

Update of Liquid Biopsy from Advanced to early disease



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**Mount
Sinai**

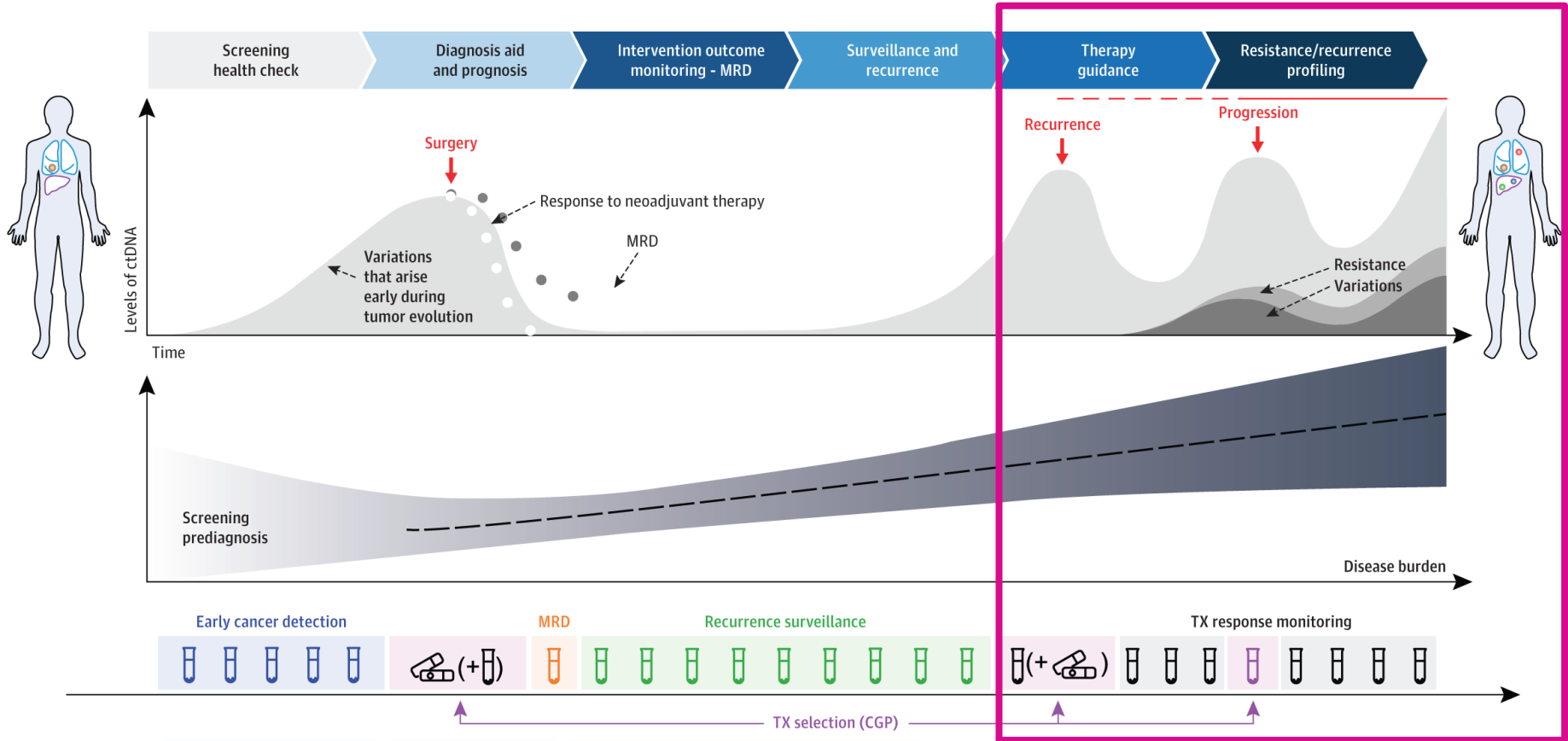
The Tisch Cancer Institute



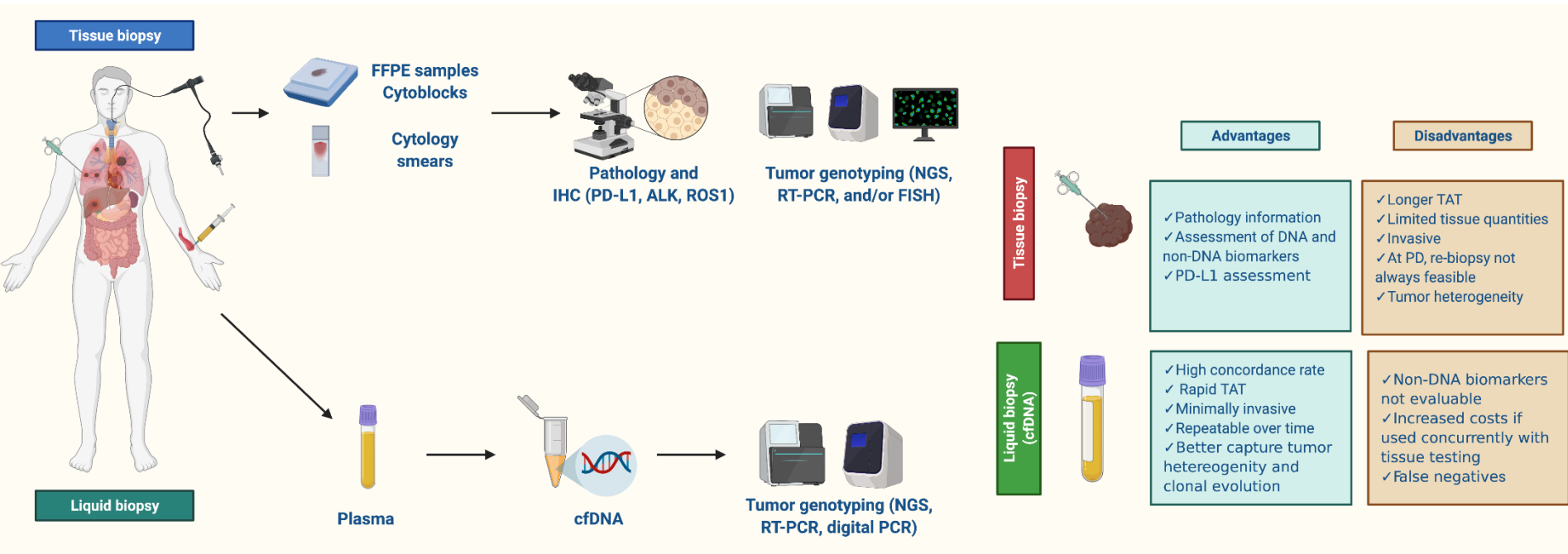
Center for Thoracic Oncology

Disclosures

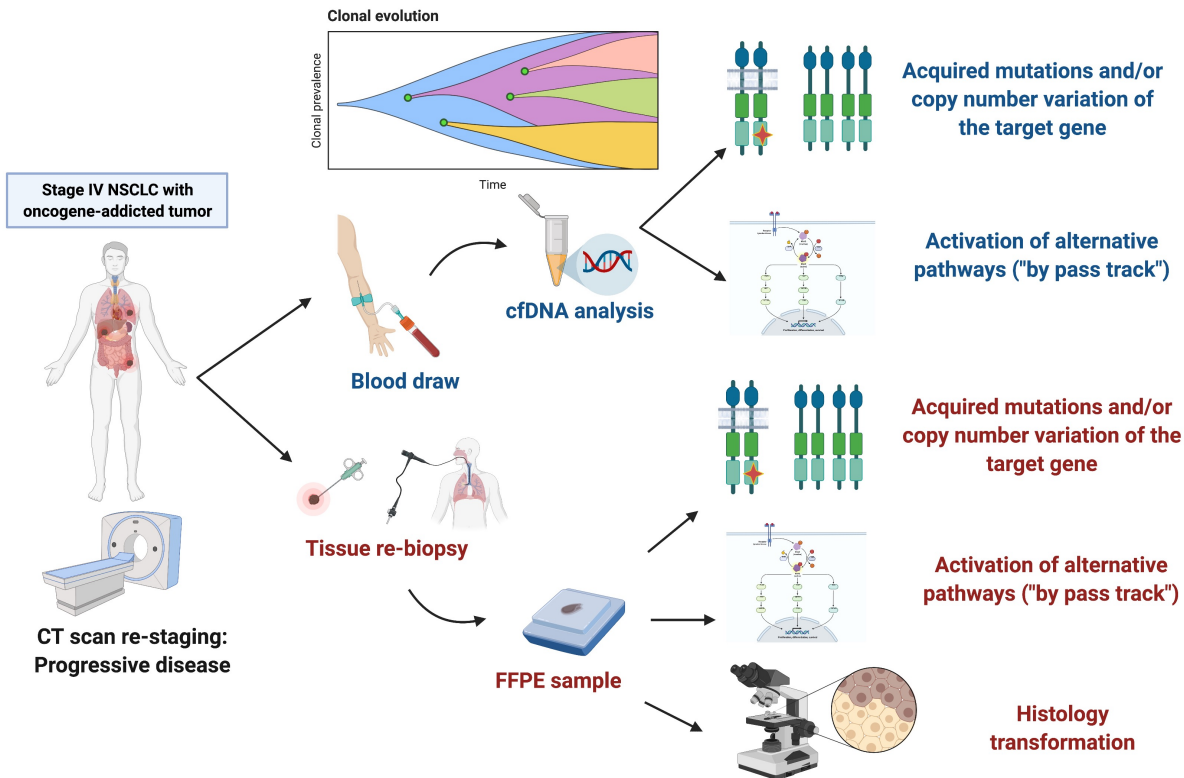
Research grants	Lung Cancer Research Foundation-Pfizer Grant 2019 American Cancer Society NIH SBIR, NCI SeroNet 2020
Personal financial interests	Speaker: MSD, Astra Zeneca, Roche, GuardantHealth
Personal financial interests	Advisory board: Inivata, ArcherDx, EMD Serono, Novartis, BMS, Boston Pharmaceuticals, Esai, BluePrint, CORE2, Pfizer
Non-financial interests	Research Collaboration: GuandantHealth (UMB)
Leadership roles	Chair Educational Committee IALSC - President ISLB (International Society of Liquid Biopsy) - Educational Chair: OLA Oncology Latin American Association Scientific Committee Member at ESO (European School of Oncology).



Tissue vs. Liquid biopsy



Clinical utility of liquid biopsy in oncogene-addicted NSCLC

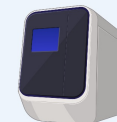


Main liquid biopsy techniques used



NGS-based approaches:

- ✓ High sensitivity
- ✓ Multiplex
- ✓ Gene rearrangements
- ✓ Gene amplifications



PCR-based approaches:

- ✓ Variable sensitivity
- ✓ Single gene testing
- ✓ Only for mutations

Main techniques used for tumor tissue



NGS-based approaches:

- ✓ High sensitivity
- ✓ Multiplex
- ✓ Gene rearrangements
- ✓ Gene amplifications



FISH:

- ✓ Gene rearrangements & amplifications



PCR-based approaches:

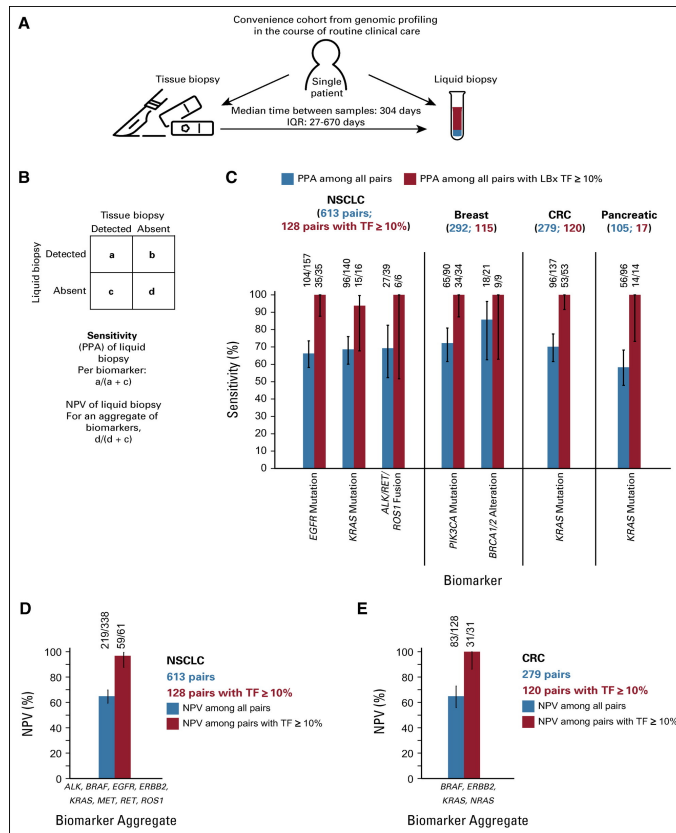
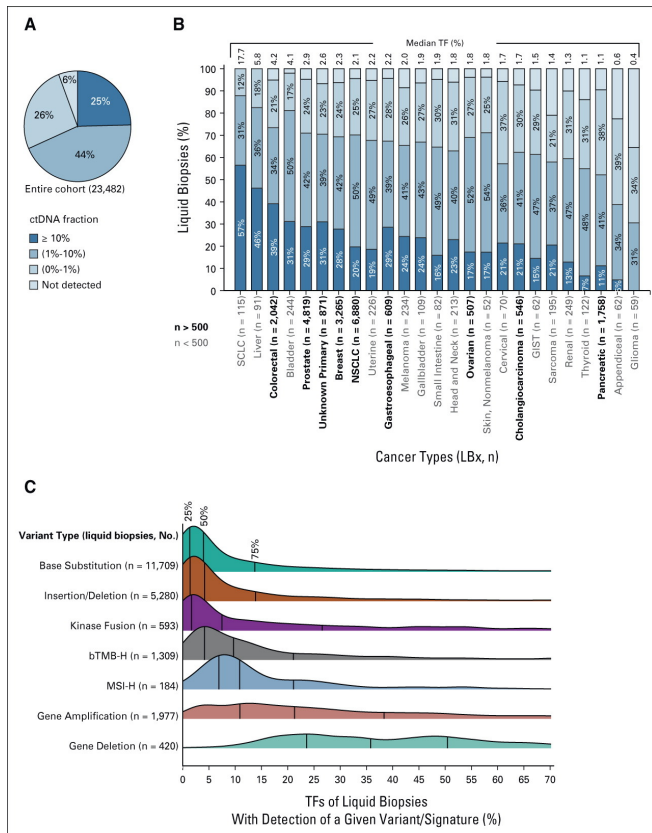
- ✓ Variable sensitivity
- ✓ Single/Multiplex gene testing
- ✓ Only for mutations



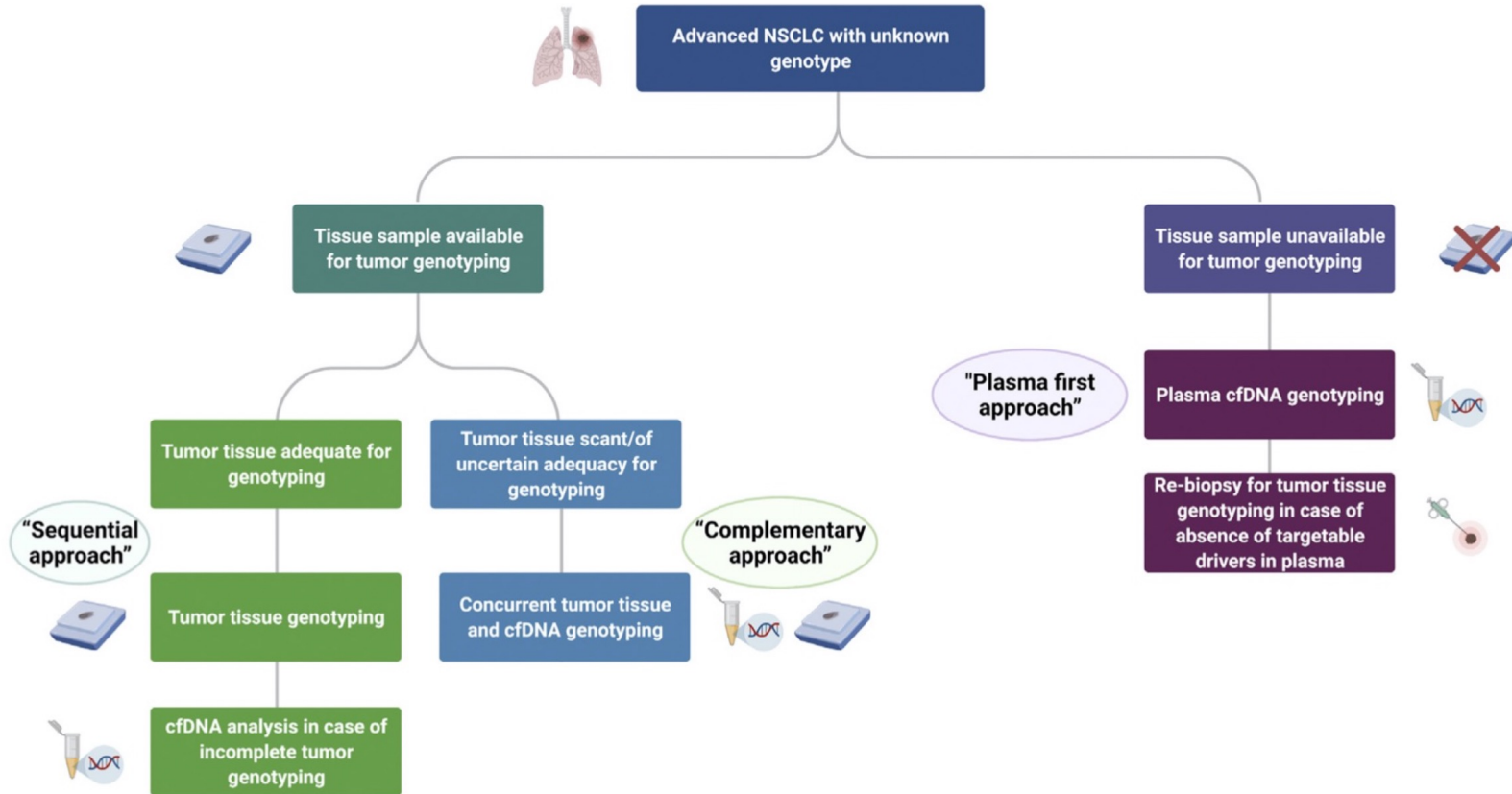
IHC:

- ✓ Protein expression

Tumor Fraction Correlates with Detection of Actionable Variants Across > 23,000 Circulating Tumor DNA Samples

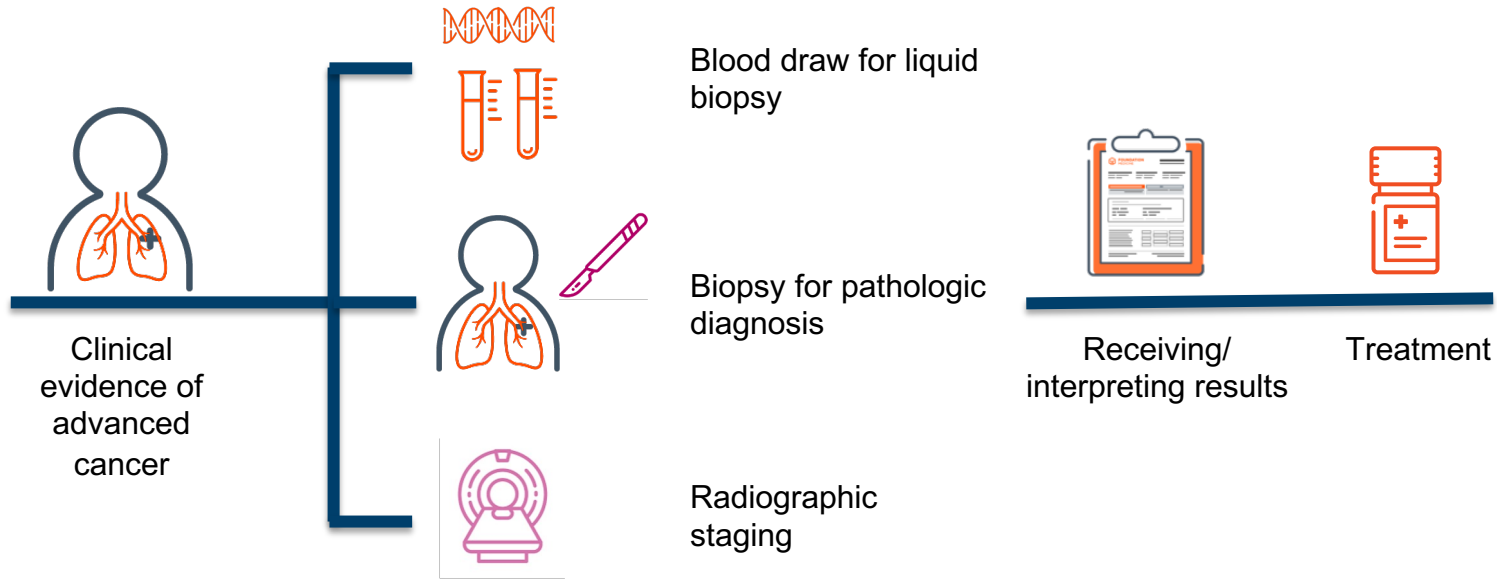


- Elevated ctDNA shed is associated with both high sensitivity and negative predictive value for detection of actionable Genomic Alterations .
- The presence of elevated TF suggests adequate tumor profiling and may reduce the value of subsequent reflex to confirmatory tissue testing in patients with negative LBx results.

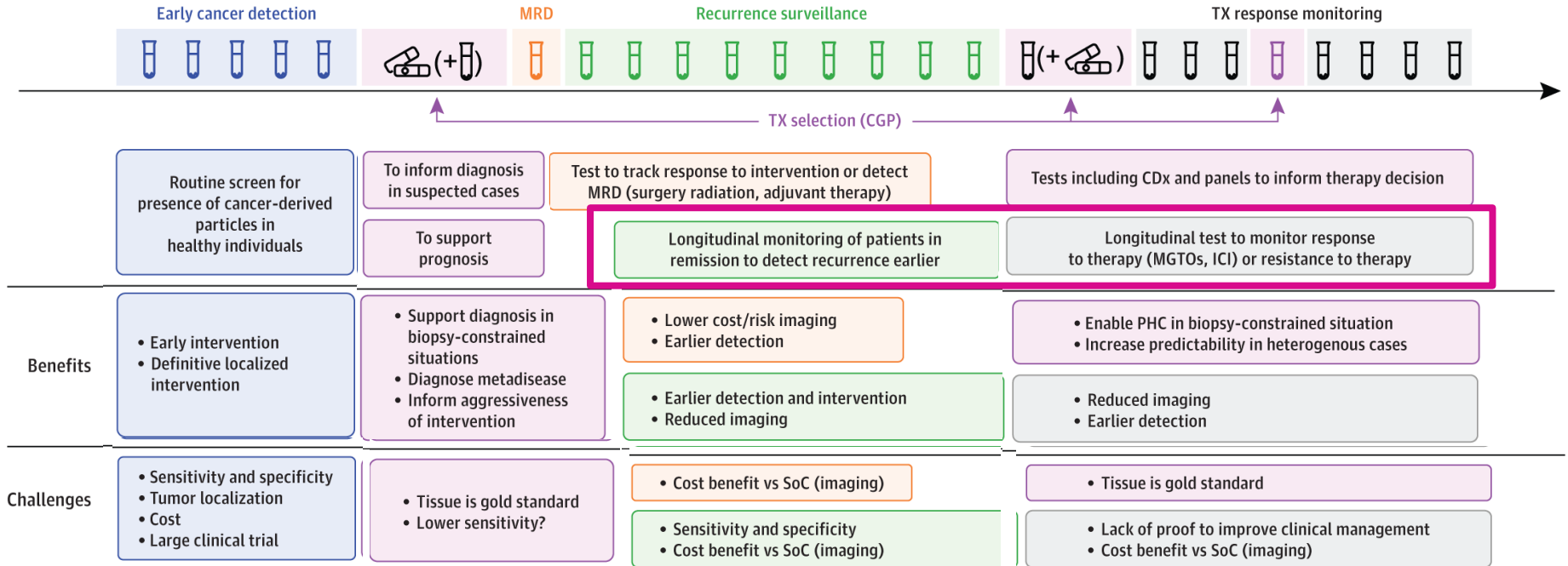


Expedited diagnostic odyssey

Stacking diagnostic steps may be able to shorten the diagnostic odyssey

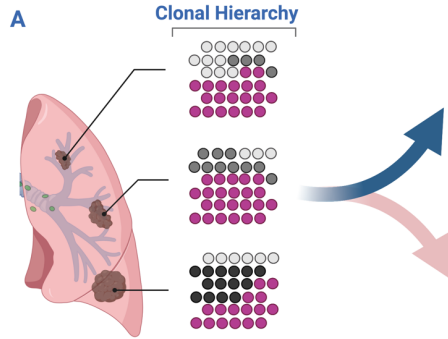


Benefits and challenges of LB in the Cancer Journey



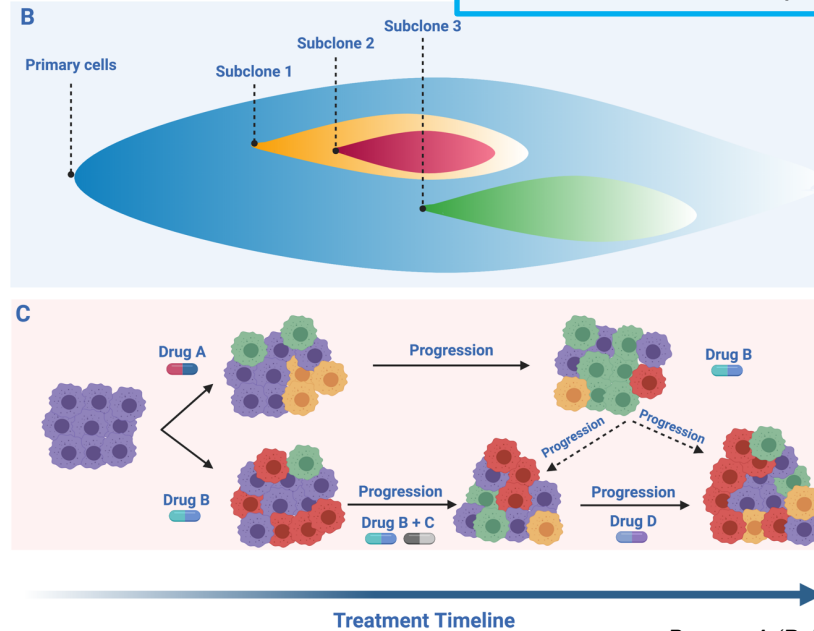
Acquired resistance is a dynamic process

Mechanisms of acquired resistance might be heterogenous and multiple mechanisms can simultaneously occur in the same patient, reflecting the clonal heterogeneity of the tumor



Tracking the clonal evolution of the tumor over time might allow the implementation of tailored therapeutic approaches

The clonal evolution of the tumor under the selective pressure of anticancer therapies



Tailoring treatment with Liquid Biopsy

Osimertinib start with an intracranial CR and extracranial PR

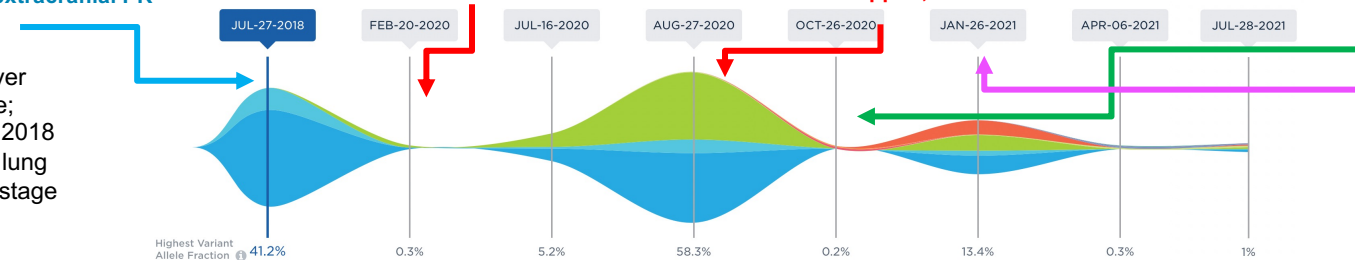
After 18 mos oligo-PD (LN mets) → SBRT and continued osi

Further disease progression → osimertinib stopped; switch to erlotinib

PET/CT in Nov. 2020: CR

Erlotinib discontinuation → platinum-based chemo start

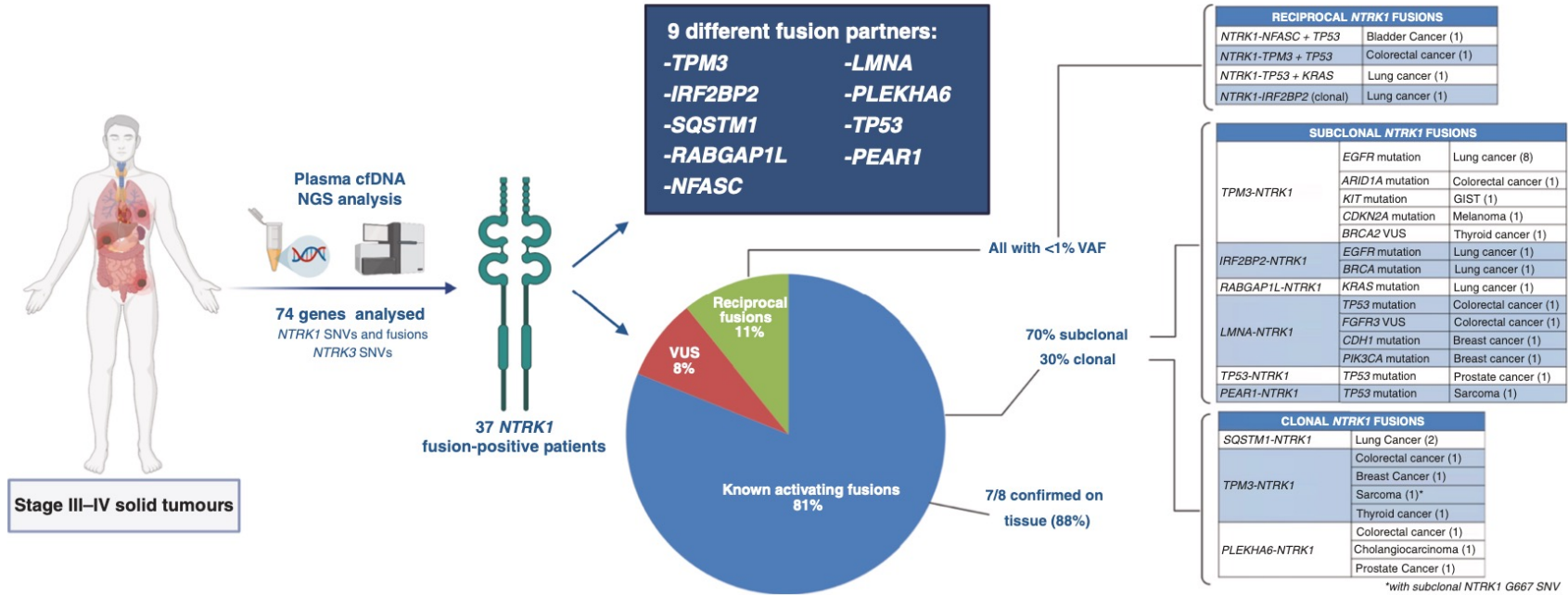
52-year-old never smoker female; diagnosed in July 2018 with cT4 N3 M1c lung adenocarcinoma (stage IVB).



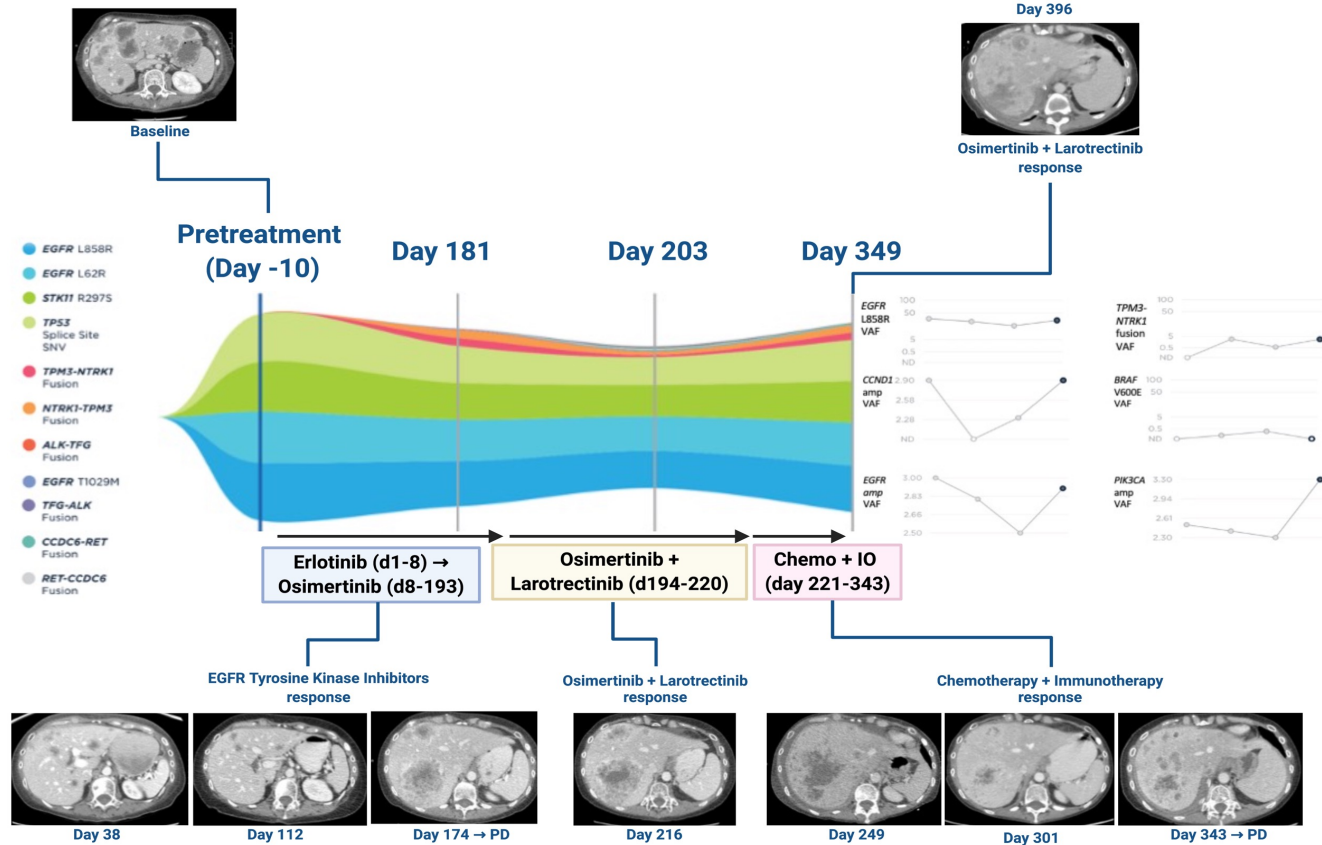
Genetic Alteration	% cfDNA or amplification							
EGFR E746_A750del	41.2%	0.2%	4.7%	58.3%	ND	13.4%	ND	1%
EGFR C797S	ND	0.3%	5.2%	5.6%	ND	10.7%	ND	0.7%
ARID1A Q456Q	ND	ND	ND	0.2%	ND	0.2%	0.3%	0.6%
EGFR T790M	ND	ND	ND	ND	ND	9.6%	ND	0.4%
TP53 C275Y	ND	ND	ND	ND	ND	ND	0.1%	0.2%
ARID1A F1728F	ND	ND	ND	ND	ND	ND	0.3%	0.2%
TP53 S127F	6.5%	ND	0.4%	7.6%	ND	2.6%	ND	0.2%
BRAF Amplification	2.2%	ND	ND	ND	ND	ND	ND	ND
CDK6 Amplification	2.2%	ND	ND	ND	ND	ND	ND	ND
EGFR Amplification	3.4%	ND	ND	4.2%	ND	ND	ND	ND
NTRK2 L699L	-	-	-	-	0.2%	ND	ND	-
EGFR N338N	ND	ND	ND	ND	0.1%	ND	ND	ND
FGFR1 V795I	ND	ND	ND	ND	ND	ND	0.1%	ND

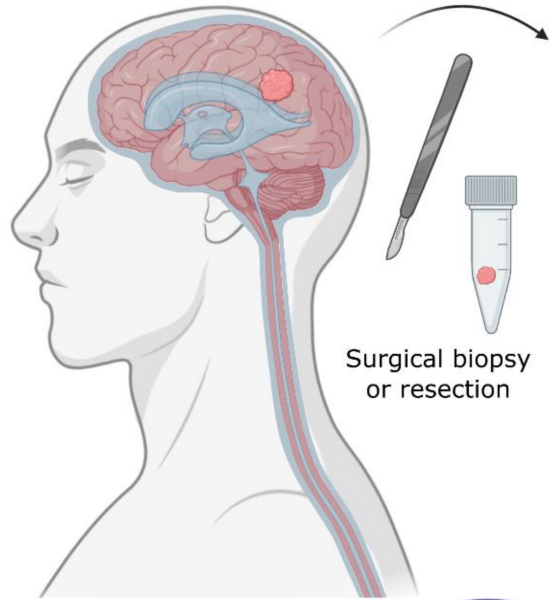


NTRK1 Fusions identified by non-invasive plasma next-generation sequencing (NGS) across 9 cancer types



NTRK fusions as mechanism of resistance





Solid biopsy (tumour specimen)

Advantages

Allow histological diagnosis

Limitations

Very invasive and risky procedure

Sometimes not feasible due to tumour anatomical location

Not representative of tumour heterogeneity

Static snapshot

Surgical biopsy
or resection

Liquid biopsy (CSF ctDNA)

Advantages

Less-invasive and easier to obtain than a tumour biopsy

CSF obtained as SOC for some patients

Concordance with tissue characterisation

Representative of intratumour and interlesion heterogeneity

Longitudinal real-time monitoring

Limitations

No histological characterisation

Lack of standardisation

Contraindications for lumbar puncture

Limited sensitivity

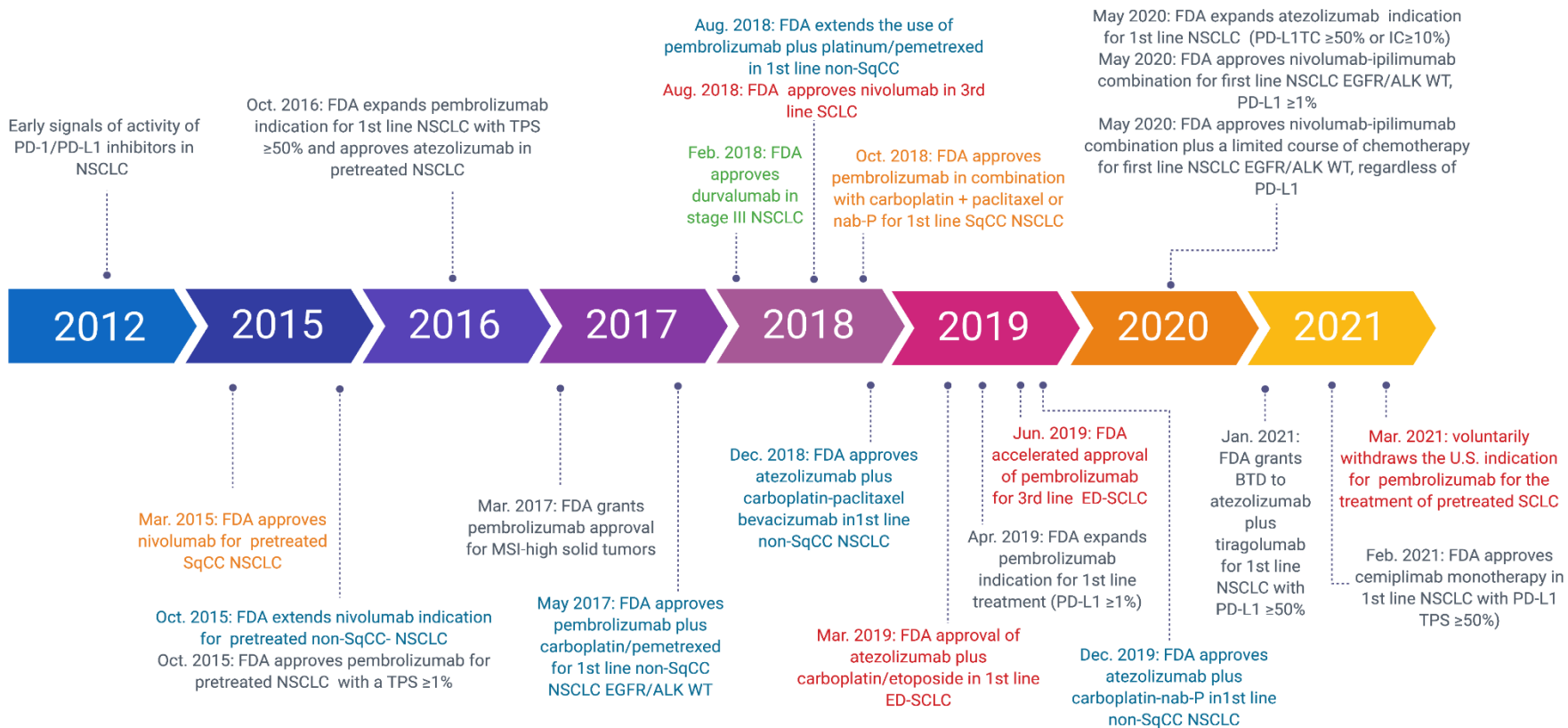
Lumbar puncture



Immunotherapy: The oncologists like a kid in a candy shop...



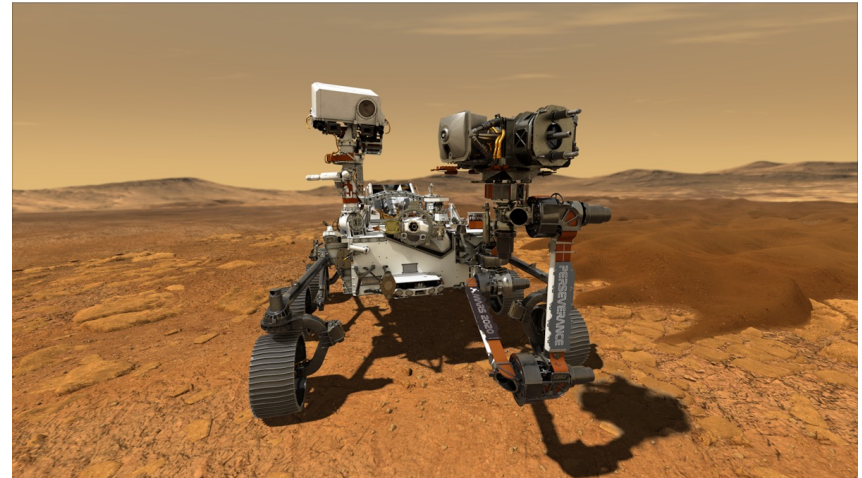
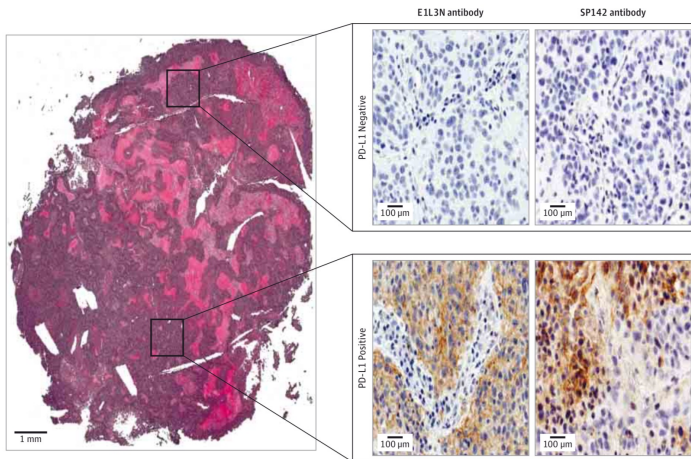
Milestones in Immunotherapy era in Lung Cancer



Adapted from Russo A (Rolfo C) et al. In: Naing A., Hajar J. (eds) Immunotherapy. Adv Exp Med Biol 2020

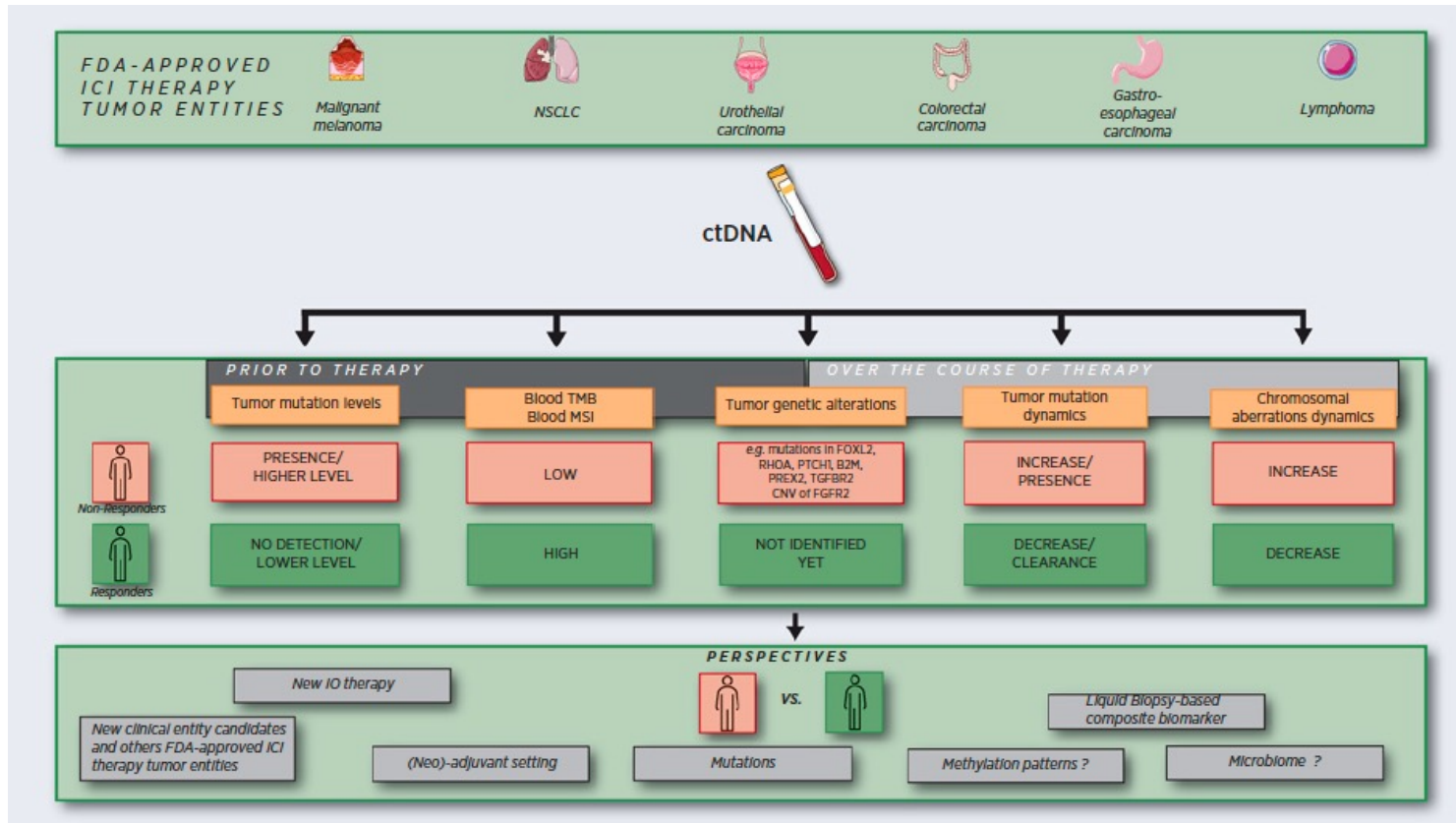
Heterogeneity of PD-L1 Expression

An imperfect but useful biomarker

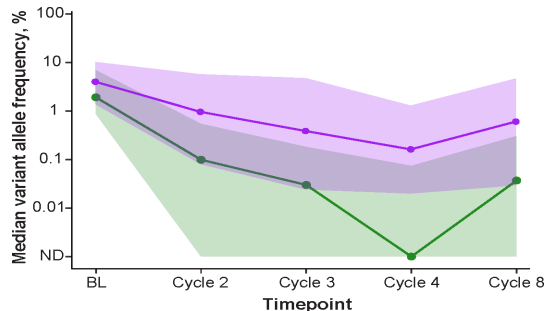
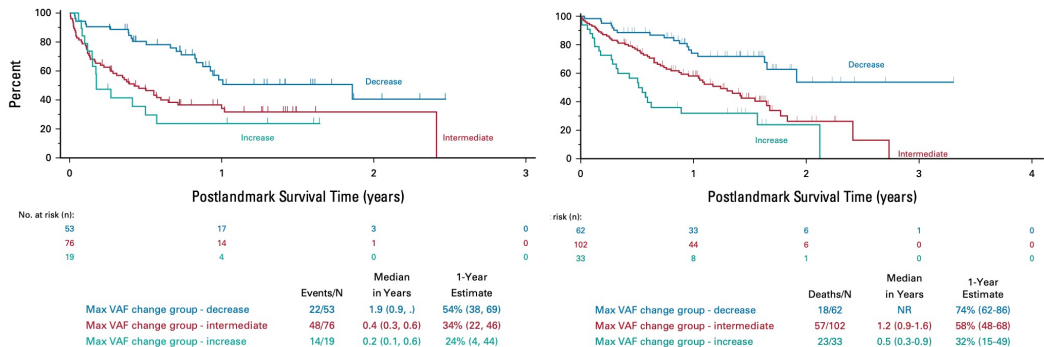
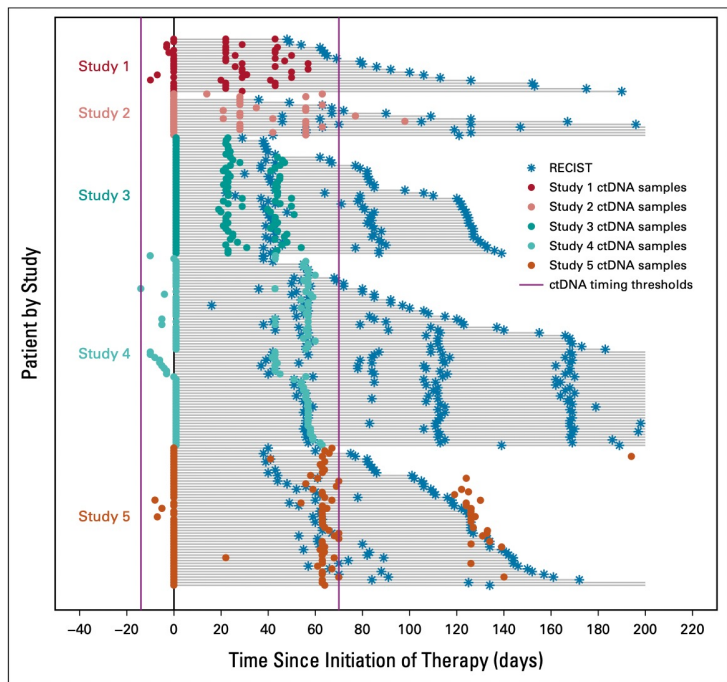


- **Intratumor** heterogeneity
- **Intrapatient** heterogeneity

Use of Liquid Biopsy in Immunotherapy



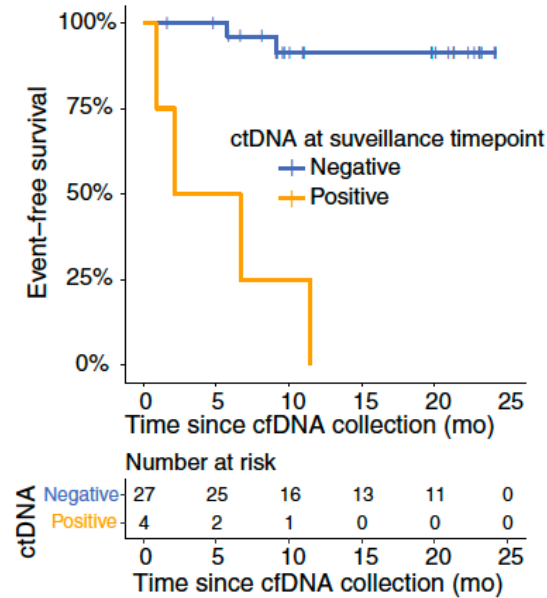
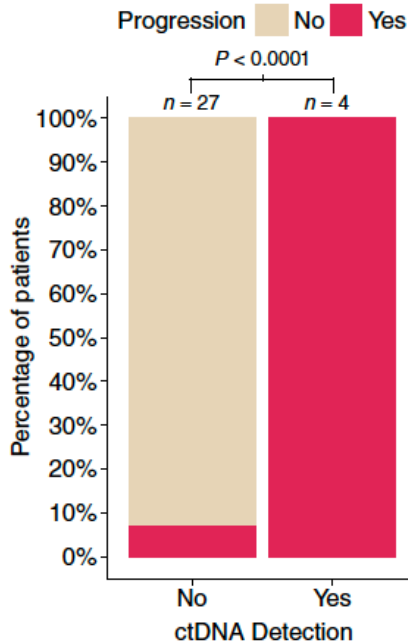
Changes in Circulating Tumor DNA Reflect Clinical Benefit Across Multiple Studies of Patients With Non-Small-Cell Lung Cancer Treated With Immune Checkpoint Inhibitors



ctDNA dynamics reflect tumour burden (SLD) and response to therapy: IMpower150

CtDNA may serve as an important tool in clinical development and an early indicator of treatment benefit

ctDNA Analysis to Assess Risk of Progression to PD-(L)1 Blockade in NSCLC

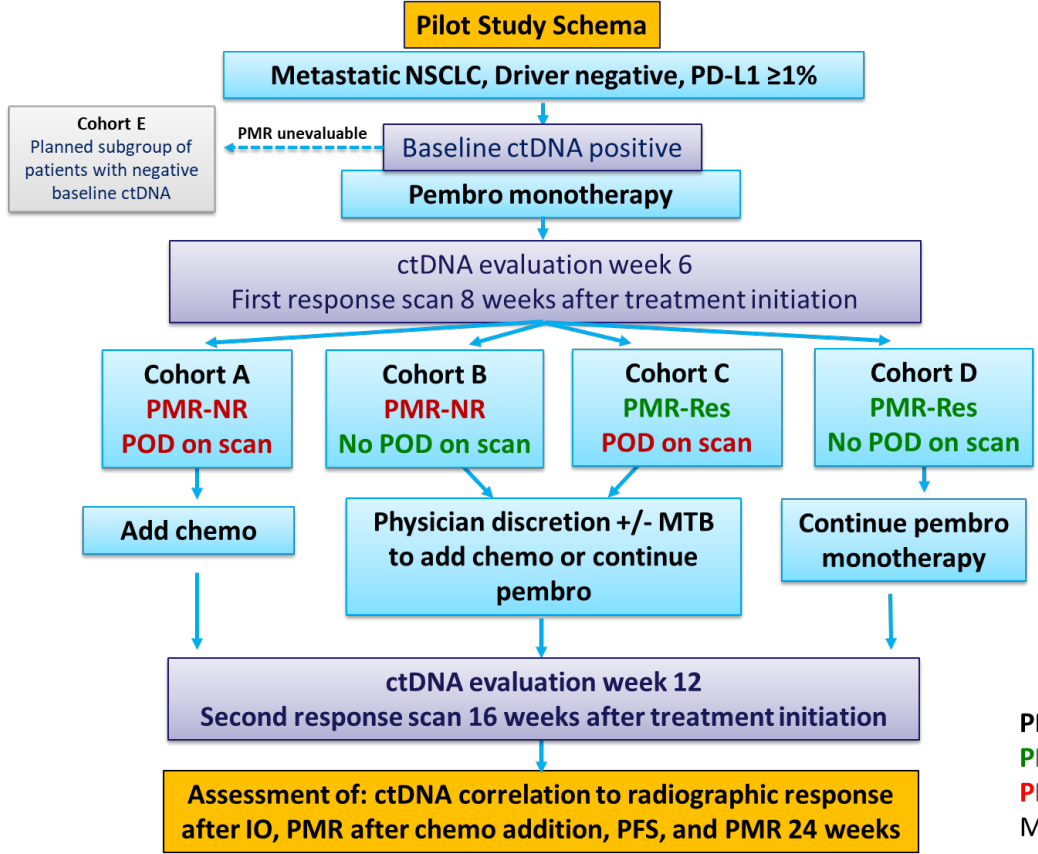


ctDNA analysis identifies patients at risk for eventual progression after long-term response to PD-(L)1 blockade.

Hellmann MD, et al. Clin Cancer Res 2020

CITAN: ctDNA-guided Immunotherapy-based Therapy in Treatment Naïve Advanced NSCLC

PI: Dr. Mack – Dr Rolfo



Translational Medicine Component

Baseline
Tissue, PBMCs

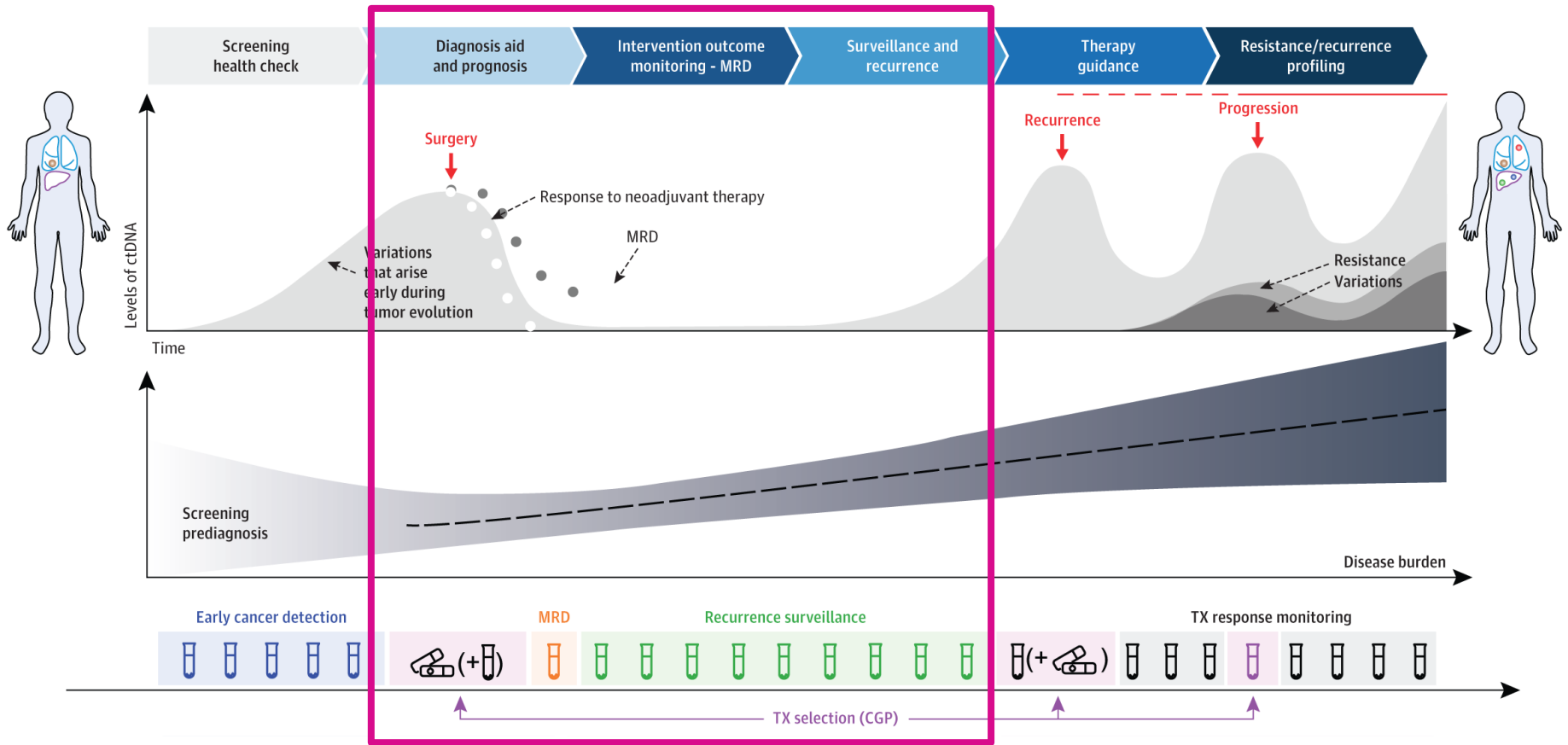
C2D1
ctDNA, PBMCs

C3D1
ctDNA, PBMCs

C4D1
ctDNA, PBMCs

Every 3 Cycles
ctDNA

PMR = Plasma Molecular Response
PMR-Res = $\geq 50\%$ PMR (responsive)
PMR-NR = $< 50\%$ PMR (non-responsive)
MTB = Molecular Tumor Board

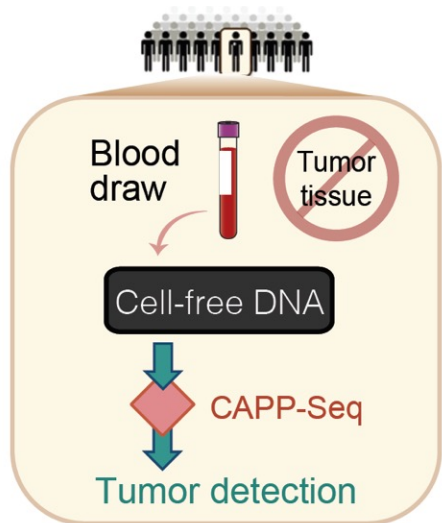


Retrospective Data From ~900 NSCLC Patients: Pre- and Post-treatment MRD strongly prognostic

Study	N	Stage	Treatment(s)	ctDNA assay
Chaudhuri <i>Cancer Discov</i> 2017	37	IB-III B	RT and/or surgery +/- chemo	CAPP-Seq
Abbosh <i>Nature</i> 2017	24	IA-III B	Surgery +/- chemo	Natera
Chen <i>CCR</i> 2019	25	I-III	Surgery +/- chemo	cSMART
Moding <i>Cancer Discov</i> 2020	48	IIB-III B	chemoRT +/- IO	CAPP-Seq
Abbosh <i>AACR</i> 2020	88	I-III	Surgery +/- chemo	ArcherDx
Zviran <i>Nat Med</i> 2020	22	I-III	Surgery +/- chemo	MRDetect
Waldeck <i>Mol Oncol</i> 2021	16	IA-III B	Surgery +/- chemo, RT	Custom NGS
Xia <i>CCR</i> 2021	329	I-III	Surgery +/- chemo	Custom NGS
Gale <i>Ann Oncol</i> 2022	59	I-III	RT and/or surgery +/- chemo	Inivata
Zhang <i>Cancer Discov</i> 2022	245	I-III	Surgery +/- chemo, IO, TKI	Custom NGS

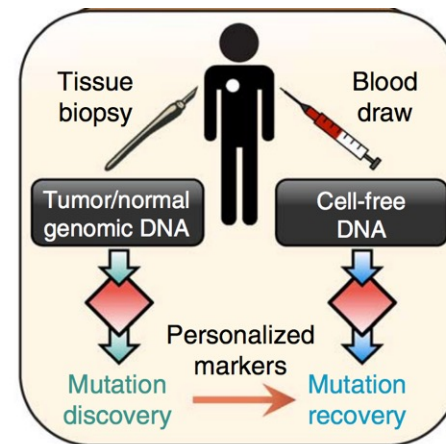
Different types of ctDNA MRD Assays

Tumor-naive



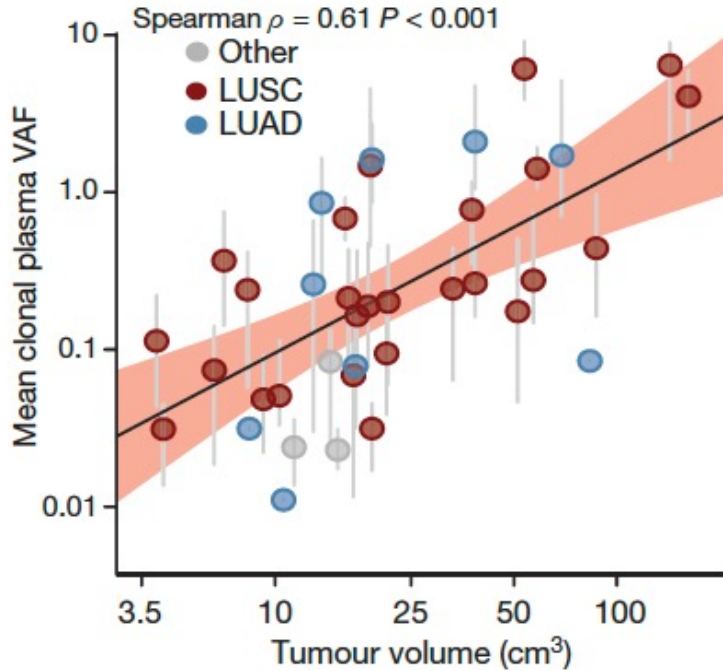
- Genotyping with no knowledge of tumor mutations (“off the shelf”)
- Faster, less expensive
- Limit of detection ~0.1%

Tumor-informed



- Tracking multiple known mutations (bespoke or personalized)
- Requires tumor tissue, time, \$\$
- Limit of detection ~0.01%

Tumour volume predicts plasma VAF



Tumour volume (cm³)

1

Predicted VAF:

0.006%

95% CI (0.001–0.03%)

10

0.1%

(0.05–0.17%)

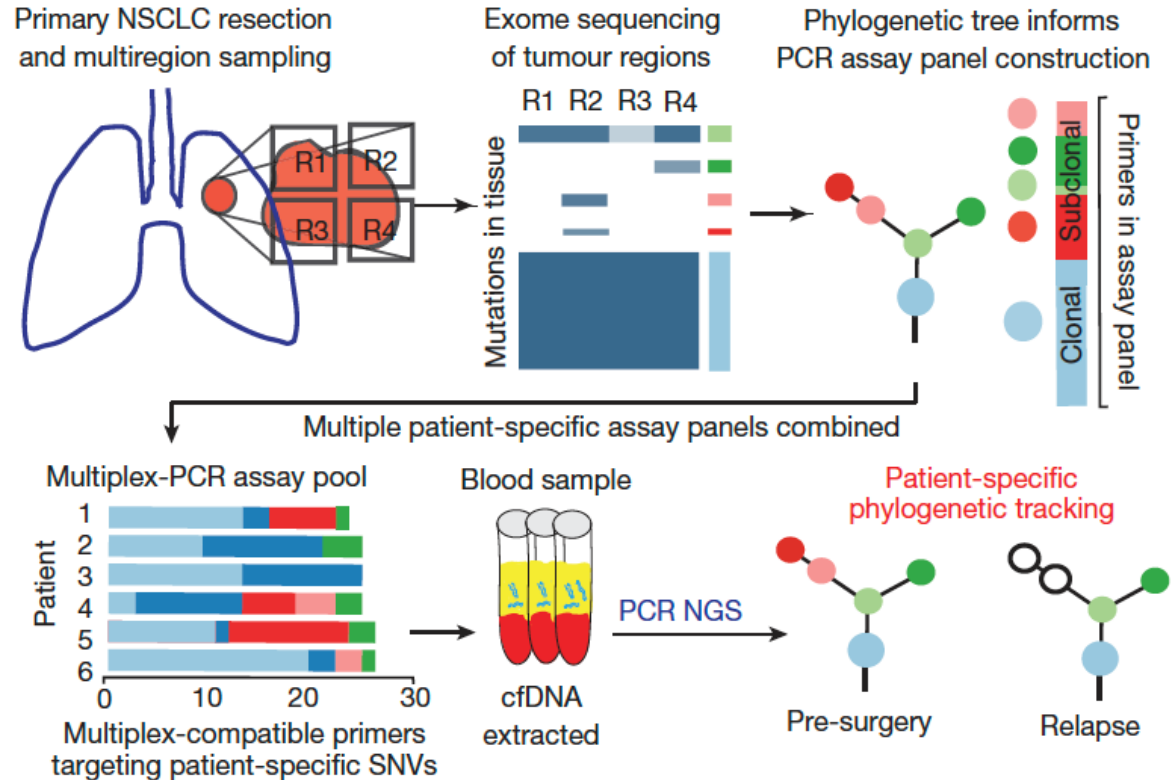
100

1.3%

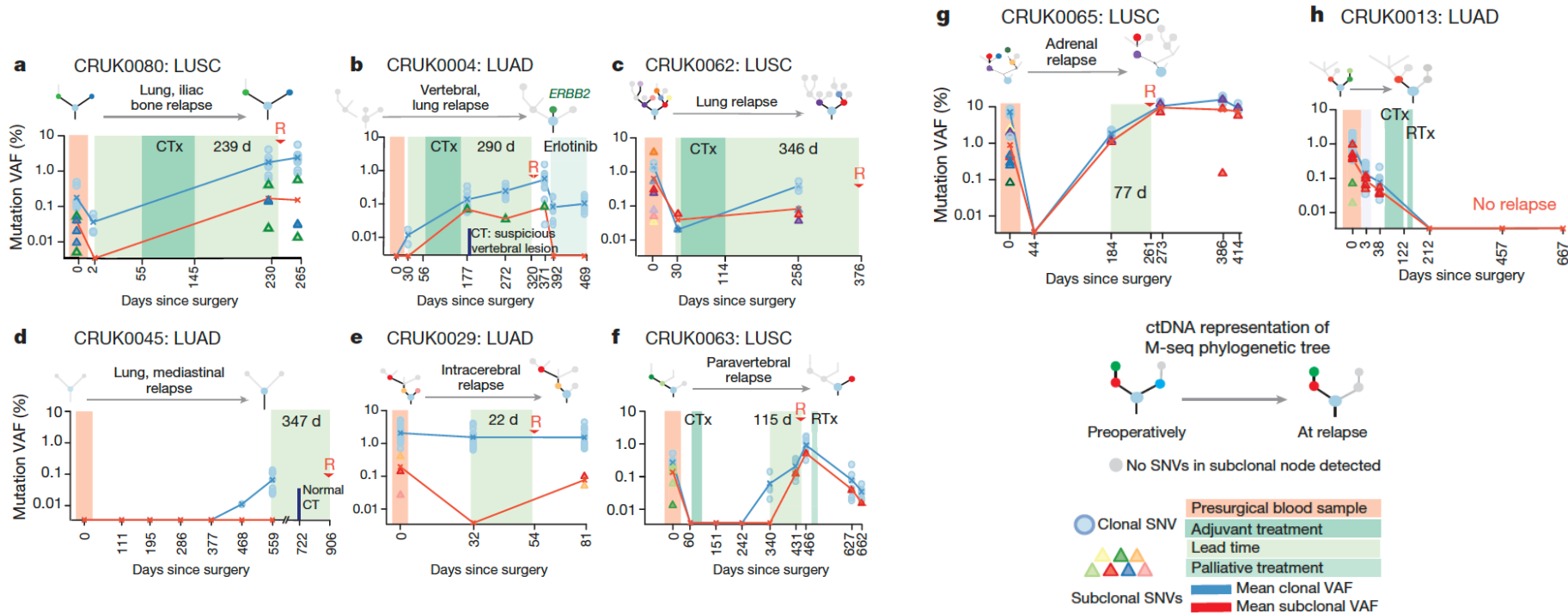
(0.57–3.1%)

▼
(Approximately 326 million malignant cells)

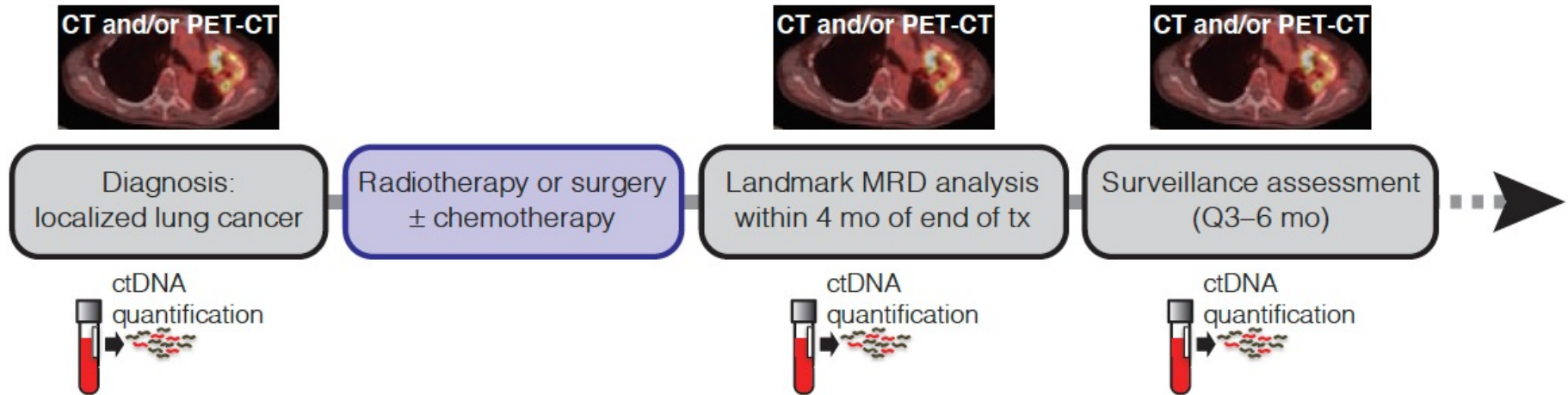
Phylogenetic approach to profile the ctDNA – TRACERx Study



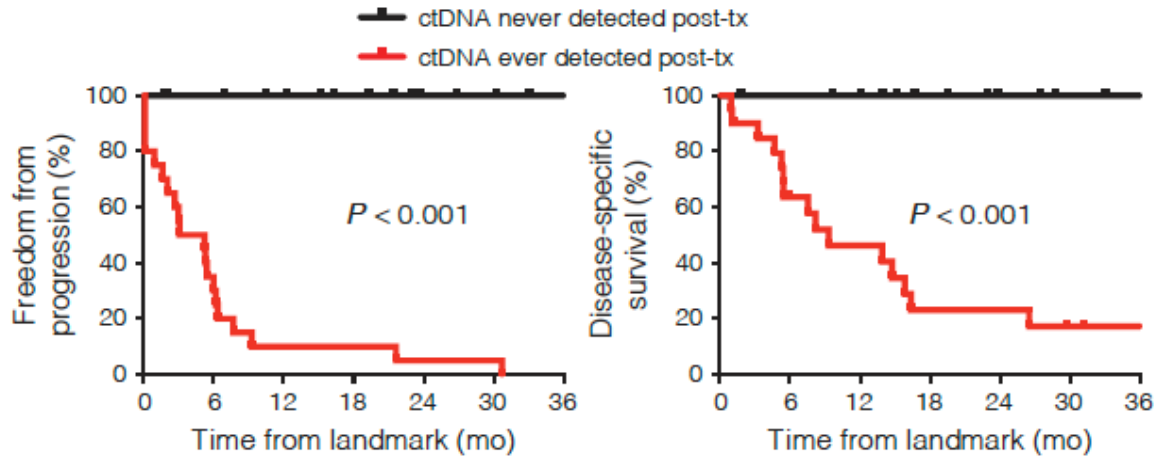
Postoperative ctDNA detection predicts and characterises NSCLC relapse



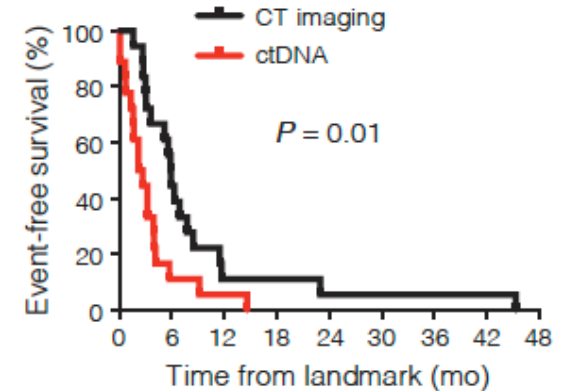
Early Detection of MRD in Localised Lung Cancer by CAPP-Seq



Application of ctDNA analysis for posttreatment surveillance in patients with localised lung cancer

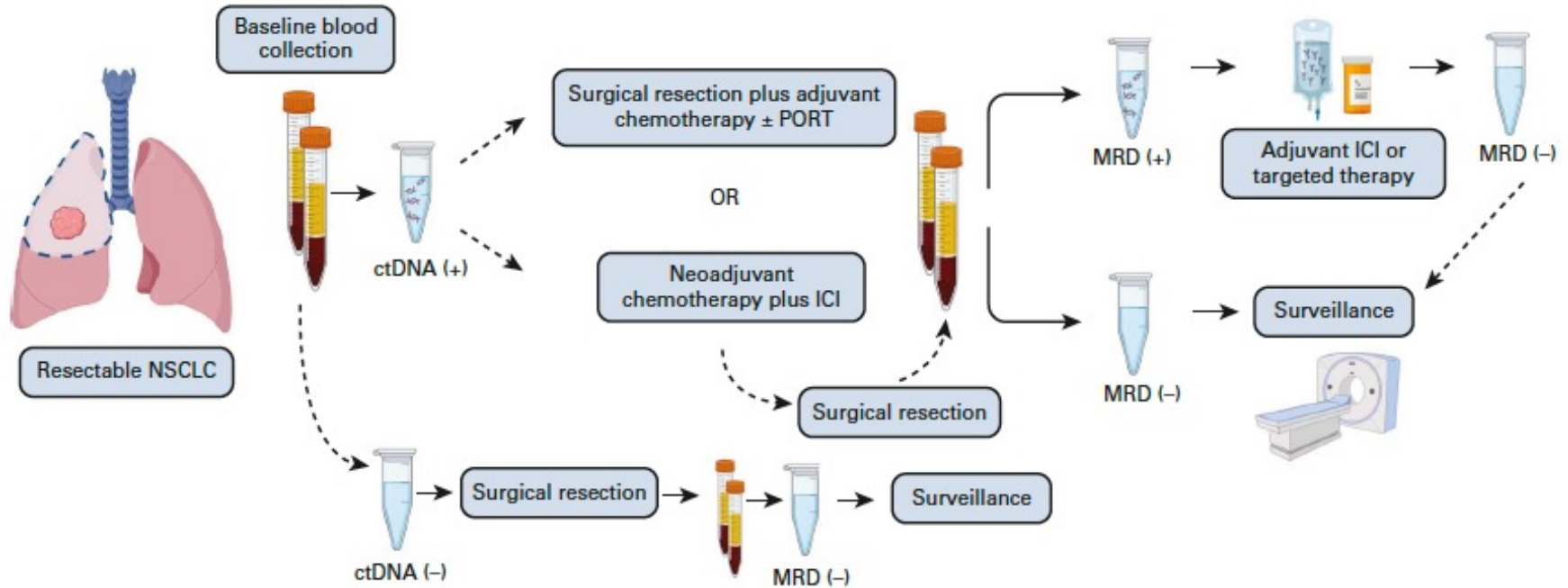


KM curves stratified by ctDNA detection status during posttreatment surveillance

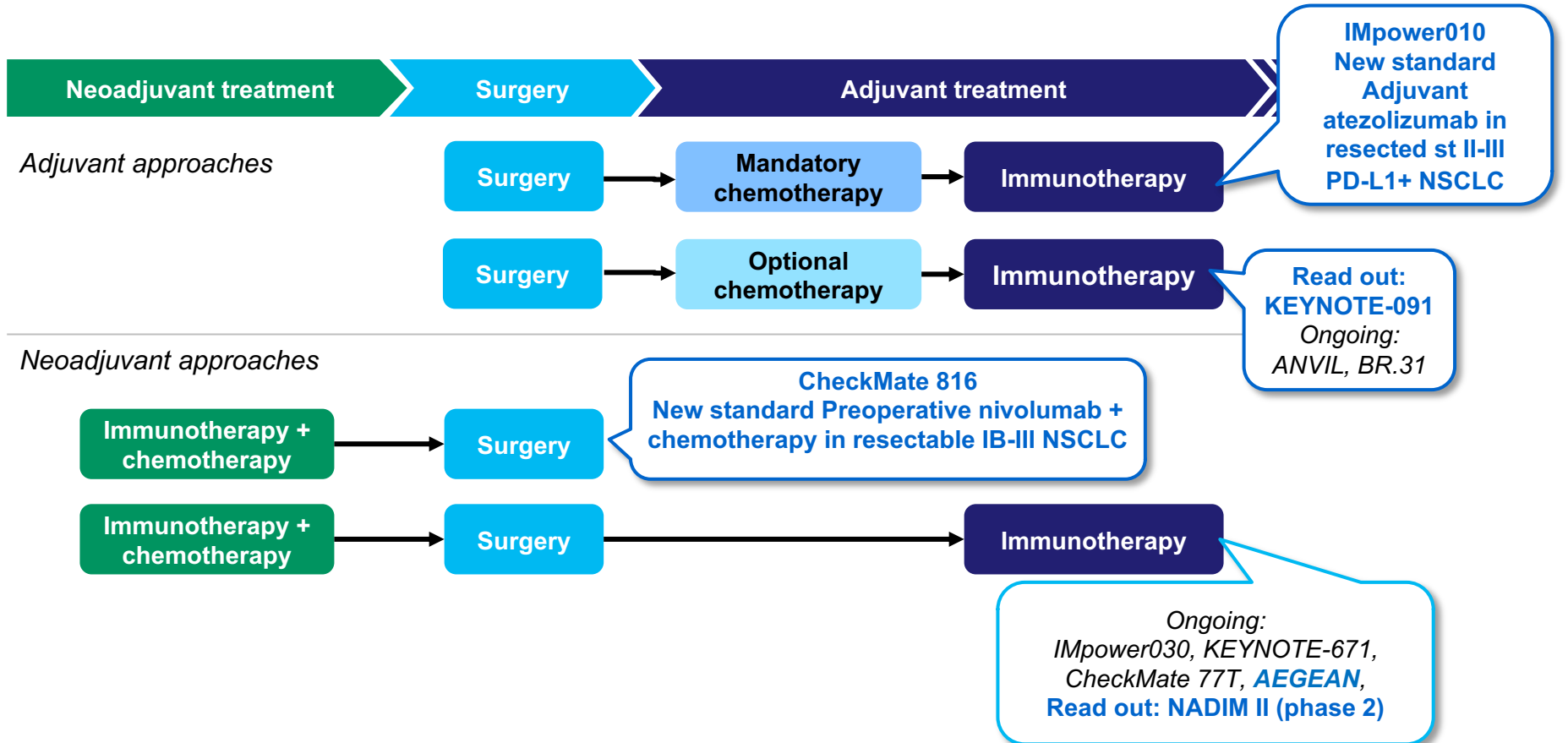


ctDNA detection and time to imaging progression

Proposed clinical trial designs for early-stage NSCLC using ctDNA as a biomarker for treatment personalization

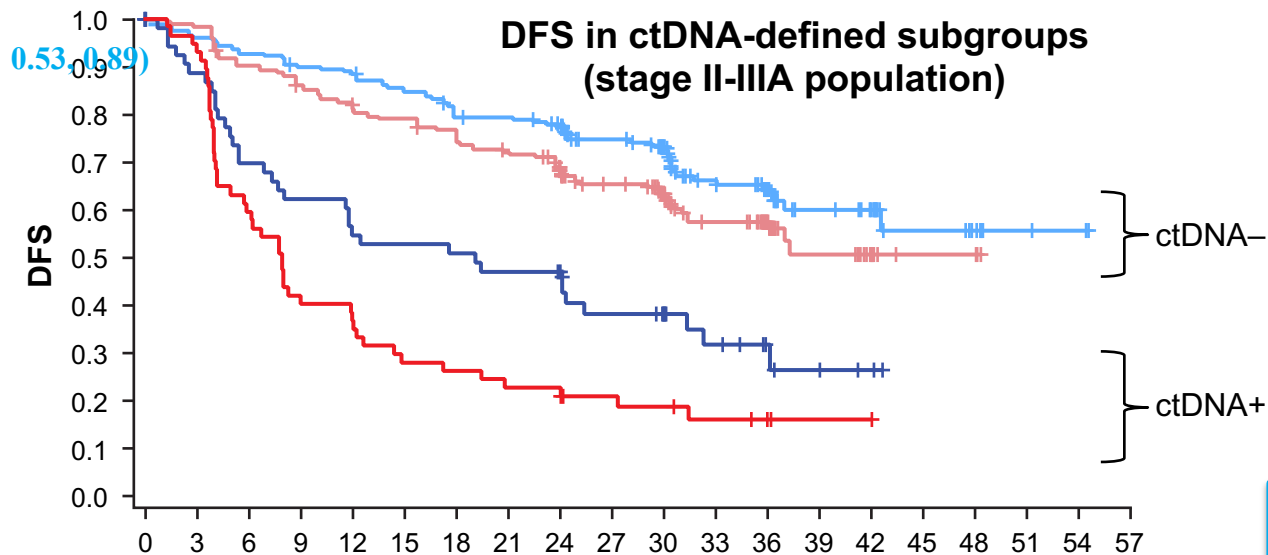


Phase III studies in resectable NSCLC



ctDNA positivity was strongly prognostic, with DFS favouring atezo in both ctDNA+ and ctDNA- patients

In all ctDNA-evaluable stage II-IIIa patients, mDFS was NR (atezo) vs 31.4 months (BSC), with an HR of 0.69 (95% CI: 0.53, 0.89)



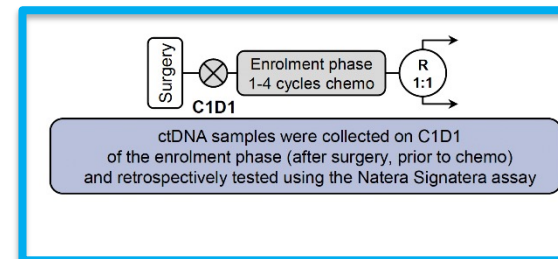
No. at risk

	0	3	6	9	12	15	18	21	24	27	30	33	36	39	42	45	48	51	54	57
Atezo, ctDNA-	218	206	199	192	189	180	170	166	151	131	112	73	58	33	24	12	8	3	2	0
Atezo, ctDNA+	53	47	37	33	29	28	27	25	23	17	14	10	6	3	2	0	0	0	0	0
BSC, ctDNA-	204	193	176	167	158	152	143	137	124	106	88	62	44	19	9	3	3	0	0	0
BSC, ctDNA+	59	53	34	24	21	16	15	13	13	9	8	6	4	1	1	0	0	0	0	0

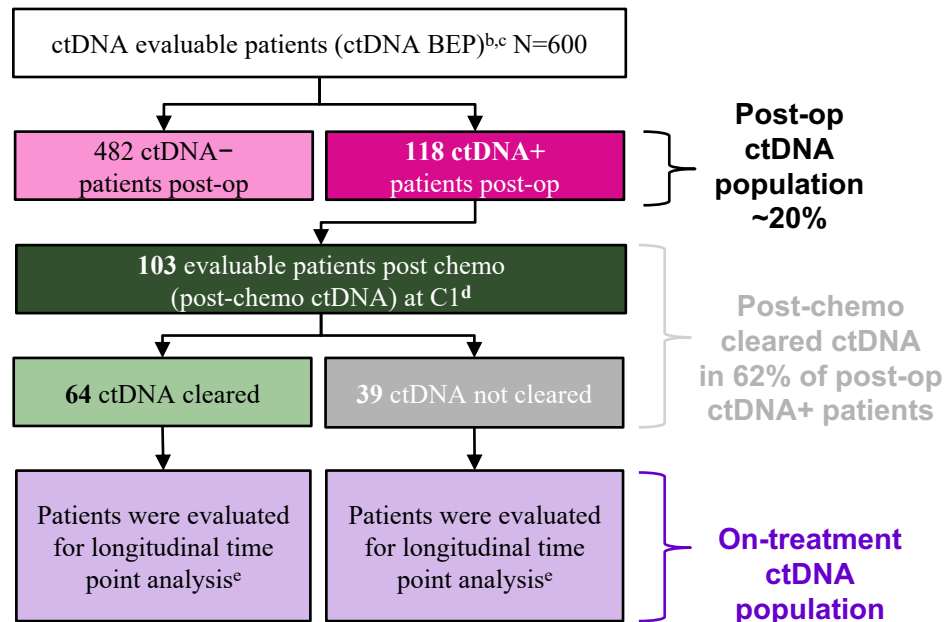
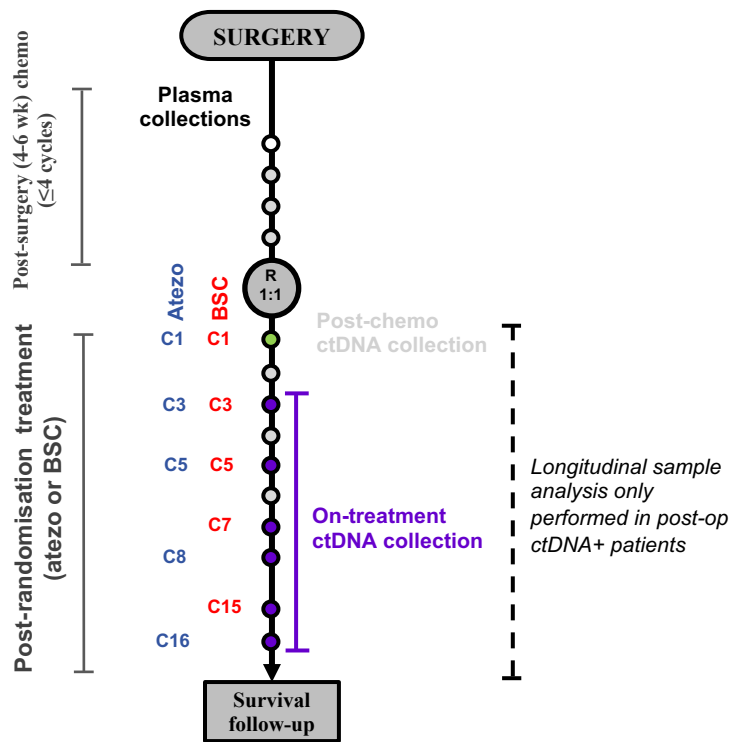
ctDNA-	Atezo (n=218)	BSC (n=204)
mDFS, mo	NR	NR
HR (95% CI)	0.72 (0.52, 1.00)	

ctDNA+	Atezo (n=53)	BSC (n=59)
mDFS, mo	19.1	7.9
HR (95% CI)	0.61 (0.39, 0.94)	

Plasma collection for ctDNA analysis



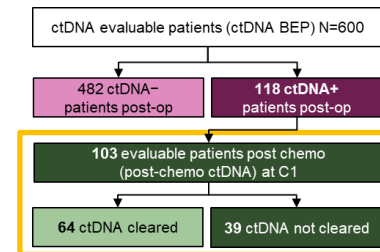
Baseline and longitudinal plasma collection for ctDNA testing^a



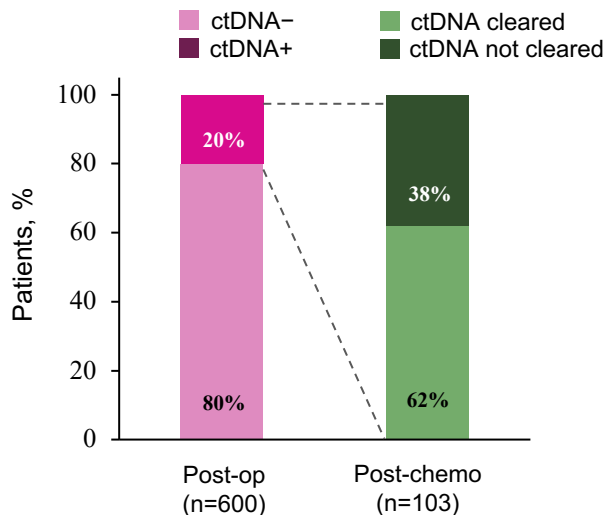
Chemo, chemotherapy; C, cycle. Clinical cutoff: 21 January 2021. ^a Using the Signatera (Natera) RUO test. ^b Treatment arms in the ctDNA BEP were balanced and comparable to the ITT population. ^c PD-L1 subgroup analyses conducted in the stage II-III A ctDNA BEP (n=532). ^d Samples in 15 patients were missing due to lack of consent or 4 mL plasma. ^e Patients with ≥1 on-treatment sample at C3, C5, C7/8 and C15/16. On-treatment analyses are shown on slides 9 (ctDNA cleared) and 10 (ctDNA not cleared).

ctDNA clearance with adjuvant chemo in post-op ctDNA+ patients

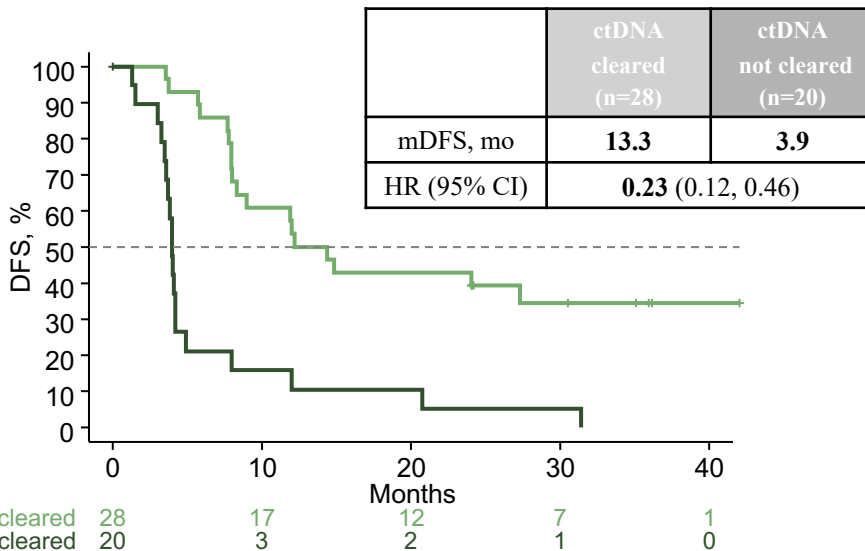
- Adjuvant chemo was effective in clearing ctDNA in $\approx 62\%$ of post-op ctDNA+ patients
- Post-chemo ctDNA positivity was linked to poor DFS outcome



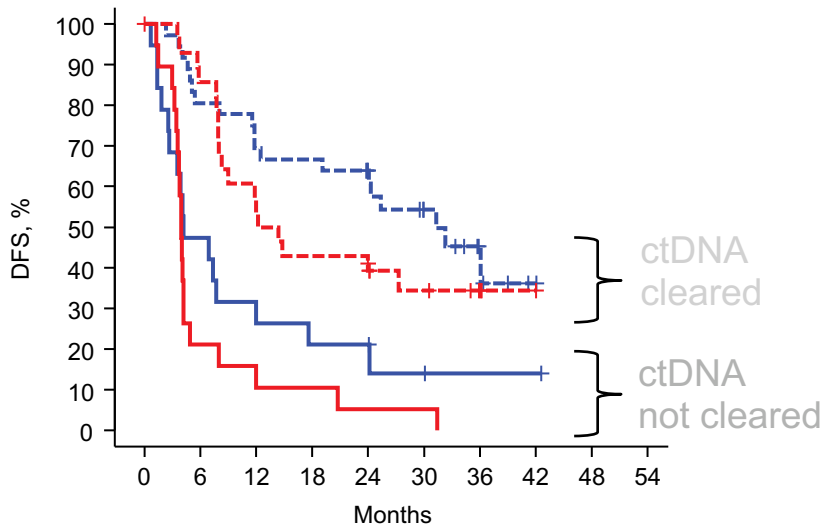
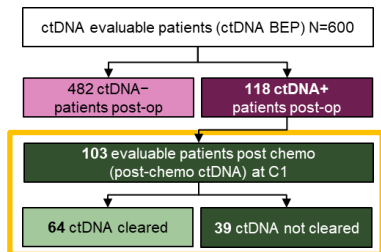
Impact of chemo on ctDNA clearance status



DFS by ctDNA clearance status in the BSC arm



DFS by treatment and post-chemo ctDNA clearance - all groups still appear to benefit from atezolizumab



ctDNA cleared	Atezo (n=36)	BSC (n=28)
mDFS, mo	31.3	13.3
HR (95% CI)	0.7 (0.37, 1.34)	

ctDNA not cleared	Atezo (n=19)	BSC (n=20)
mDFS, mo	4.2	3.9
HR (95% CI)	0.67 (0.34, 1.32)	

Atezo, ctDNA cleared	36	35	29	28	25	24	24	23	21	17	12	10	5	2	1	0	0	0	0
Atezo, ctDNA not cleared	19	13	9	6	5	5	4	4	4	2	2	1	1	1	1	0	0	0	0
BSC, ctDNA cleared	28	28	24	18	15	12	12	12	12	8	7	6	4	1	1	0	0	0	0
BSC, ctDNA not cleared	20	16	4	3	2	2	2	1	1	1	1	0	0	0	0	0	0	0	0

Clinical cutoff: 21 January 2021.

Data are hypothesis generating and should be interpreted with caution due to the exploratory nature of the analysis and small sample size.

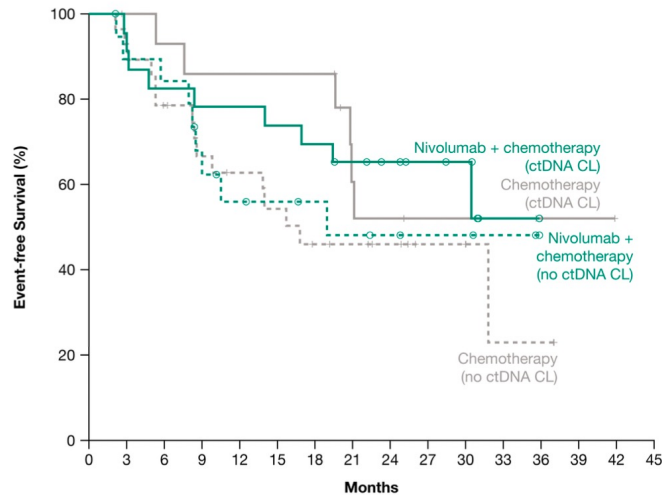
Modified from Dr. Felip, ESMO IO 2022

Courtesy Dr. Natasha Leigh

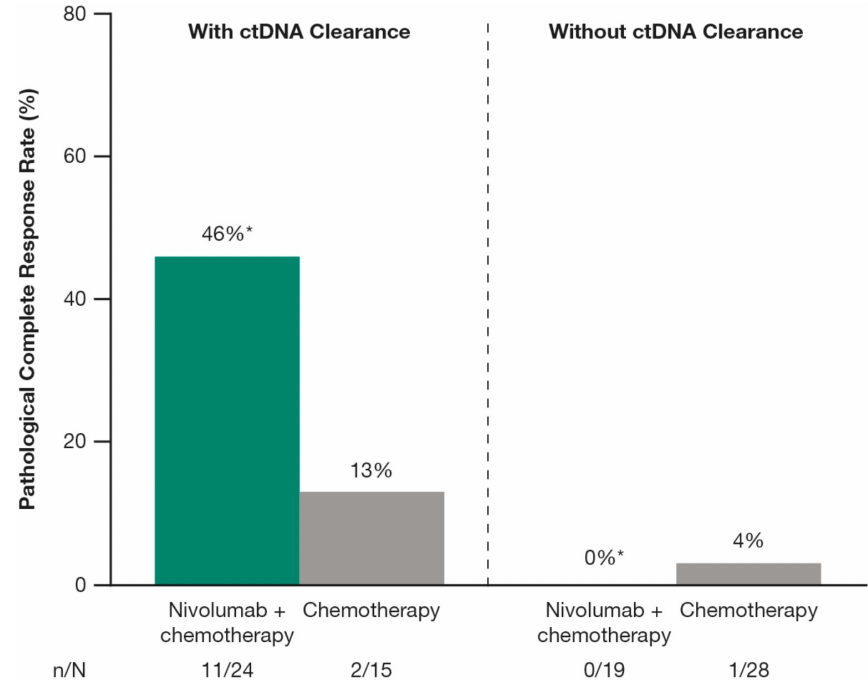
Liquid Biopsy in Neoadjuvant IO + chemo combination

WES ctDNA in 89 pts

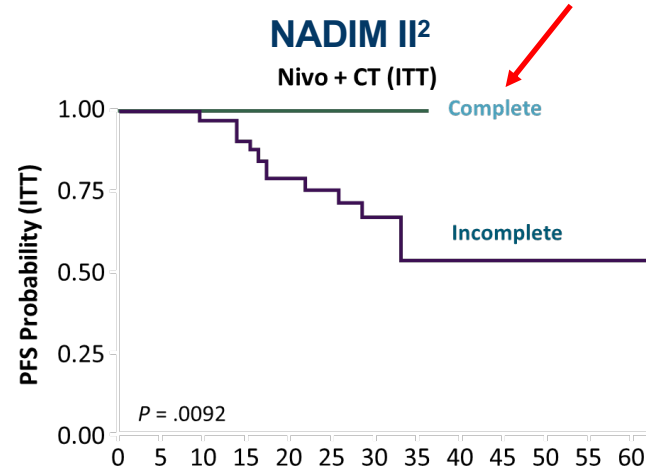
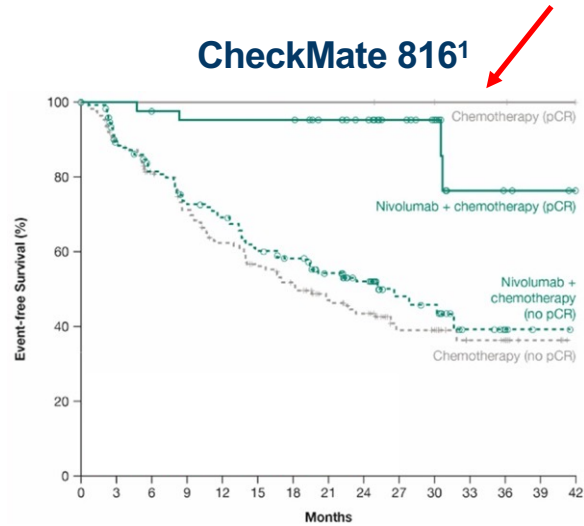
	Nivolumab + chemotherapy		Chemotherapy	
	ctDNA CL (n=24)	No ctDNA CL (n=19)	ctDNA CL (n=15)	No ctDNA CL (n=28)
Median EFS, mo (95% CI)	NR (16.8–NR)	18.9 (8.3–NR)	NR (19.6–NR)	16.8 (8.3–NR)
HR (95% CI)	0.60 (0.20–1.82)		0.63 (0.20–2.01)	



	No. at Risk															
	24	21	19	18	18	17	16	13	11	8	7	1	0	0	0	0
Nivolumab + chemotherapy (ctDNA CL)	24	21	19	18	18	17	16	13	11	8	7	1	0	0	0	0
Chemotherapy (ctDNA CL)	15	14	13	12	12	12	12	7	6	5	5	5	3	1	0	0
Nivolumab + chemotherapy (no ctDNA CL)	19	17	16	12	9	8	7	6	5	3	3	2	0	0	0	0
Chemotherapy (no ctDNA CL)	28	26	21	17	15	13	10	9	7	4	3	1	1	0	0	0



Pathologic complete response - a more promising surrogate endpoint



	Nivo + CT		CT	
	pCR	No pCR	pCR	No pCR
mEFS, months	NR	26.6	NR	18.4
HR (95% CI)	0.13 (0.05, 0.37)		Not computed*	

Months from randomisation

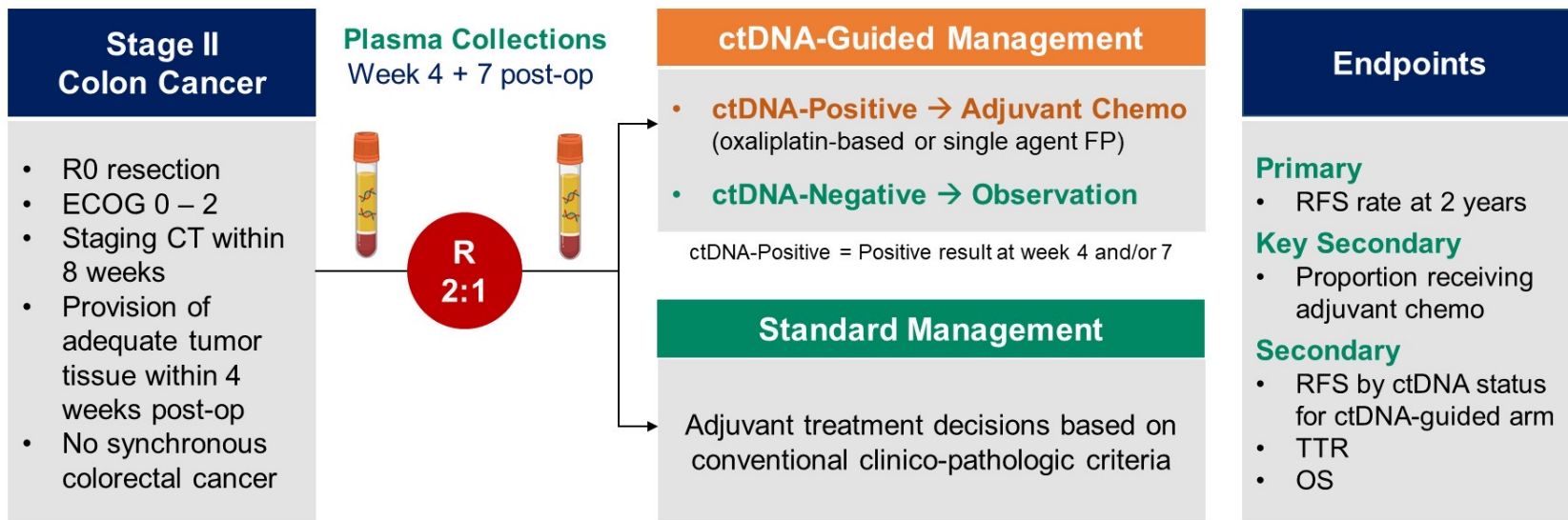
Patients at Risk, n	0	5	10	15	20	25	30	35	40	45	50	55	60
Complete	21	21	21	21	15	10	5	1	0	0	0	0	0
Incomplete	35	35	34	32	22	21	10	4	1	1	1	1	1

Current prospective interventional trials in early stage lung cancer

Number	Prior tx	Stage	N	ctDNA-positive intervention	ctDNA-negative intervention	Phase	Primary Endpoint	Site(s)
NCT04585477	Surgery or RT +/- chemo	I-III	80	Durvalumab	None	II	ctDNA change	Stanford
NCT04585490	chemoRT + several cycles durvalumab	III	48	Durvalumab + chemo	None	II	ctDNA change	Stanford
NCT04966663	Surgery	I	66	Nivolumab + chemo <u>vs.</u> No treatment	None	II	RFS	Toronto

DYNAMIC Study Design

ACTRN12615000381583



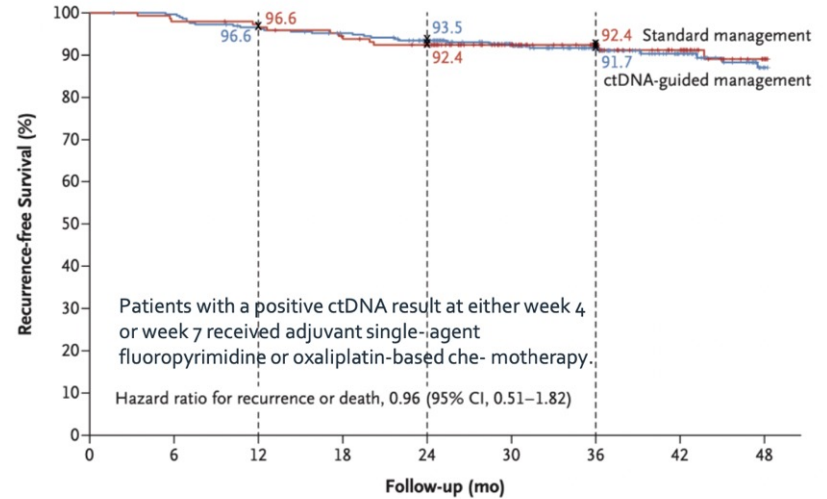
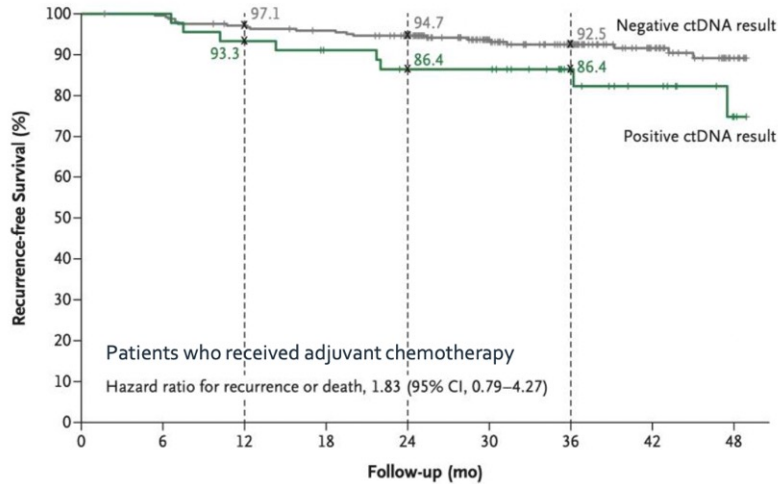
Stratification Factors

- T stage (T3 vs T4)
- Type of participating center (metropolitan vs regional)

Surveillance:

- CEA → 3-monthly for 24M, then 6-monthly for 36M
- CT C/A/P → 6-monthly for 24M, then at 36M

ctDNA-guided adjuvant therapy had similar outcomes to stage-directed treatment



No. at Risk

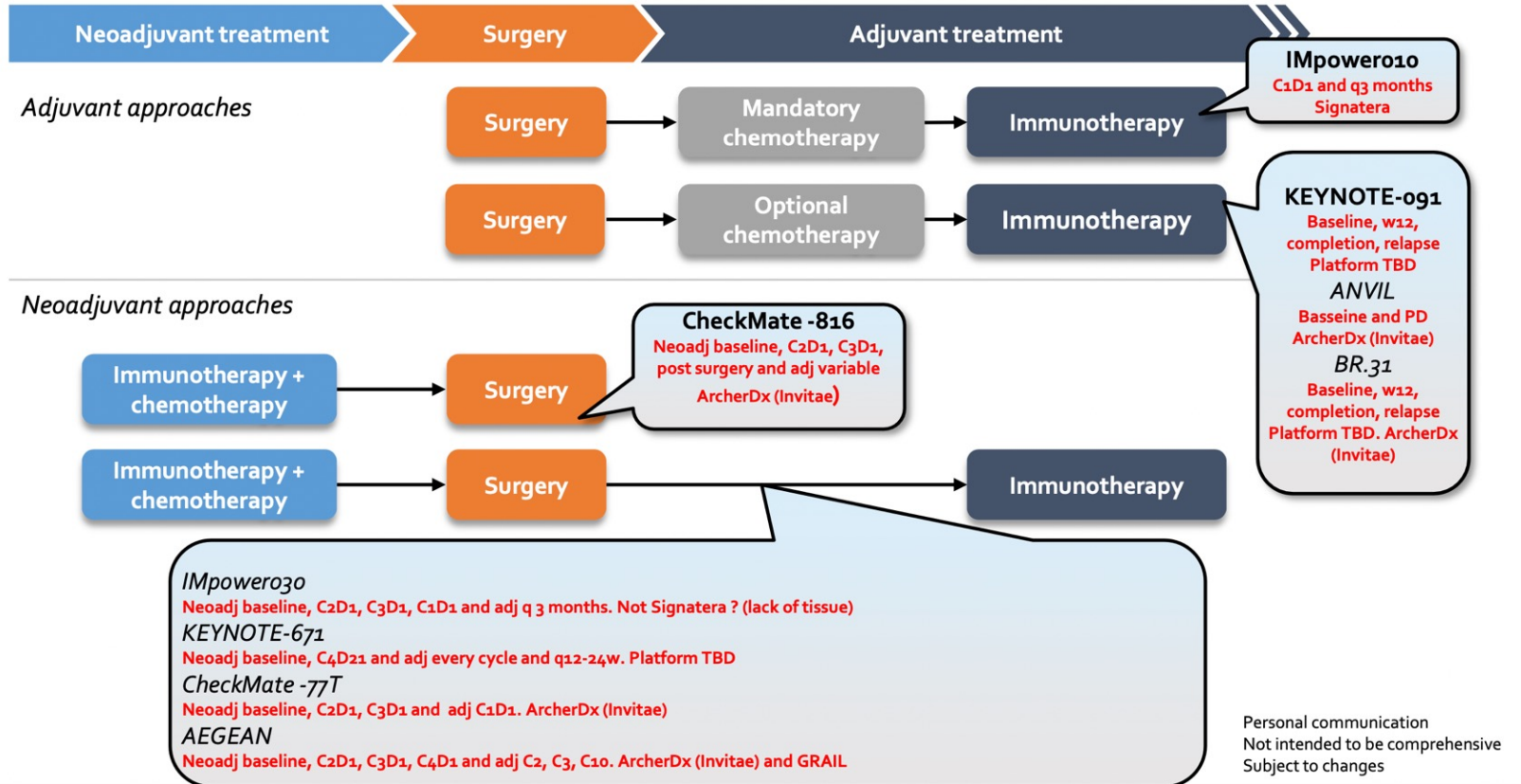
Negative ctDNA result

Positive ctDNA result

246	244	236	231	220	169	131	93	55
45	45	42	39	36	36	22	16	9

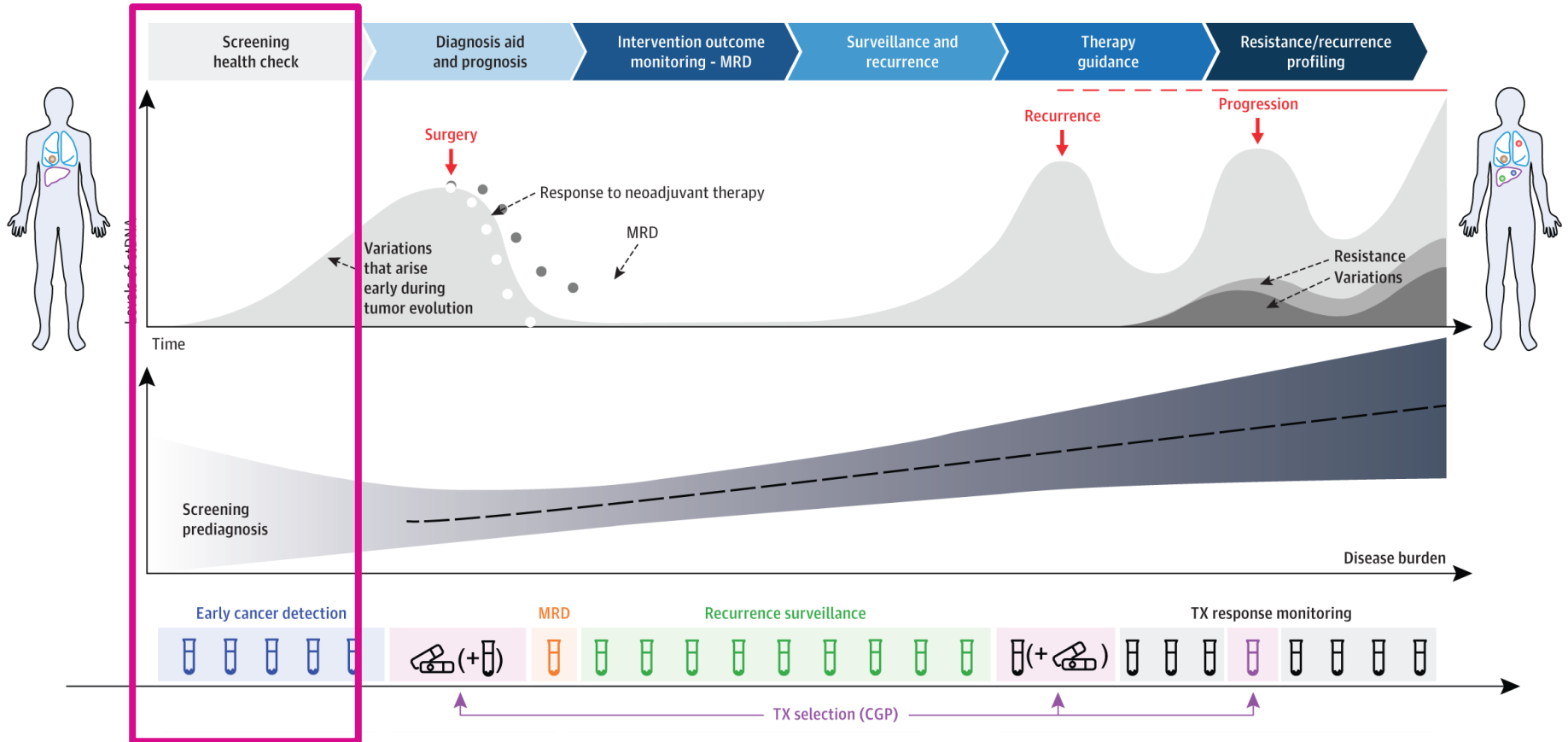
- 455 patients randomized, 302 were assigned to ctDNA-guided management and 153 to standard management
- 15% of patients in the ctDNA-guided group vs 28% in standard-management group received adjuvant chemotherapy
- ctDNA-guided management was noninferior to standard management
- Safe-Sequencing System tumor-informed personalized ctDNA assays (tumor-informed personalized approach)

More data are on the way!

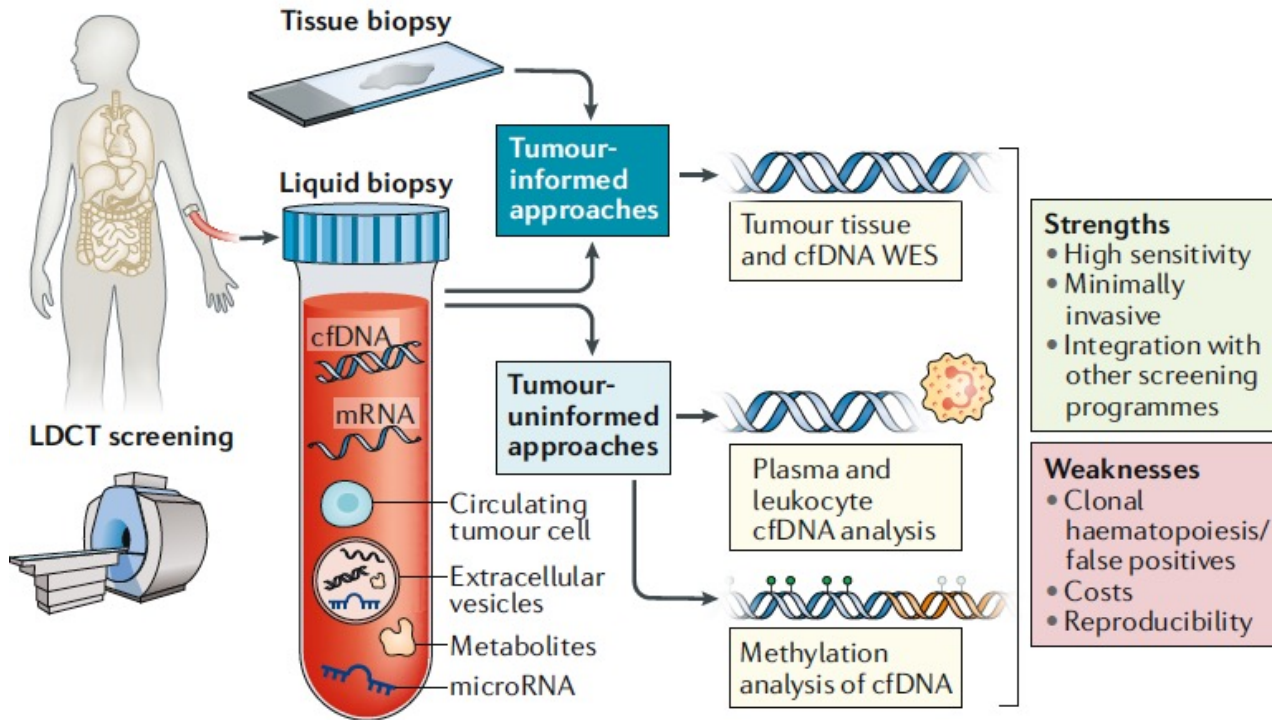


Take Home Message on MRD

- Detecting MRD is crucial to improve survival and disease control rates
- Liquid Biopsy is a perfect tool for MRD
- MRD at difference of early detection, counts with tissue and liquid biopsy as a source of information, increasing the possibilities
- Integrating liquid biopsy in clinical trials is a necessity
- Real time monitoring in patients with high risk of recurrence requires improved technology in liquid biopsy
- Other analytes in liquid biopsy as exosomes or CTCs can go beyond cfDNA and offer opportunities in research and possible in clinical practice.



Liquid biopsy & early detection: Strengths and weaknesses of currently used approaches



**Sensitivity at
98% specificity**

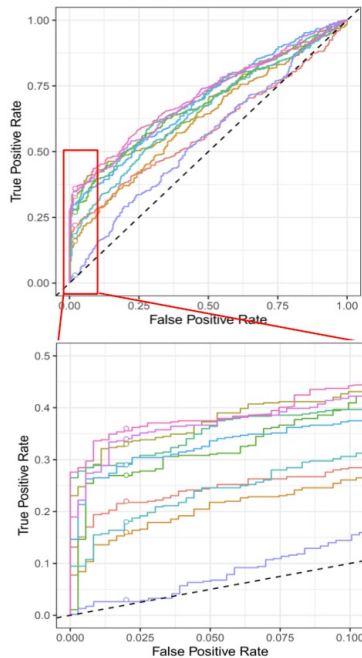
WGBS

● **WG methylation**

34% (30%-39%)

158/464

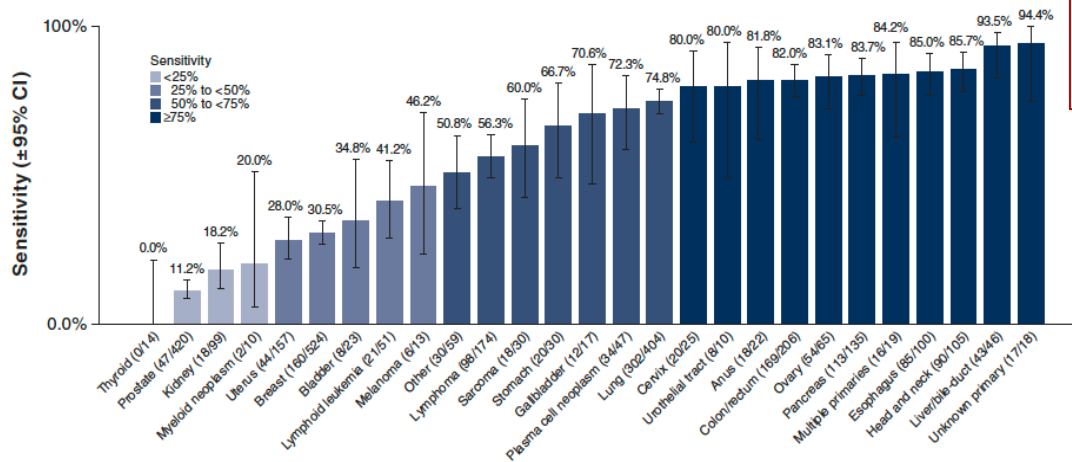
Performance



	Cancer	Non-cancer	Total
	2823	1254	4077
Test positive	1453	6	1459
Test negative	1370	1248	2618
	Sensitivity = 1453/2823 51.5% (49.6%-53.3%)	Specificity = 1248/1254 99.5% (99.0%-99.8%)	

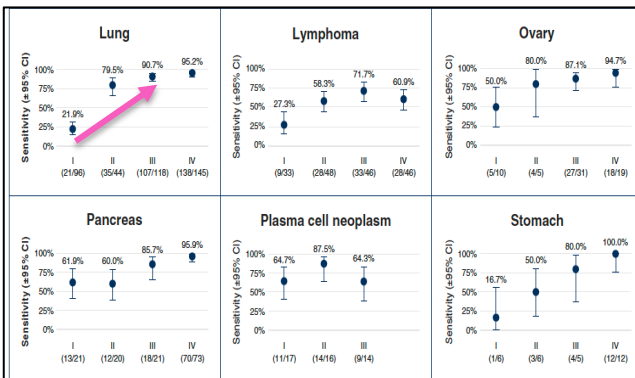
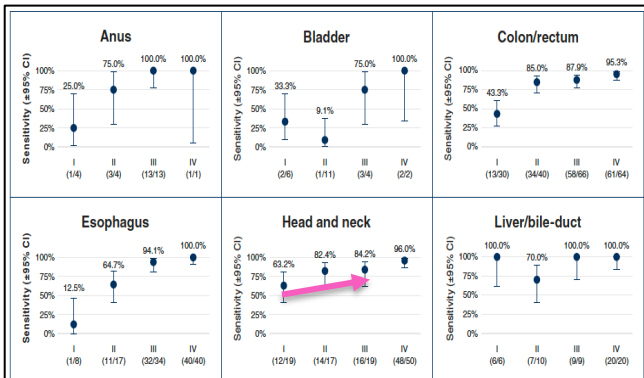
Two-sided 95% Wilson confidence intervals were calculated.

**Confirmed status analysis set, n = 4077
cancer, n = 2823; non-cancer, n = 1254**



Clinical cancer stage, n (%)

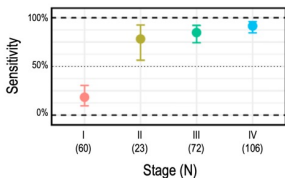
I	849 (30.1)
II	703 (24.9)
III	566 (20.0)
IV	618 (21.9)



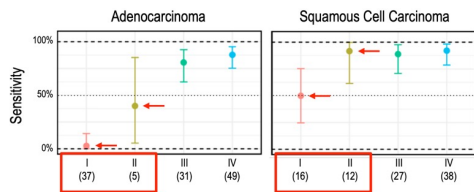
Clinical stage	Total N	Test positive	Sensitivity % (95% CI) ^a
All	2823	1453	51.5 (49.6% to 53.3%)
I	849	143	16.8 (14.5% to 19.5%)
II	703	284	40.4 (36.8% to 44.1%)
III	566	426	77.0 (73.4% to 80.3%)
IV	618	557	90.1 (87.5% to 92.2%)
I-II	1552	427	27.5 (25.3% to 29.8%)
I-III	2118	863	40.7 (38.7% to 42.9%)
I-IV	2736	1420	51.9 (50.0% to 53.8%)
III-IV	1184	993	83.9 (81.7% to 85.9%)
Not expected to be staged	67	23	34.3 (24.1% to 46.3%)
Missing	20	10	50.0 (29.9% to 70.1%)

All subtypes have the same sensitivity?

Lung Cancer Detection Varies by Subtype at 99.4% Specificity

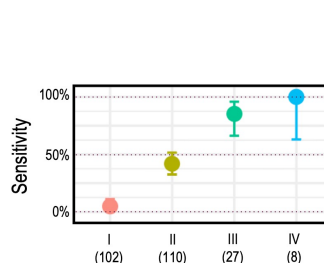


- Detection rate affected by early-stage adenocarcinomas
 - Detection higher in squamous cell carcinoma
- Consistent with prior report showing ctDNA detection was higher in squamous cell carcinoma than adenocarcinoma¹

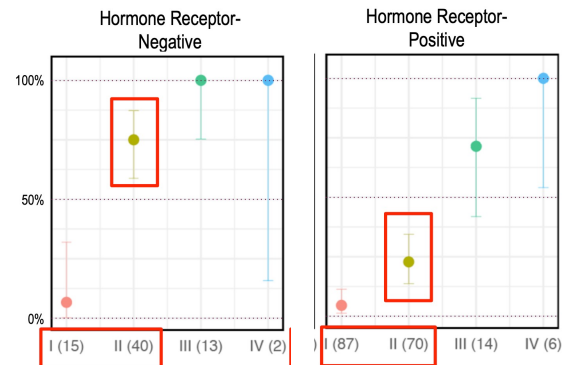


- Overall lung cancer sensitivity: 71.6% (95% CI: 65.8-77.0%)

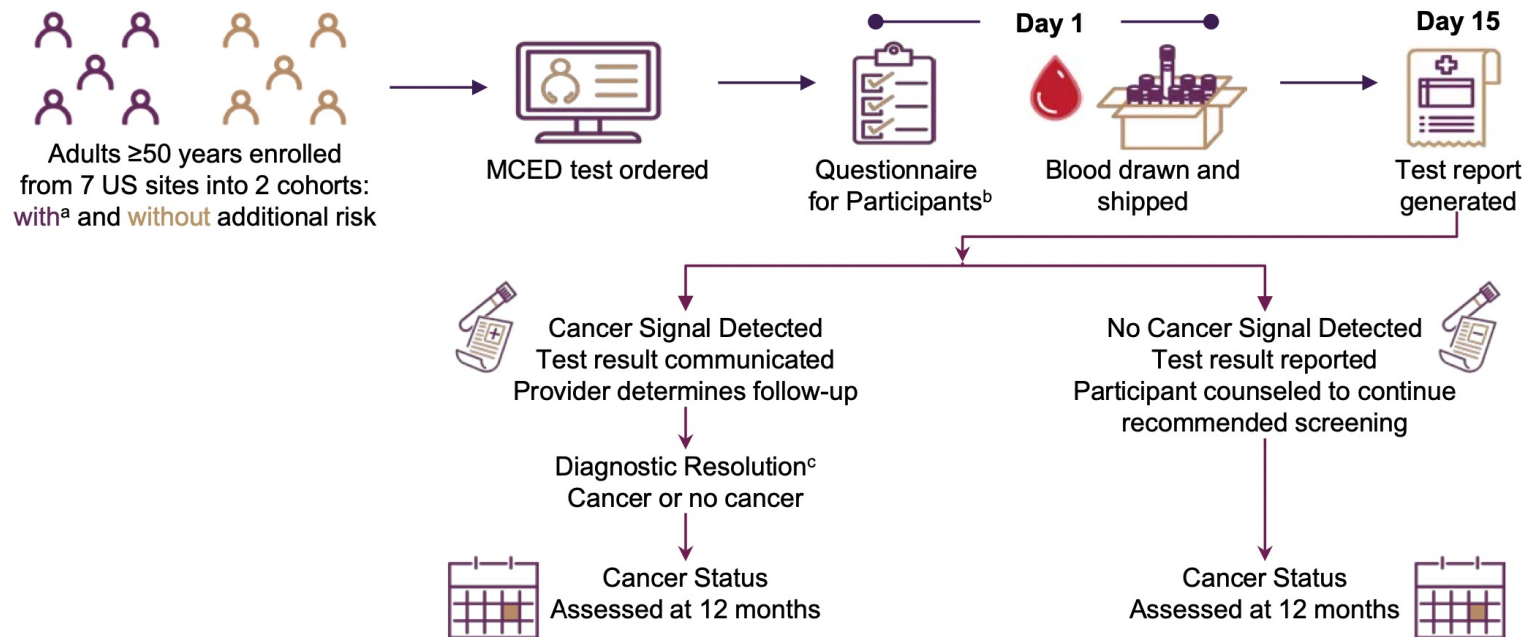
Breast Cancer Detection Varies by Subtype at 99.4% Specificity



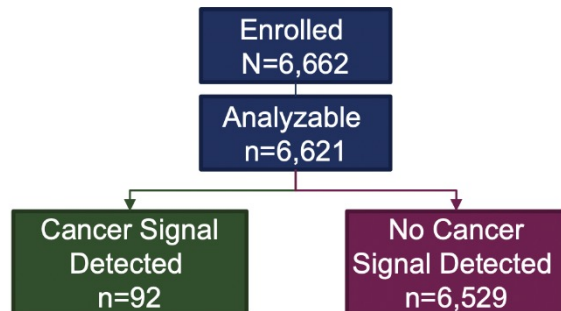
- Overall breast cancer sensitivity: 33.2% (95% CI: 27.4-39.4%)



PATHFINDER: A Prospective Cohort Study to Return the Results of MCEd Tests to Participants



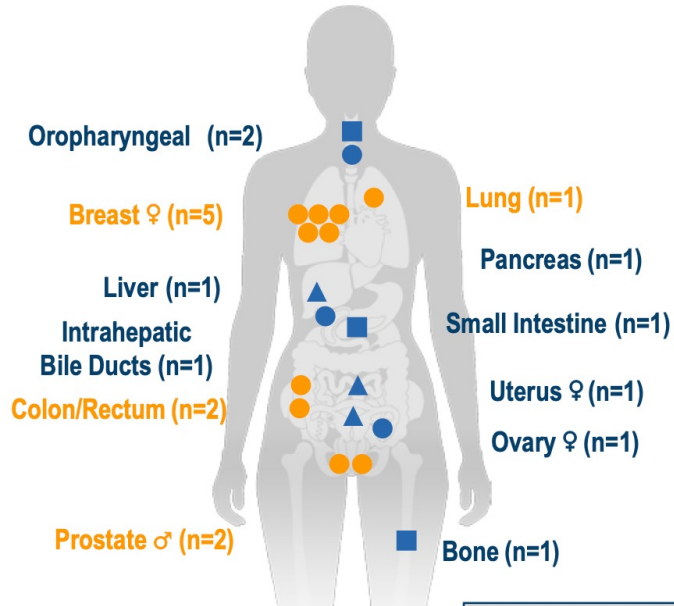
	With Additional Risk ^a n = 3,681	Without Additional Risk n = 2,940	Total N = 6,621
Age ^b , in years, mean (SD)	64.7 (8.7)	61.6 (8.1)	63.4 (8.6)
Female	65%	62%	63%
White, Non-Hispanic	93%	89%	92%
College Degree or Higher	59%	71%	65%
Up to Date With Standard Cancer Screening Prior to MCED Testing			
Colorectal Cancer ^c	91%	92%	92%
Breast Cancer ^d	78%	83%	80%



	With Additional Risk ^a n = 3,681	Without Additional Risk n = 2,940	Total N = 6,621
Signal Detected	1.5%	1.2%	1.4%
No Signal Detected	98.5%	98.8%	98.6%

Cancers Diagnosed After a True Positive MCED Signal

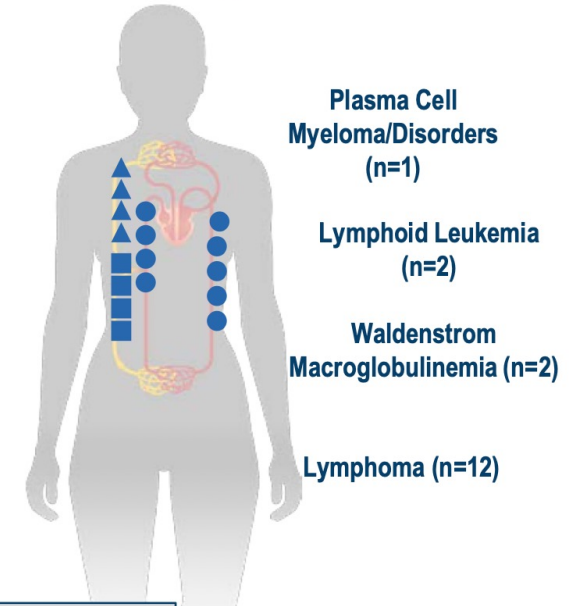
19 Solid Tumors



35 people were diagnosed with 36 cancers

- 24 in high-risk cohort
- 11 in not-high-risk cohort
- 7 recurrent cancers
- 14 early-stage cancers
- 26 cancers lacking standard screening

17 Hematologic Cancers



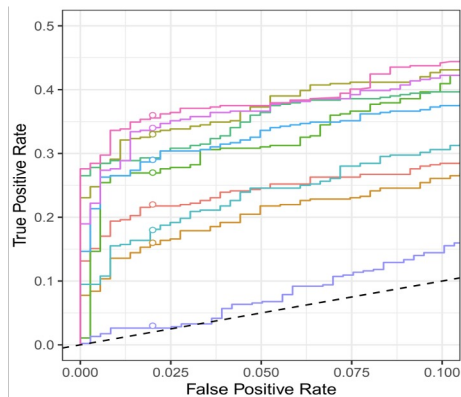
AJCC Staging: ▲ Stage I ■ Stage II ● Stage III/IV/No Stage/Recurrent

Available Screening: **USPSTF cancer screening** or **No standard screening**

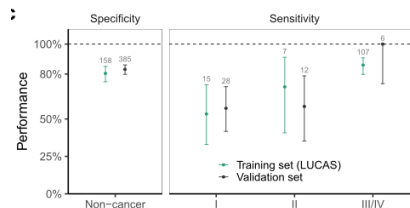
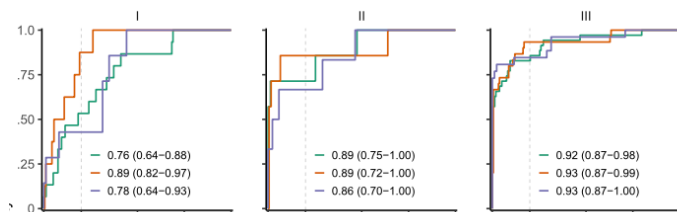
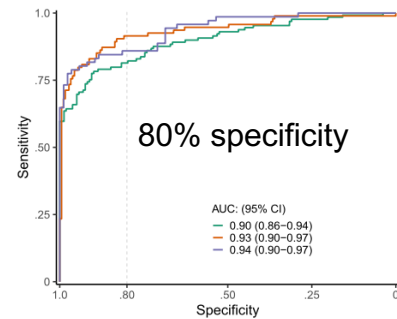
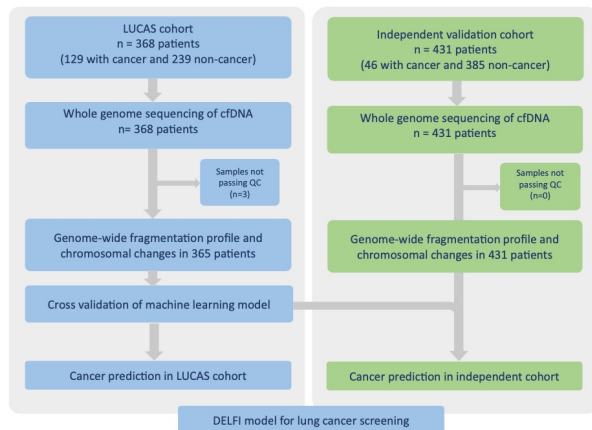
Fragmentomics in a Single-tumor test

Sensitivity at 98% specificity

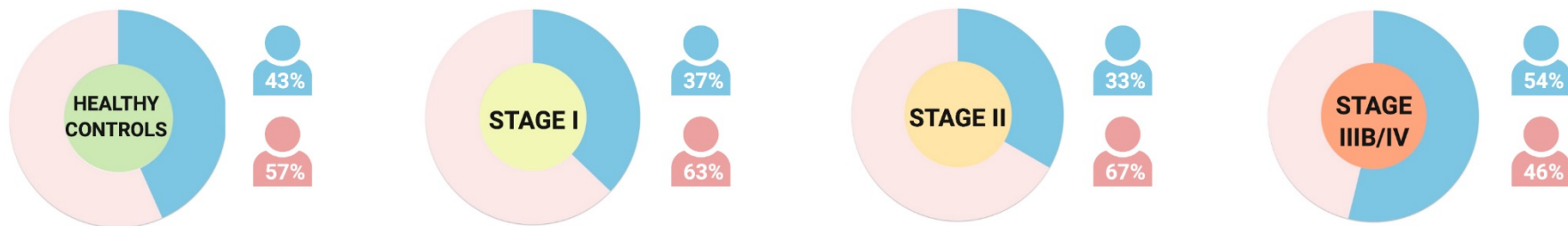
● Fragment endpoints	18% *** (15%-22%)	84/464
● Fragment lengths	29% * (25%-34%)	135/464



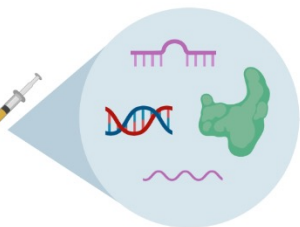
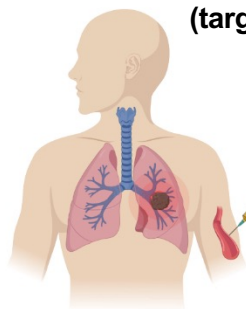
Non-cancer individuals: 236
Patients with Lung cancer 129



A High-Performing Plasma Metabolite Panel for Early-Stage Lung Cancer Detection

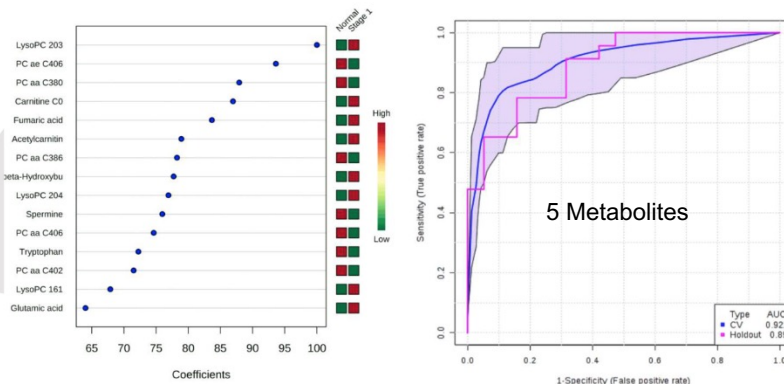


quantitative targeted mass spectrometry (MS) analysis
(targeting 138 metabolites) was performed on all samples

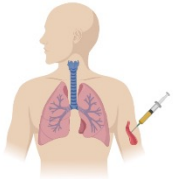


PLASMA SAMPLES
(n=156 NSCLC patients)

LC-MS

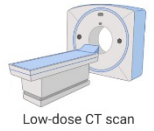


NSCLC Early detection



112 Patients with malignant nodules

39 Patients with benign nodules



Low-dose CT scan



Blood



Plasma



Size-exclusion chromatography

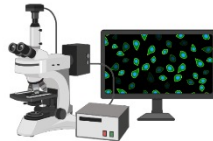


Extracellular vesicles



Circulating tumor cells

FISH CTC detection



10µL Plasma

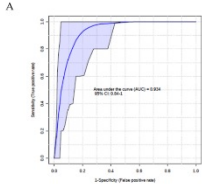
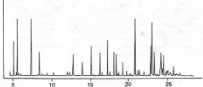


LC-MS/MS



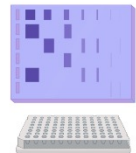
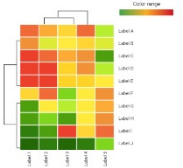
ICP MS

Metabolomics



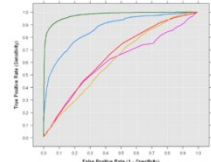
LC-MS/MS

EV Proteomics



Immunoblot & ELISA validation

Multomics detection





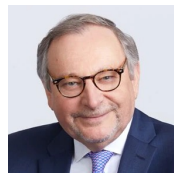
CTO As facilitator of Multidisciplinary Expert Multi-omics approach in Lung Cancer Early Detection

Screening Program

Biomarkers Lab: Liquid & Tissue

Pulmonology

Surgery



Claudia Henscke

David Yankelevitz

Christian Rolfo

D. De Miguel

Phil Mack

Fred Hirsch

Javier Zulueta

David Steiger

Daniel Nicastrì

Radiomics

Proteomics and RNAsq
in EVs

cfDNA

RNAsq Tissue

Lung Nodule Clinic

External Collaboration

Statistician

Epidemiologist

Internal Support

CTCs

Metabolomics



Hsin-Hui (Vivien) Huang



Emanuela Taioli



**Mount
Sinai**

*Innovation
Partners*

Take home message

- ▶ Early detection is crucial to increase survival rates in cancer
- ▶ Liquid Biopsy is an ideal tool to make it possible
- ▶ We need methods to complement the screening programs
- ▶ But also methods standing by their self
- ▶ Multi-cancer detection or single tumor? Not clear yet
- ▶ Important to include risk populations in trials
- ▶ We are not yet in the best scenario that we could be...
- ▶ But we are not far to get it!



@ChristianRolfo



**Join us in 2023
Madrid, Spain**



5TH ANNUAL CONGRESS

Liquid Biopsy

19 - 21 November 2023 | Madrid, Spain

**Save the Date
19-21, November
2023**