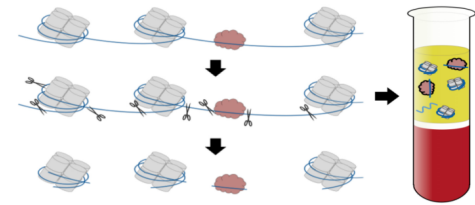
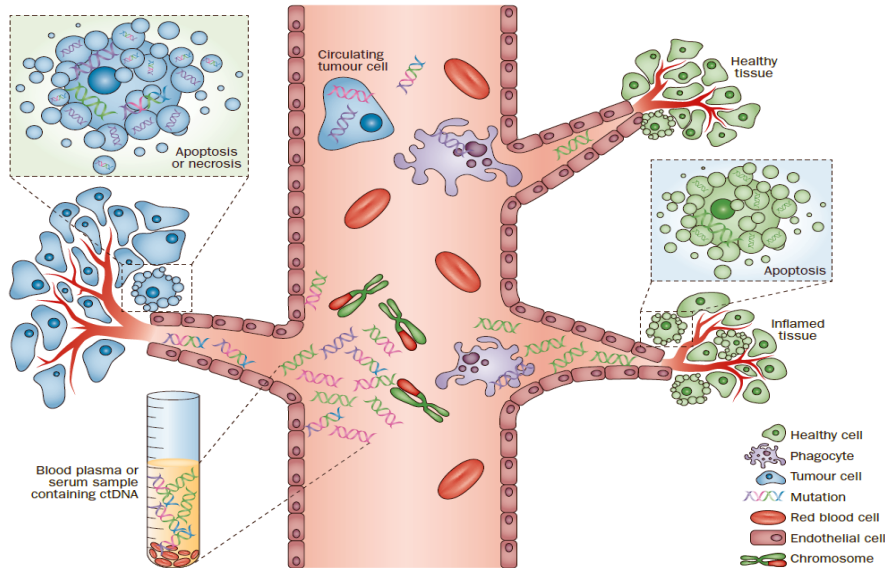


The role of ctDNA to decide postoperative treatment in MIBC: are we ready for prime time?

M. Carmen Mir, MD, PhD, FEBU

Department of Urology; Mount Sinai Icahn School of Medicine, Elmhurst/Queens Hospitals

What is ctDNA?



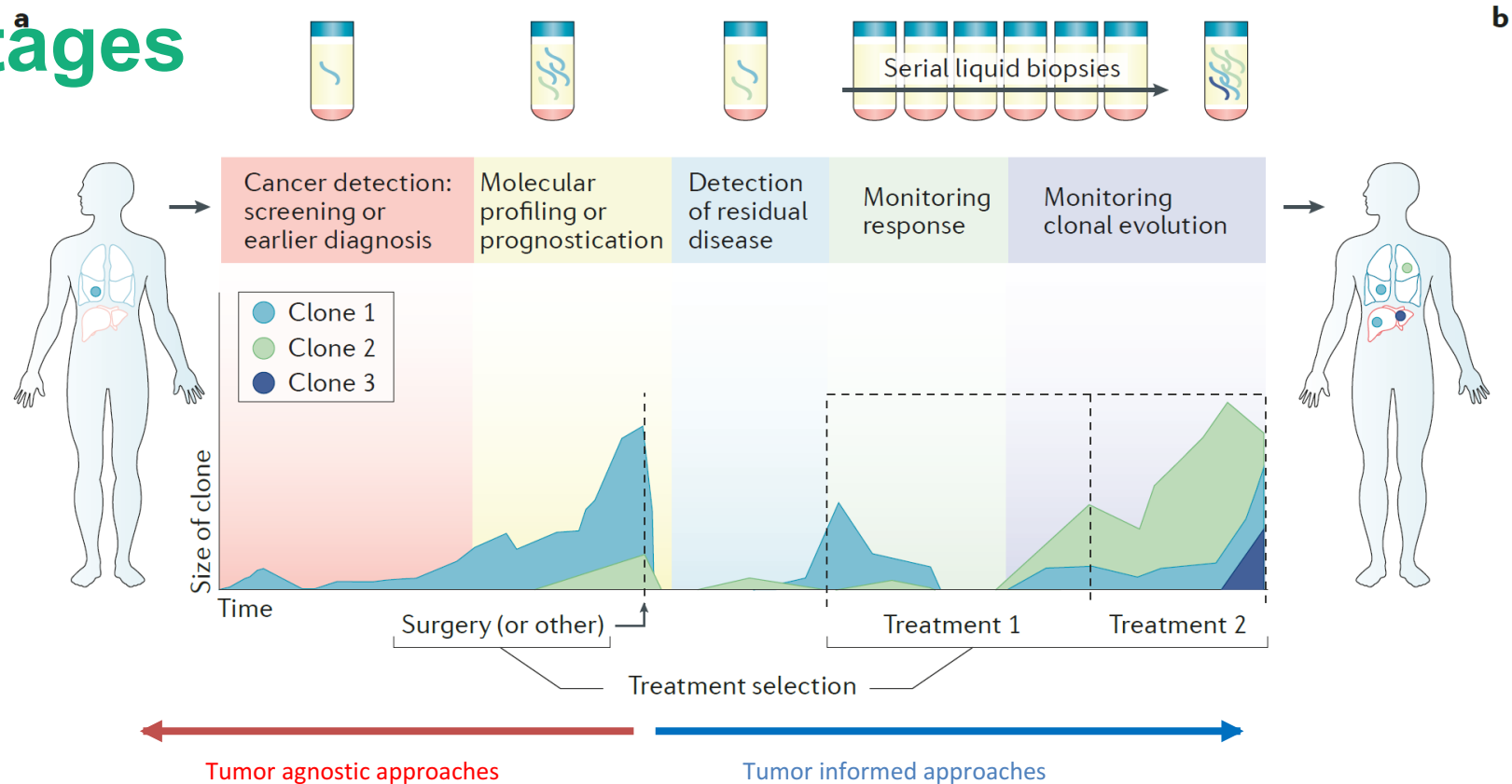
DNA fragments (<146bp)



Urine cfDNA
from renal clearance

cfDNA half life: <2 hours → real time monitoring of tumor burden

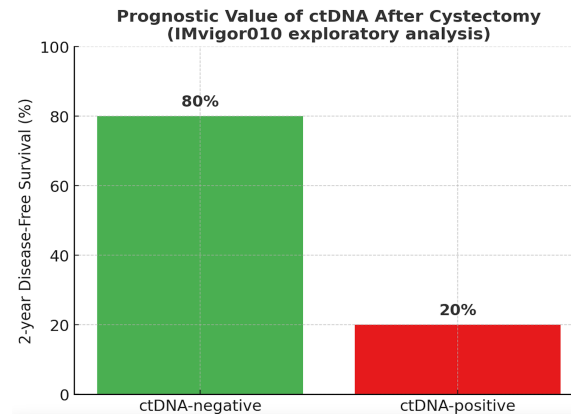
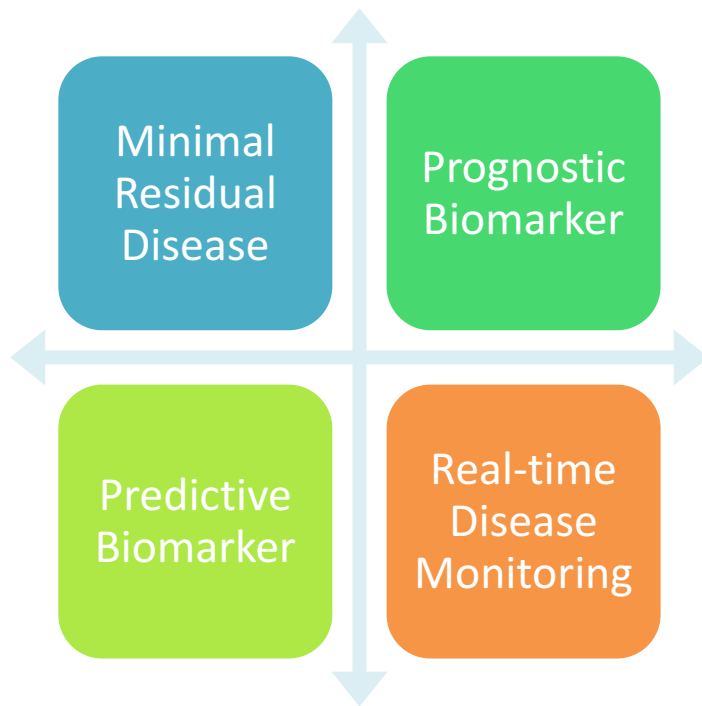
ctDNA has potential to be used thru disease stages^a



Why is ctDNA important as a Biomarker in Urothelial Carcinoma?

IMvigor 010

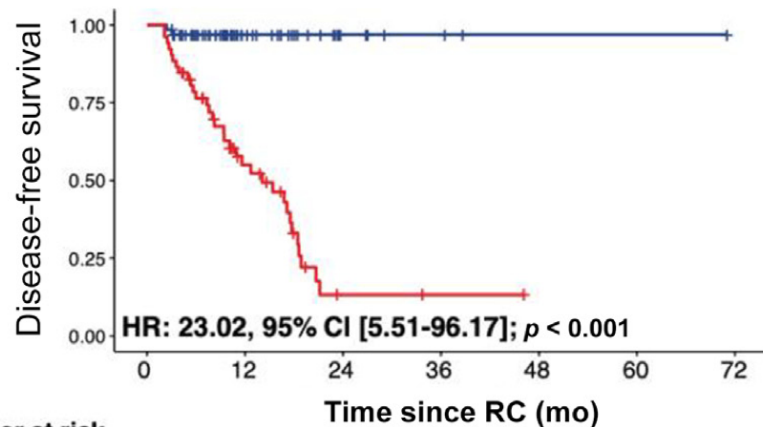
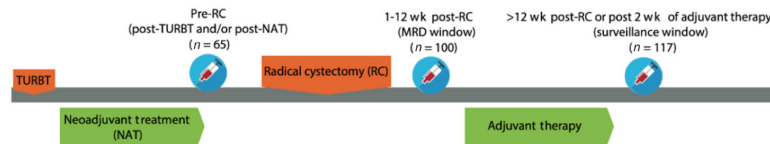
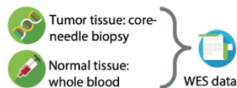
**IMVigor 011
MODERN
TOMBOLA**



ctDNA Assessment → Risk Stratification

- ✓ Strong association between ctDNA detectability and poor DFS
- ✓ ctDNA status is better predictor of DFS than pCR

167 patients with bladder cancer



Number at risk

ctDNA undetectable	65	24	6	3	1	1	0
ctDNA detectable	52	20	2	1	0	0	0

Status	ctDNA undetectable	ctDNA detectable
Events %	3.1 (2/65)	63.5 (33/52)
24-mo DFS %	96.9 (88.2-99.2)	13.2 (3.8-28.6)

Association of Tumor-informed Circulating Tumor DNA Detectability Before and After Radical Cystectomy with Disease-free Survival in Patients with Bladder Cancer

John P. Sfakianos^{a,†}, Arnab Basu^{b,†}, George Laliotis^c, Shivaram Kumarasamy^a, Jordan M. Rich^a, Ajitha Kommalapati^b, Michael Glover^d, Tamara Mahmood^c, Neeraja Tillu^a, Christopher J. Hoimes^e, Grayce Selig^a, Revathi Kollipara^f, Tyler F. Stewart^g, Samuel Rivero-Hinojosa^c, Punashi Dutta^c, Mark Calhoun^c, Shruti Sharma^c, Meenakshi Malhotra^c, Adam C. ElNaggar^c, Minetta C. Liu^c, James E. Ferguson 3rd^b, Marcio Diniz^h, Reza Mehrhazin^h, Peter Wiklund^a, Alan Tan^f, Sumit Shah^d, Matthew D. Galsky^{b,†}

Limitations of ctDNA

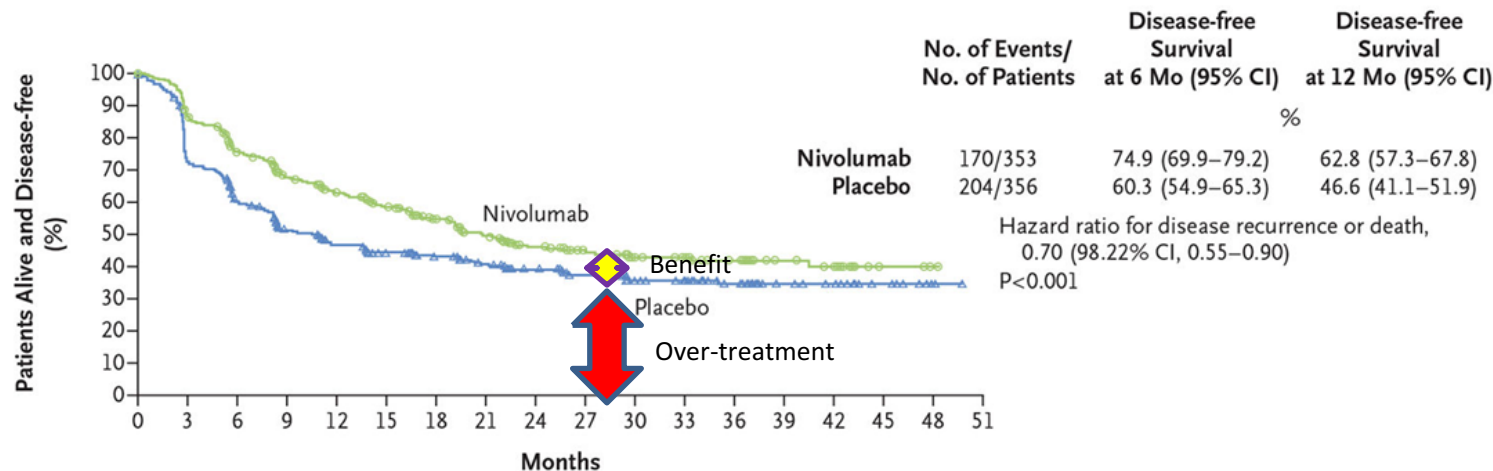
1. **Sensitivity** depends on tumor fraction (low shedding in some NMIBC, small-volume, or papillary disease)- MDR??
2. **Standardization** of assays is lacking (different panels, thresholds, platforms)-
Tumor informed ctDNA assays vs non-tumor informed ctDNA
3. Not yet fully integrated into guidelines — but rapidly moving toward clinical implementation. IMvigor 011?



ctDNA to guide Adjuvant Therapy

Adjuvant IO For high-Risk TCC- Pathological Definition/Imaging FII

Intention-to-Treat Population



No. at Risk															
Nivolumab	353	296	244	212	178	154	126	106	85	68	57	51	36	23	20
Placebo	356	248	198	157	134	121	105	94	80	65	54	50	37	22	19

CM-274
(nivolumab)

- DFS
- OS trending

Ambassador
(Pembrolizumab)

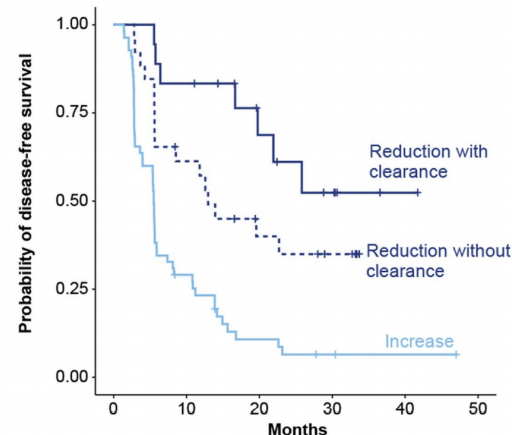
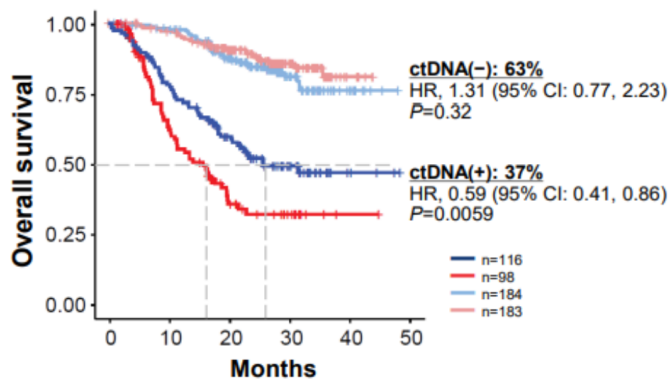
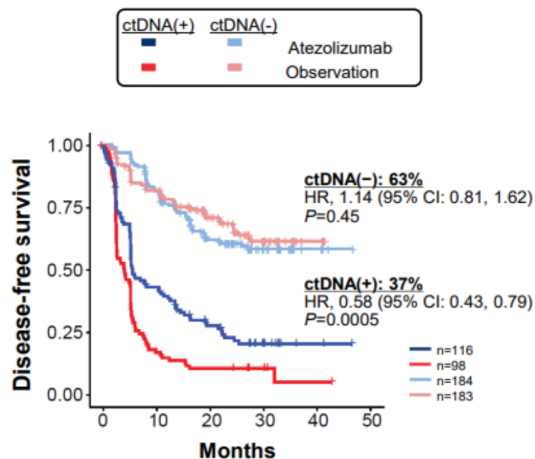
- DFS

Invigor 010
(Atezolizumab)

- NO DFS/OS

ctDNA prognostic/predictive of OS w/ adjuvant atezolizumab- Imvigor 010

DFS based on ctDNA clearance (C1D1+, atezolizumab arm)

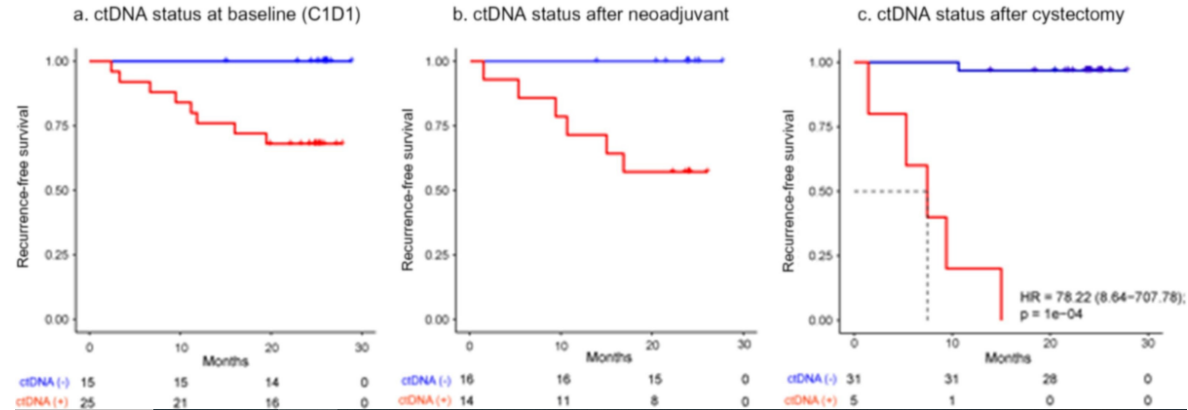
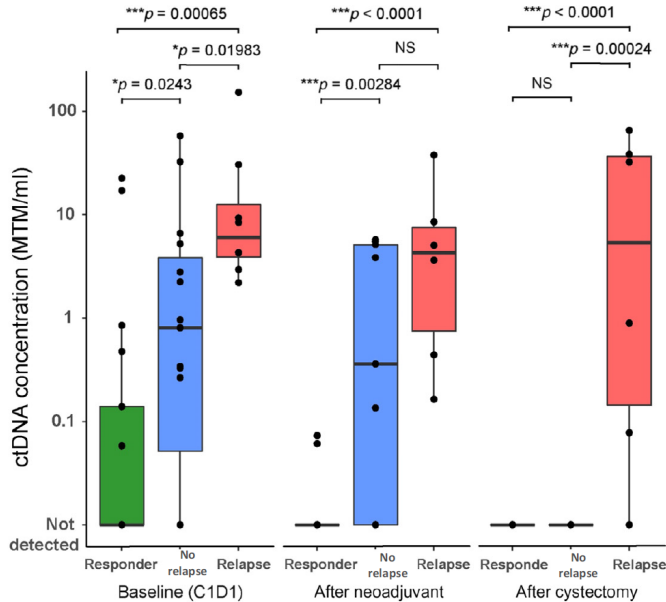


No. at Risk

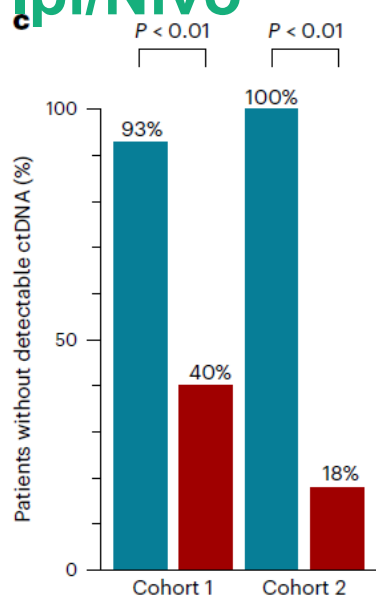
Reduction with clearance	18	15	9	5	1	0
Reduction without clearance	26	15	8	5	0	0

	ctDNA(+) patients	
	Atezolizumab	Observation
Median DFS (95% CI), mo	5.9 (5.6, 11.2)	4.4 (2.9, 5.6)
Median OS (95% CI), mo	25.8 (20.5, NR)	15.8 (10.5, 19.7)

ctDNA prognostic in ABACUS Trial → Neoadjuvant Atezolizumab



ctDNA Dynamics in NABUCCO Trial → Neoadjuvant Ipi/Nivo



d

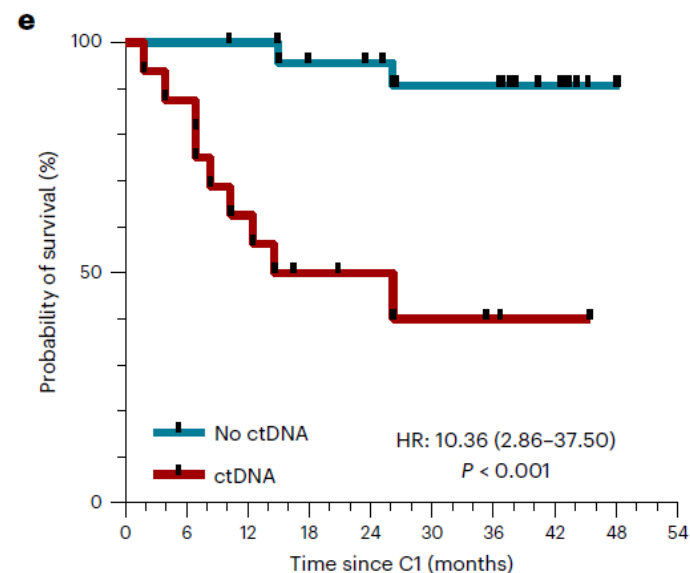
	No detectable ctDNA, <i>n</i>	Detectable ctDNA, <i>n</i>	Total, <i>n</i>
Cohort 1, <i>n</i>			
Responders	13	1	14
Non-resp.	4	6	10
Total	17	7	24
Cohort 2, <i>n</i>			
Responders	5	0	5
Non-resp.	2	9	11
Total	7	9	16

Legend Fig. 2a-c

■ ypT0/Tis/Ta/T1N0

■ ≥ypT2 or N⁺

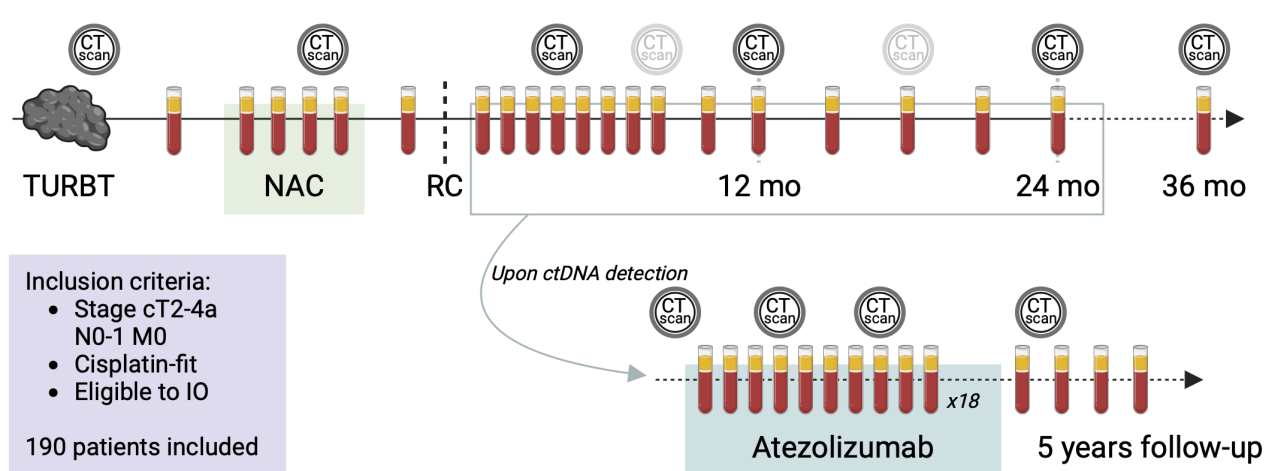
■ Non-evaluable



at risk

No ctDNA	25	25	24	21	20	16	16	8	1	0
ctDNA	16	14	10	6	5	4	3	3	0	0

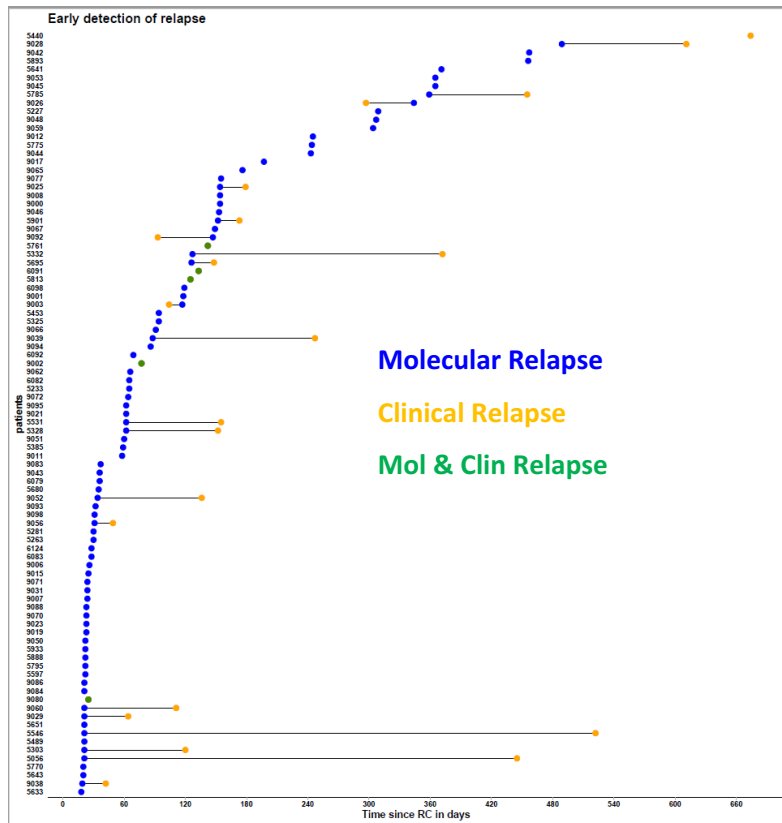
TOMBOLA Trial → ctDNA Guided intervention Trial in MIBC



Primary objective:

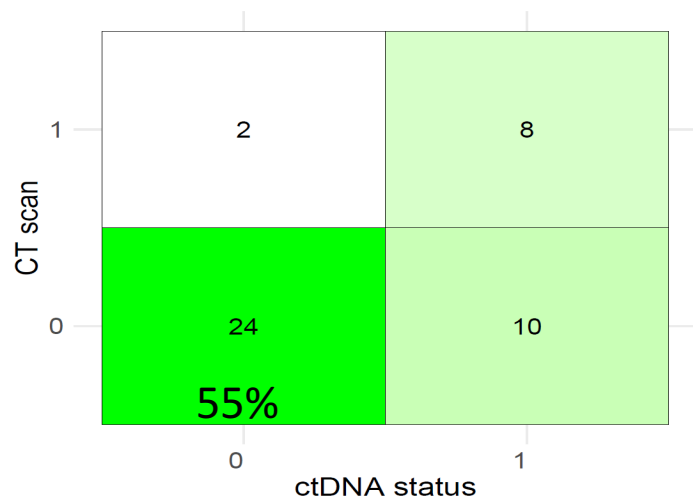
- Complete response (CR) after treatment with investigational agent initiated by ctDNA positive status after radical cystectomy.

CR defined as NED - negative ctDNA and no visible metastasis on CT



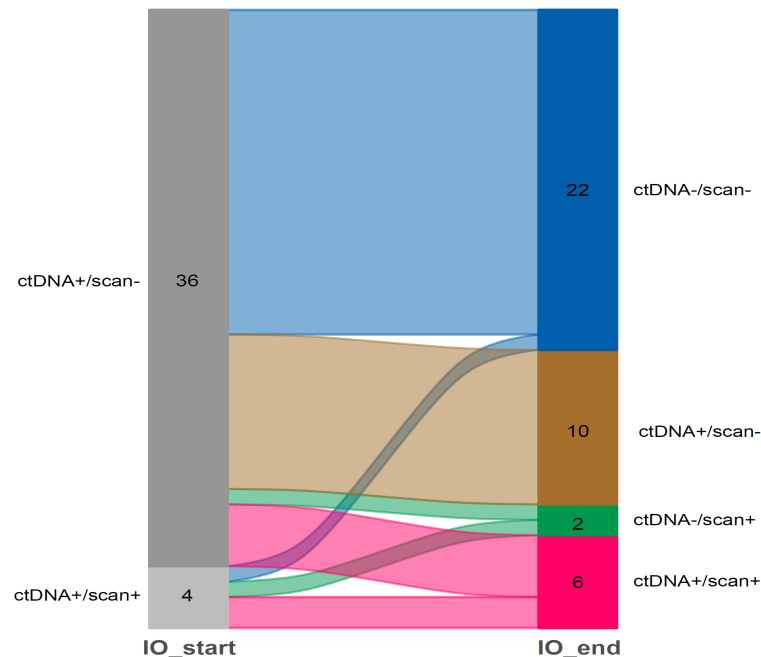
- ✓ 93 of 166 patients (56%) were ctDNA+ post-RC
- ✓ 75% were detected < 4 months post RC
- ✓ Of the ctDNA- patients, only 2 (3%) developed metastases on CT-scan during follow-up

NED (No evidence of disease) (CT and ctDNA-) following immunotherapy

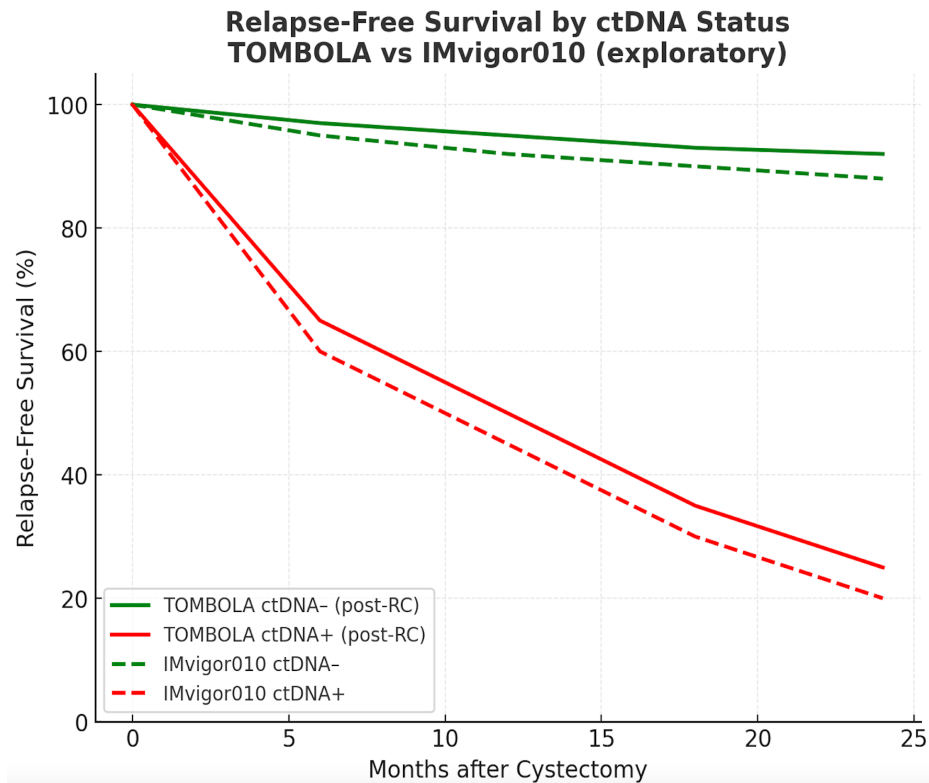


Count
(p-value = 0.0042) 5 10 15 20

ctDNA and imaging during IO



Serial measurements of ctDNA following NAC and RC is a highly specific method to identify patients that might benefit from early immunotherapy at a time of minimal metastatic disease

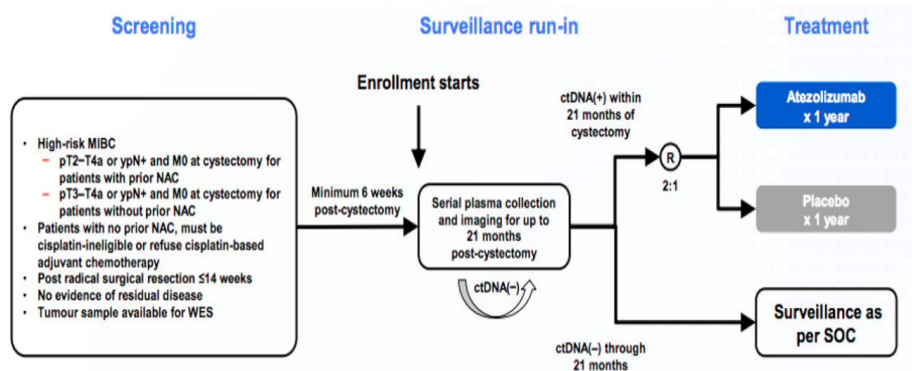


- Prognostic power
- Lead Time to intervention
- Potential to guide adjuvant therapy

ctDNA-guided Adjuvant IO Trials

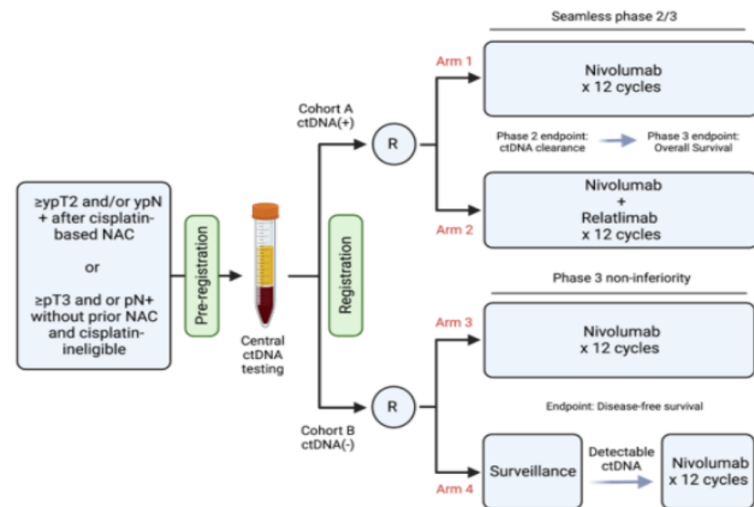
IMvigor011- ctDNA positive pts randomized to atezo vs placebo

MODERN- pts randomized to immediate nivo vs when they become ctDNA positive



Stratification factors

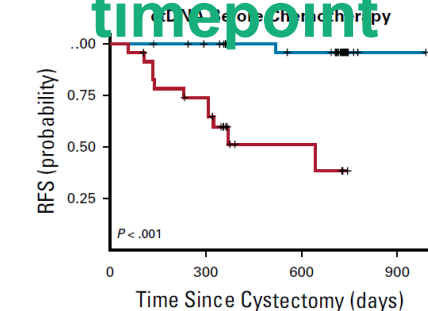
- Nodal status (positive vs negative)
- Tumour stage after cystectomy (spT2 vs pT3/pT4)
- PD-L1 IHC status (IHC score of IC0/1 vs IC2/3)
- Time from cystectomy to first ctDNA(+) sample (≤20 weeks vs >20 weeks)



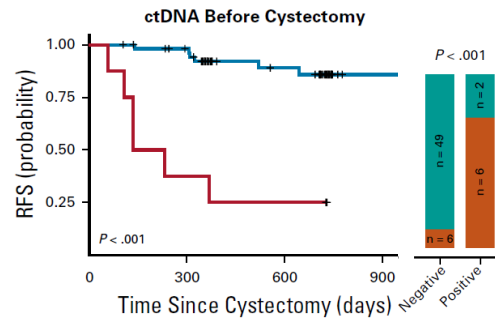
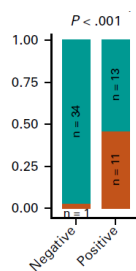


ctDNA to guide Bladder Preservation

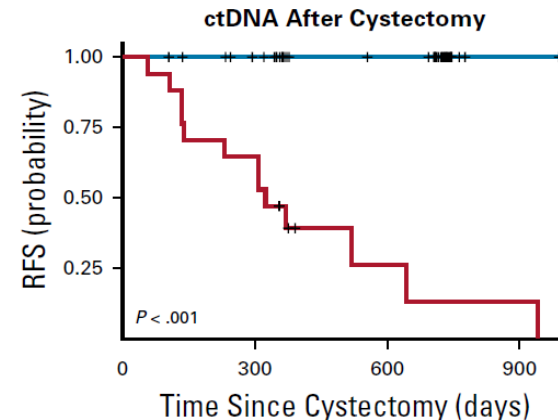
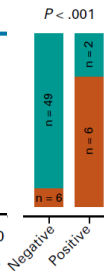
ctDNA is Prognostic of Recurrence in MIBC @ any timepoint



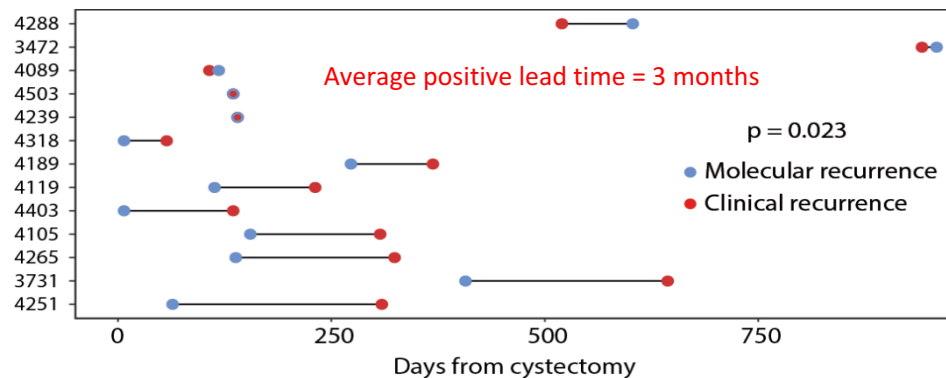
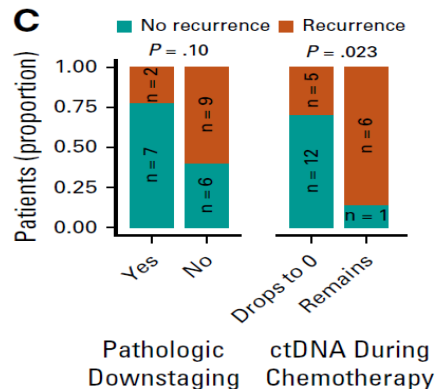
No. at risk	35	32	23	1
Negative	35	32	23	1
Positive	24	16	4	0



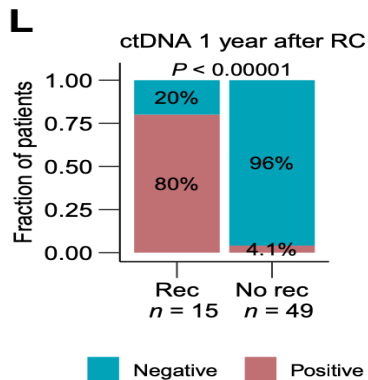
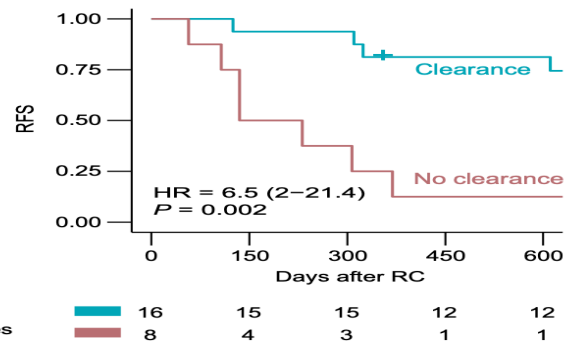
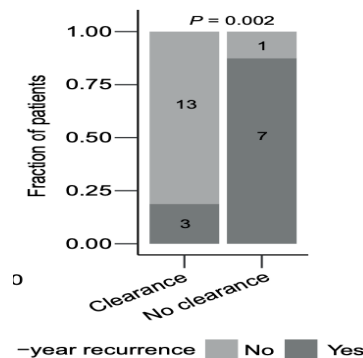
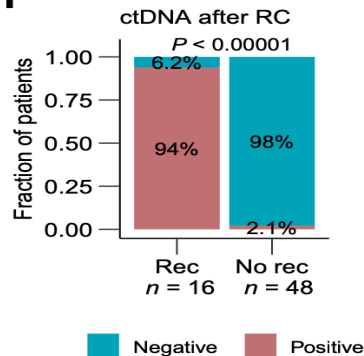
No. at risk	55	49	28	1
Negative	55	49	28	1
Positive	8	3	2	0



No. at risk	47	42	29	1
Negative	47	42	29	1
Positive	17	11	2	1



ctDNA Dynamics during NAC prognostic in MRD



- ✓ N=68; FU 5yr- to predict treatment response w/ ctDNA and early detection of metastasis
- ✓ Includes non-High-risk cohort
- ✓ ctDNA after RC identifies Metastatic relapse (sensitivity 94%)
- ✓ ctDNA Dynamic during NAC are associated w/ pts outcomes
- ✓ **NPV ctDNA predict MRD 89% preNAC/80% preRC**

ctDNA and Outcome after NAC

RESEARCH ARTICLE | OCTOBER 02 2023

Circulating tumor DNA analysis in advanced urothelial carcinoma: insights from biological analysis and extended clinical follow-up

Sia V. Lindskrog¹; Karin Birkenkamp-Demtrodt¹; Iver Nordentoft¹; George Laliotis¹; Philippe Lamy¹; Emil Christensen¹; Derrick Renner¹; Tine G. Andreassen¹; Naja Lange¹; Shruti Sharma¹; Adam C. ElNaggar¹; Minetta C. Liu¹; Himanshu Sethi¹; Alexey Aleshin¹; Mads Agerbæk¹; Jørgen B. Jensen¹; Lars Dyrskjot¹

CLINICAL CANCER RESEARCH | TRANSLATIONAL CANCER MECHANISMS AND THERAPY

Cell-Free Urine and Plasma DNA Mutational Analysis Predicts Neoadjuvant Chemotherapy Response and Outcome in Patients with Muscle-Invasive Bladder Cancer

Anders Bjørndal^{1,2}, Sara K. Elbæk², Sia V. Lindskrog^{1,2}, Mads Agerbæk¹, Philippe Lamy¹

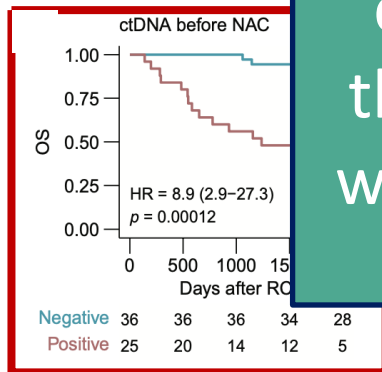


+ Author & Article Information

Clin Cancer Res (2023)

Correlations and associations in a cohort are not enough to guide therapy in an individual patient – we need near 100% NPV to justify bladder preservation.

(Custom Panel)



ctDNA negative : 89% \leq ypTa/CISN0

Negative	55	54	52	47	34
Positive	10	6	2	2	2

ctDNA negative : 80% \leq ypT1N0

Clearance	23	21	21	21	20
Remains	34	27	27	25	21

All ctDNA positive patients had residual disease
All pCR were ctDNA negative

(Signatera)

RETAIN Trial

Major Inclusion Criteria:

- cT2-T3 N0M0
- ECOG 0-1
- Urothelial Predominant Histology

MFS is defined as the absence of a recurrence of urothelial carcinoma that is >cN1 (more than one clinically suspicious pelvic lymph node) or surgically unresectable local recurrence (e.g., >cT4a) or M1 disease.



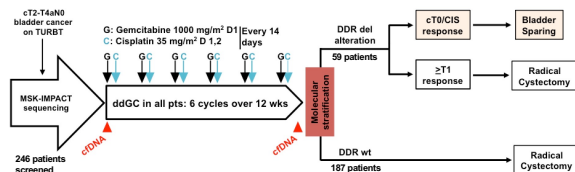
Sequencing (Caris)

Mutation positive

- defined as any alterations in:
- ATM
 - RB1
 - FANCC
 - ERCC2

Primary Endpoint: Metastasis-free survival (MFS) at 2 years.
Non-inferiority design with a 14% margin between risk-adapted design (MFS=78%) and standard-of-care (MFS=64%).
Sample size=70 with an 82% power, Type I error=0.045

A031701: A phase II study of dose-dense Gemcitabine plus Cisplatin in patients with muscle-invasive bladder cancer with bladder preservation for those patients whose tumors harbor deleterious DNA damage response (DDR) gene alterations



Modified DDR gene

panel

ERCC2
ERCC5
BRCA1
BRCA2
RAD51C
ATR
RECQL4
ATM
FANCC

Deleterious alterations in one or more of these genes will allow patients to be potentially eligible for the bladder-sparing arm of the study



HOOSIER

HCRN GU16-257

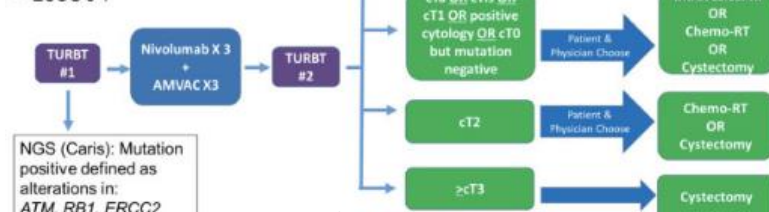


• Determine association between DDR panel and "benefit" in cCR patients

DDR Panel: ERCC2, FANCC, ATM, RB1 and TMB

Major Inclusion Criteria:

- cT2-T3 N0
- Predominant Urothelial Carcinoma of Bladder
- ECOG 0-1



RETAIN 2

Metastasis-free survival (MFS) is defined as the absence of a recurrence of urothelial carcinoma that is >cN1 (more than one clinically suspicious pelvic lymph node) or surgically unresectable local recurrence (e.g., >cT4a) or M1 disease).

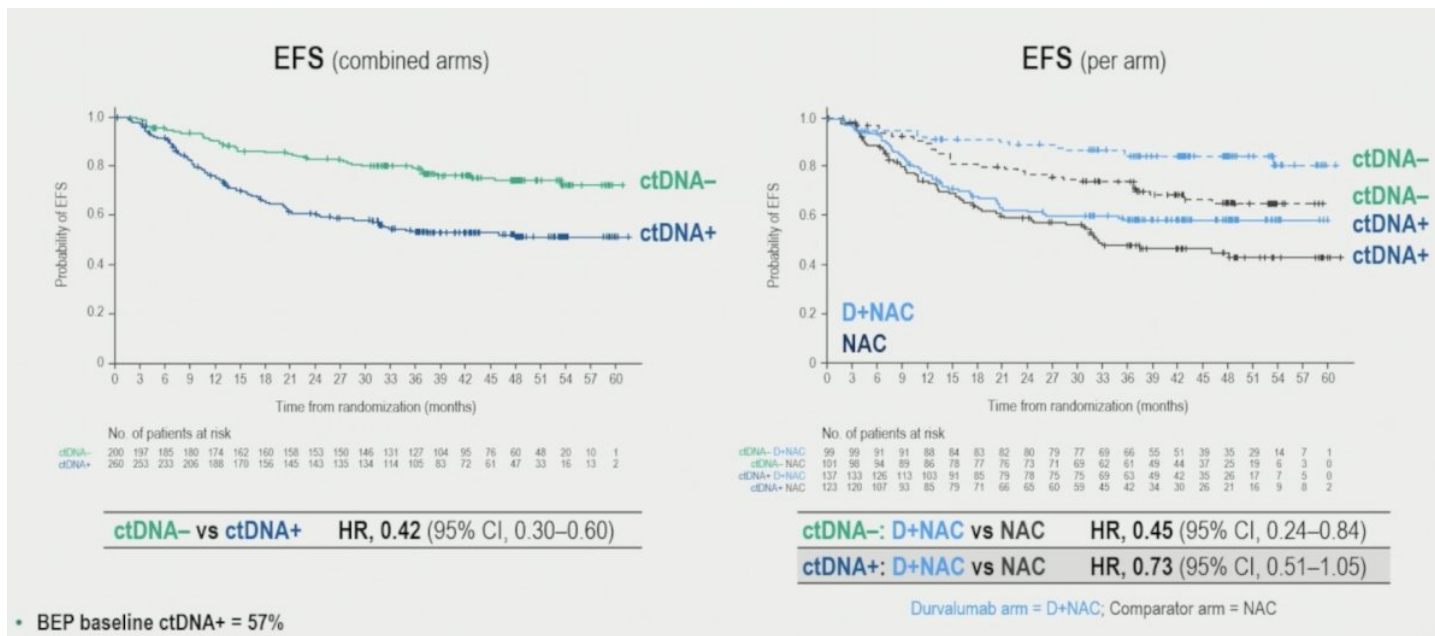
Primary endpt:
2-yr Metastasis-free survival
Follow-up: 5 years

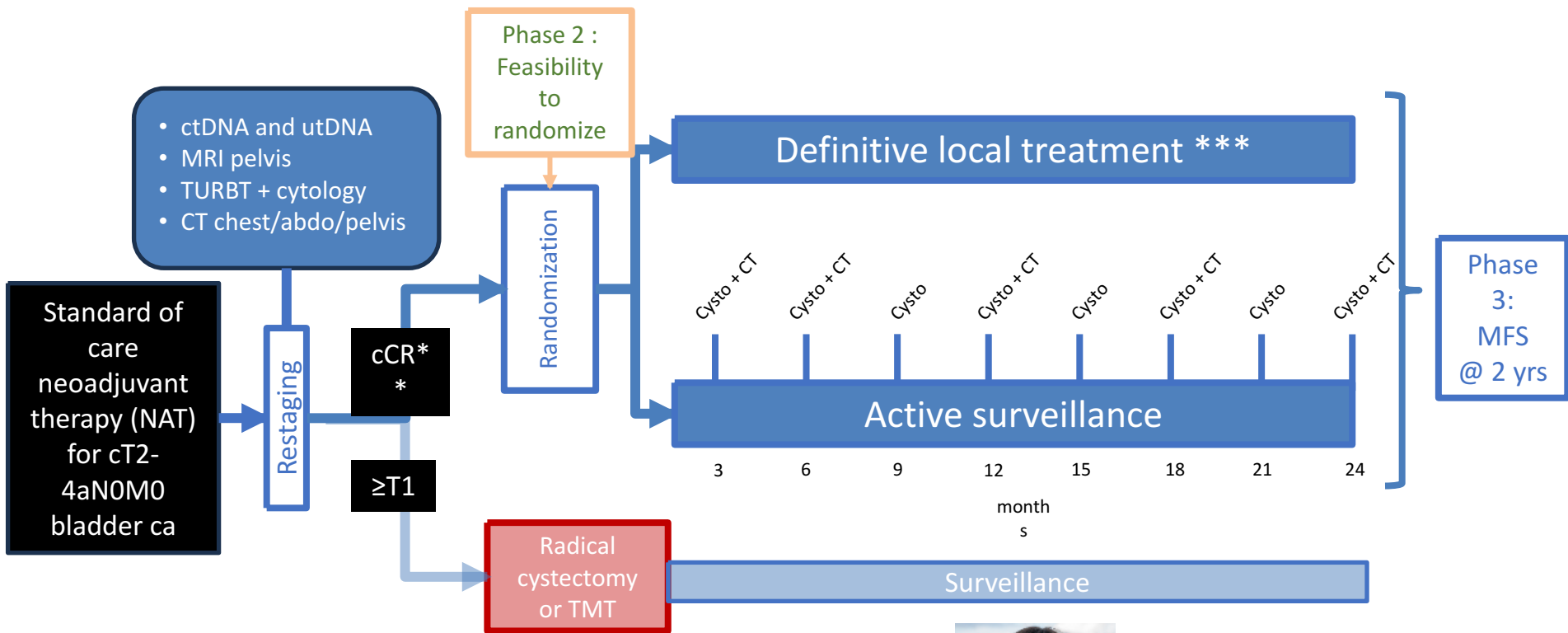
Abbreviation: TURBT= transurethral resection of a bladder tumor, AMVAC=accelerated MVAC, NGS= next generation sequencing, chemo-RT= chemoradiation

ctDNA to guide Perioperative Treatment

NIAGARA Trial- Durvalumab NAC

- ctDNA preRC correlated with no CR
- EFS Benefit if ctDNA clearance specially with D





**Definition of cCR:

- Negative MRI (VI-RADS 0-1-2)
- Negative repeat TURBT (< cT1, no extensive CIS)
- Negative ctDNA
- Negative utDNA
- No mets (negative conventional CT or PET-CT)



Marie-Pier St-Laurent
UBC

*NAT = any SOC regimen approved at time of enrolment (Cisplatin, EV, IO, etc.)

*** Patient/investigator's choice; Radical cystectomy or TMT

Take Home Message→ ctDNA in MIBC

- ctDNA is the most promising tool we have to tailor adjuvant therapy in MIBC→ risk adapted adjuvant therapy.
- Current evidence shows **strong prognostic value** and signals of **predictive benefit** in guiding immunotherapy.
- But: **not ready for “prime time” yet** — still awaiting results from ongoing randomized ctDNA-guided trials (IMvigor 011 @ESMO 2025)
- Limitations apply → standardization assays; definition negative test; timing; costs

For now→ use in clinical trial settings, while preparing infrastructure for eventual adoption.