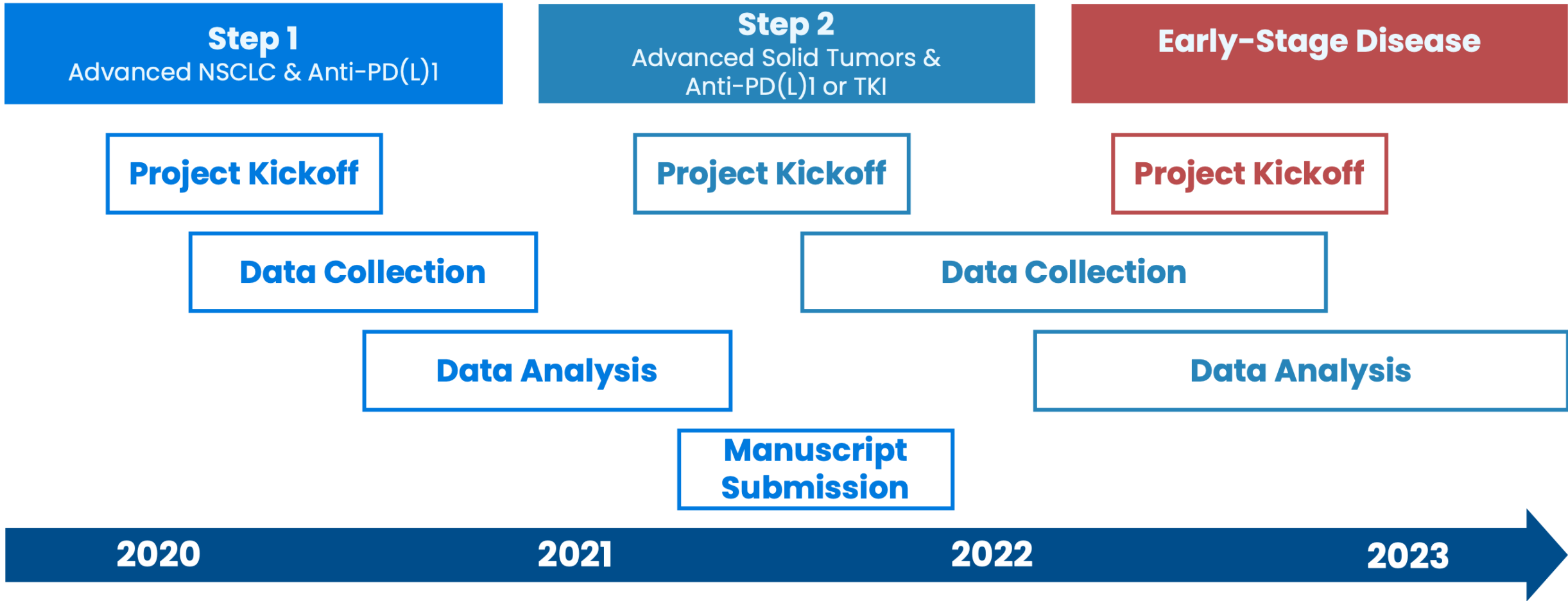
Systematic literature review completed to identify clinical trials to include in the meta-analysis

# MRD in AML Project Team

Year 1 Milestone > Year 2 Milestone > Year 3 Milestone > Year 4 Completion



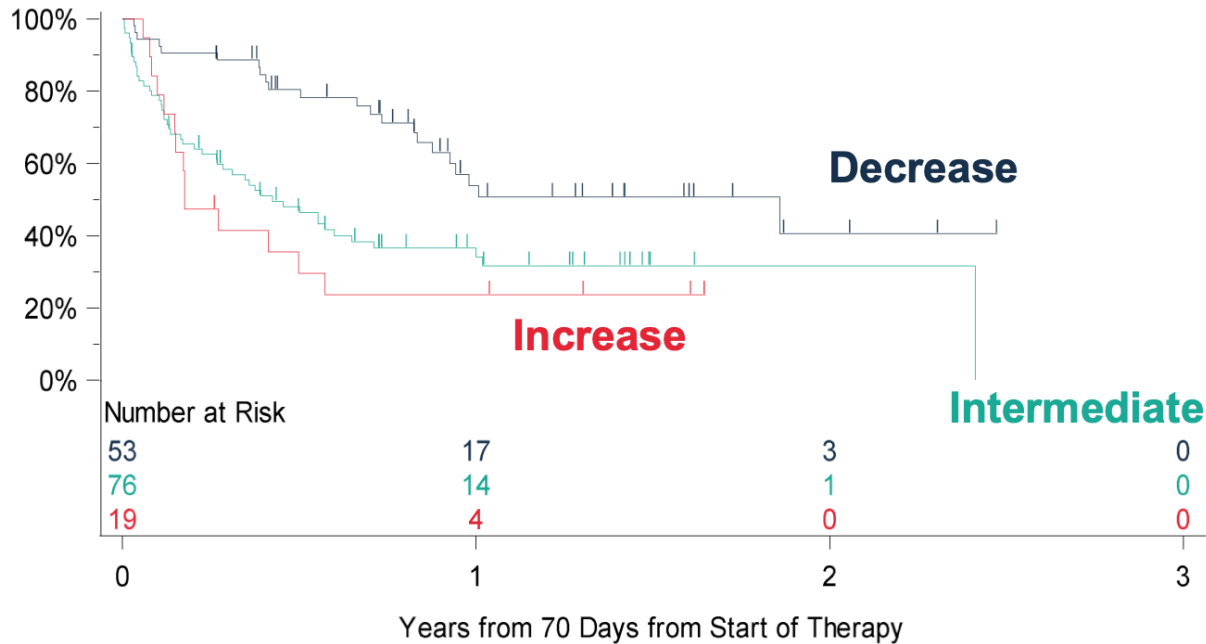
# ctMoniTR Project Timeline





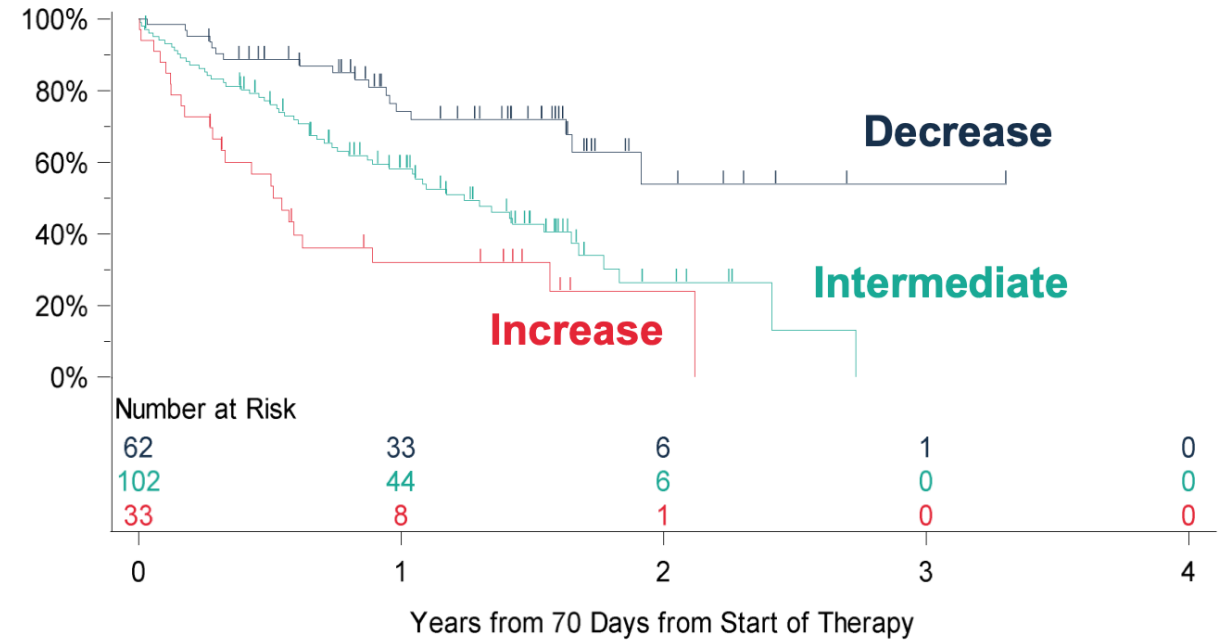
# Robust Association Observed Between Strong Decreases in ctDNA and Patient Survival

## Progression-Free Survival by Max VAF\*



Log-rank Pairwise p-value	Decrease	Intermediate	Increase
Decrease	-		
Intermediate	<b>0.001</b>	-	
Increase	<b>&lt;0.001</b>	0.426	-

## Overall Survival by Max VAF



Log-rank Pairwise p-value	Decrease	Intermediate	Increase
Decrease	-		
Intermediate	<b>&lt;0.001</b>	-	
Increase	<b>&lt;0.001</b>	<b>0.014</b>	-

## Survival Outcomes (3-Level)

Kaplan-Meier Curves

J. Allen: Presentation at CDDF Workshop, 2022

\*Note: patients with progression within 70 days were excluded from the PFS plots.



# Important Perspectives

- **Patient Perspective (Hans Scheuer, MPE)**

- Understand the impact of MRD results in research on patients
- Preference of peripheral blood versus bone marrow samples
- Consider the influence of accelerated approval on patient expectations
- Consider patient input sufficiently early in the development path

- **Health Technology Assessment (Carole Longson, UK)**

- Limited evidence is presented supporting the validity of the relationship between the biomarker surrogate endpoint and endpoints/outcomes of most interest to HTA decision-making: HRQoL and survival
- Whilst expression of the biomarker may be associated with efficacy/survival, it is not usually related to HRQL
- This creates high levels of uncertainty around the real incremental impact of innovative cancer drugs

# Conclusions and Next Steps

- **Methods for MRD and ctDNA:**
  - Further standardisation and quality control
- **Validation for regulatory use:**
  - Endpoint: Demonstration of patient- and trial-level surrogacy
  - Patient selection: Diagnostic or prognostic biomarker (enrichment/stratification)
  - Treatment modification: Escalation or de-escalation of therapy
- **Important next steps:**
  - Continued collaboration towards standardisation and validation (MPAACT, FoCR, NIH)
  - Early inclusion of patient input
  - Models that allow HTA assessment after approval based on surrogate endpoint
  - Improve response assessment and definition by including MRD and ctDNA