

The importance to Use ctDNA to Detect Minimal Residual Disease (MRD)

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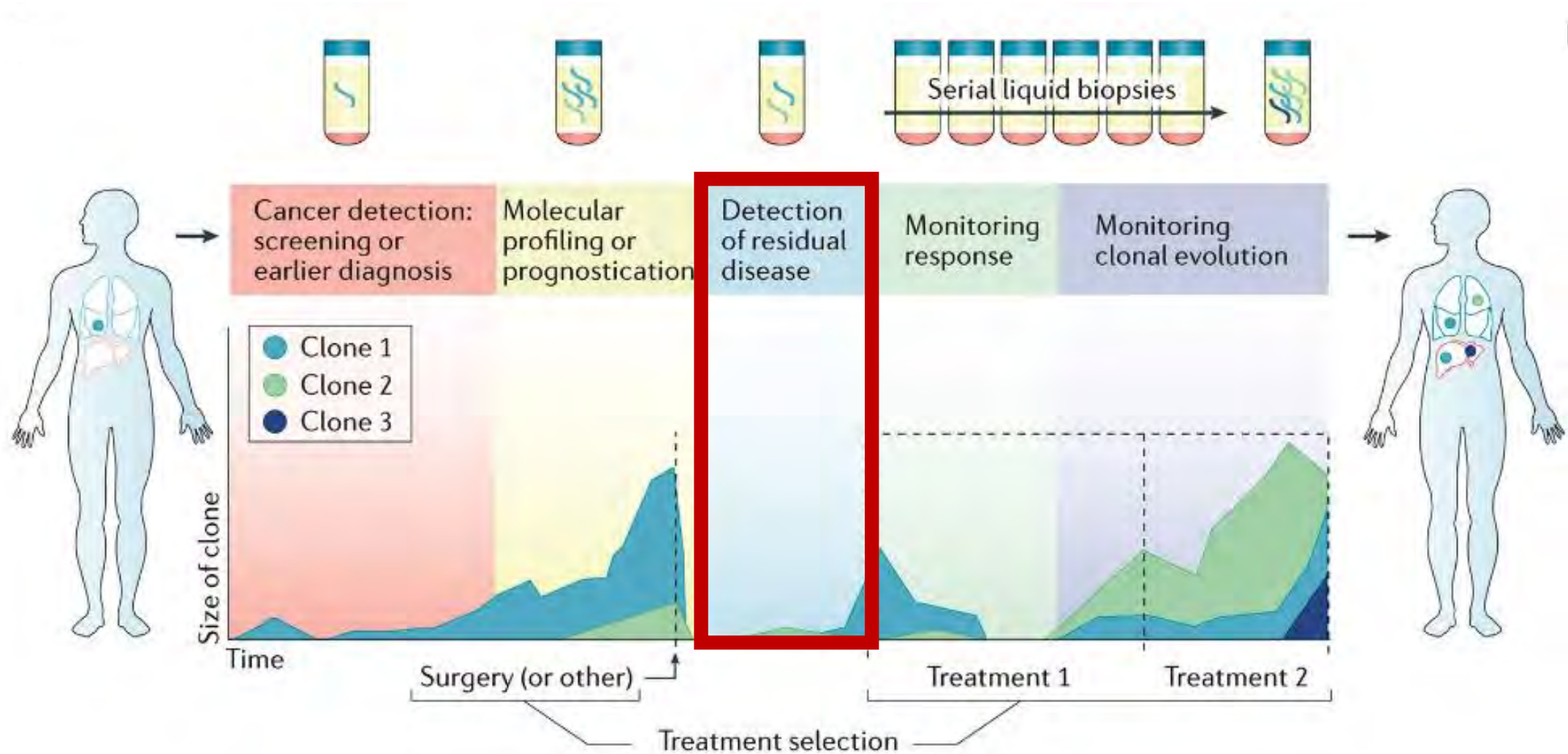
@BrunaPellini



Disclosures:

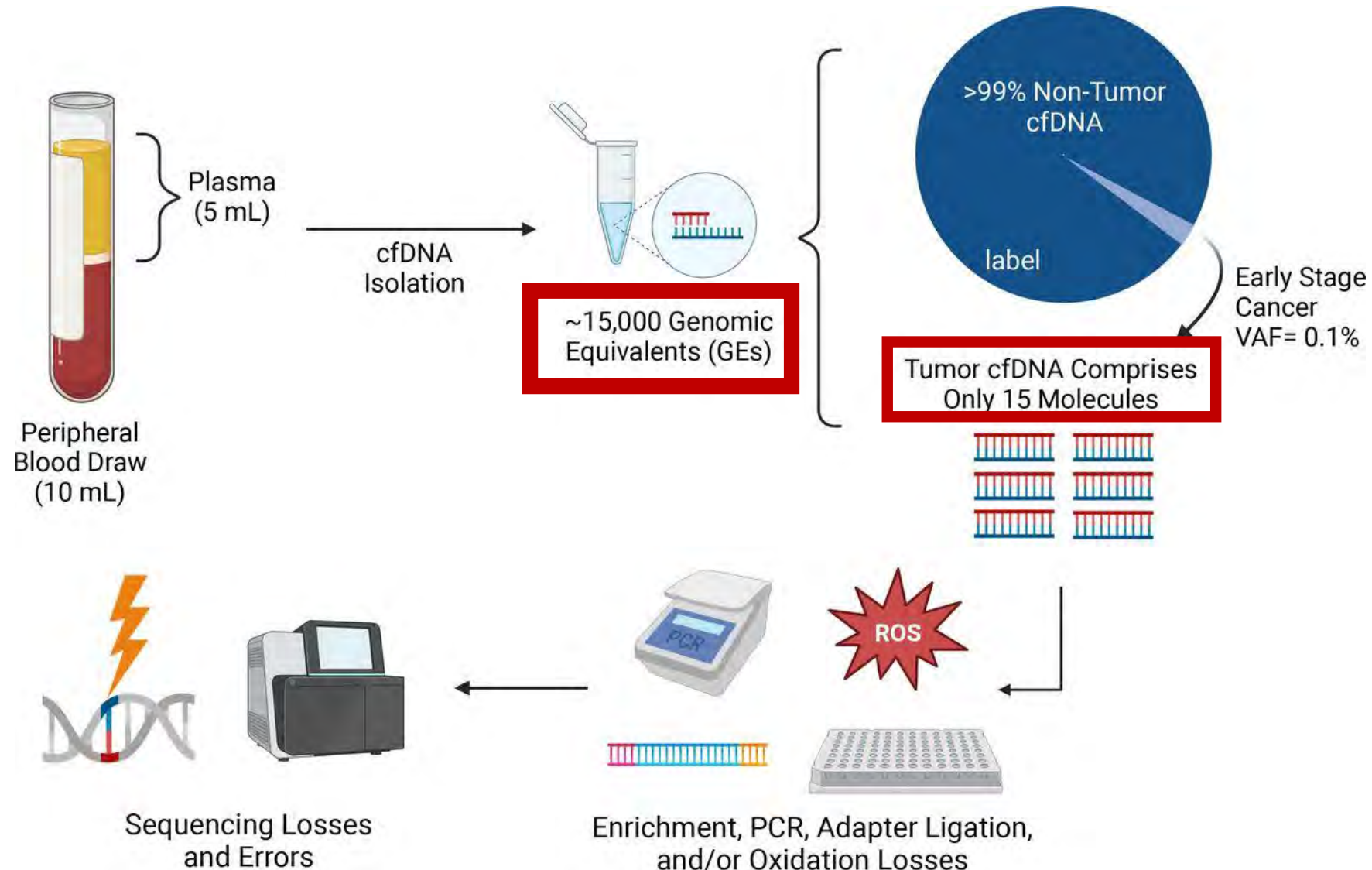
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ctDNA applications in thoracic oncology



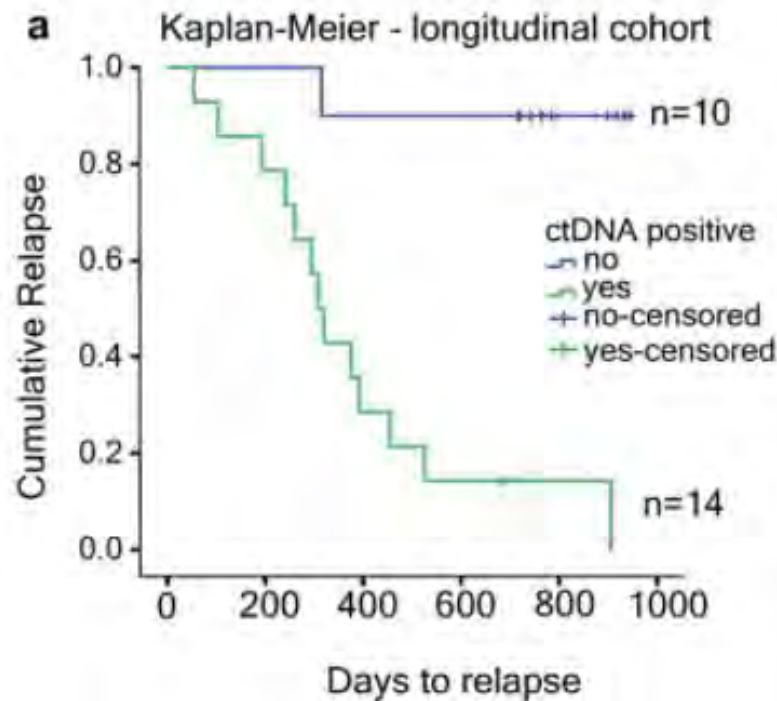


Challenges of ctDNA detection in MRD settings



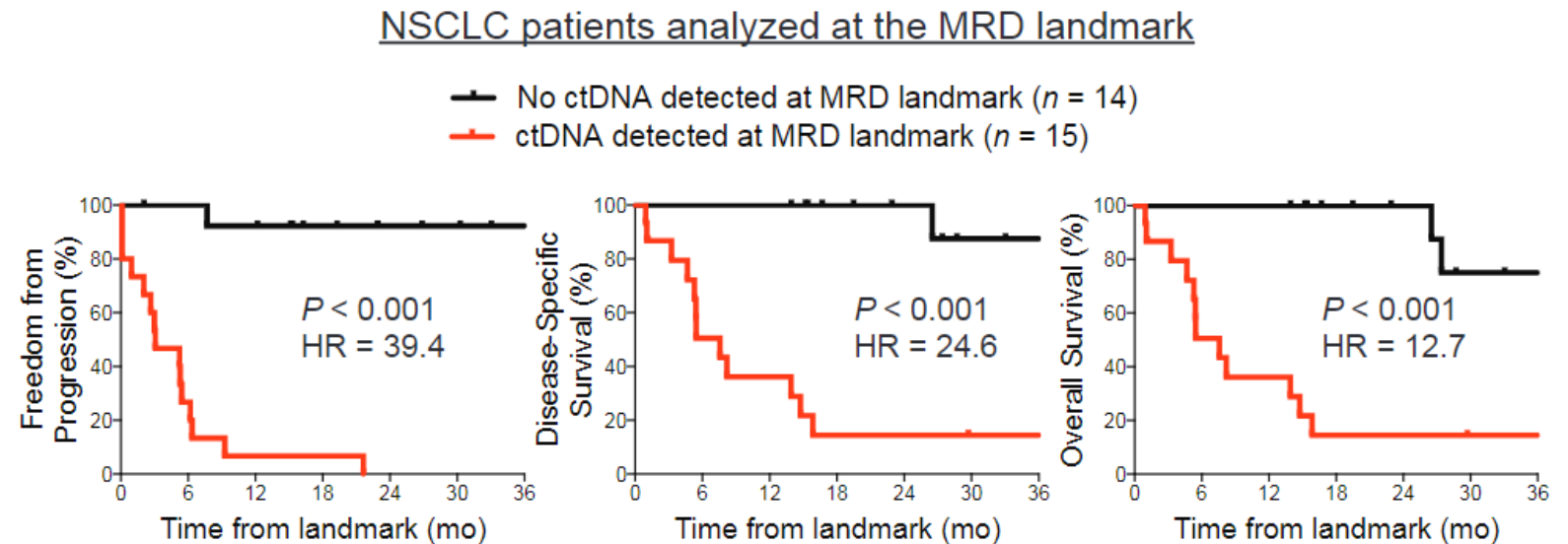
Multiple studies have shown that ctDNA can be used to detect minimal residual disease (MRD) and it is a powerful prognostic biomarker

Stages I-III NSCLC
Tumor-informed assay
(Signatera™)



LOD= 0.01%

Stages I-III NSCLC
Tumor-naïve assay
(CAPP-Seq)



LOD= 0.002%
(0.1%)

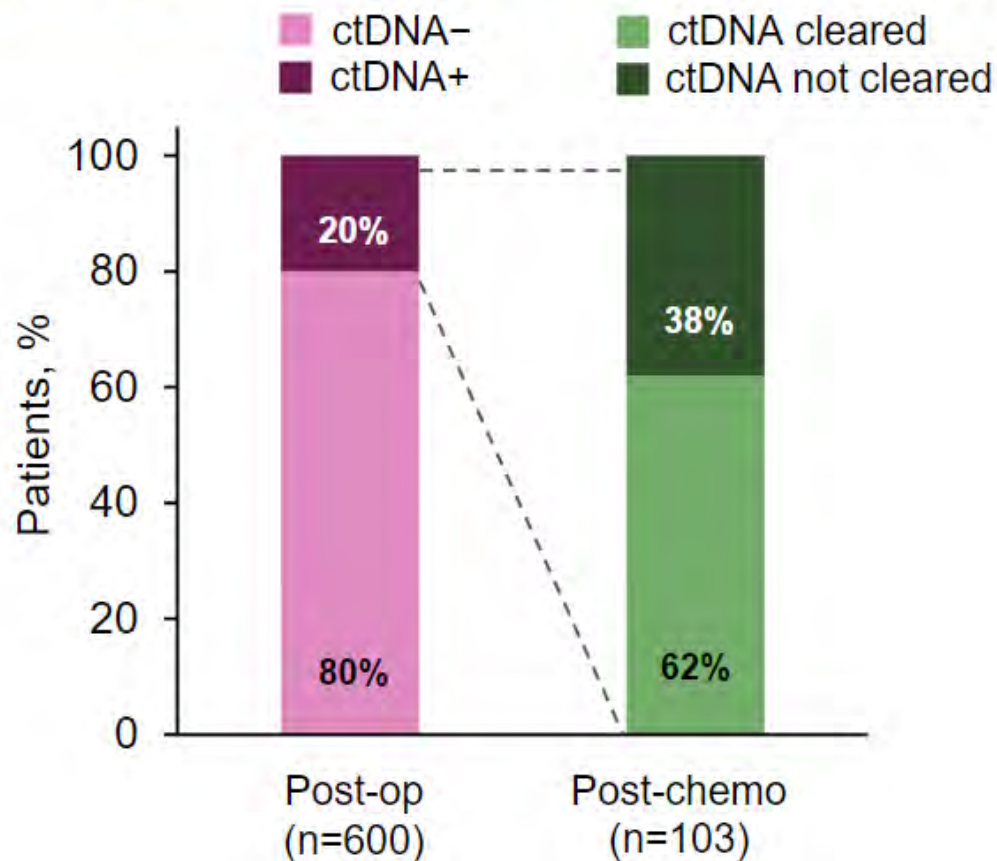
Abbosh C et al. *Nature*. 2017
Chaudhuri A et al. *Cancer Discov*. 2017



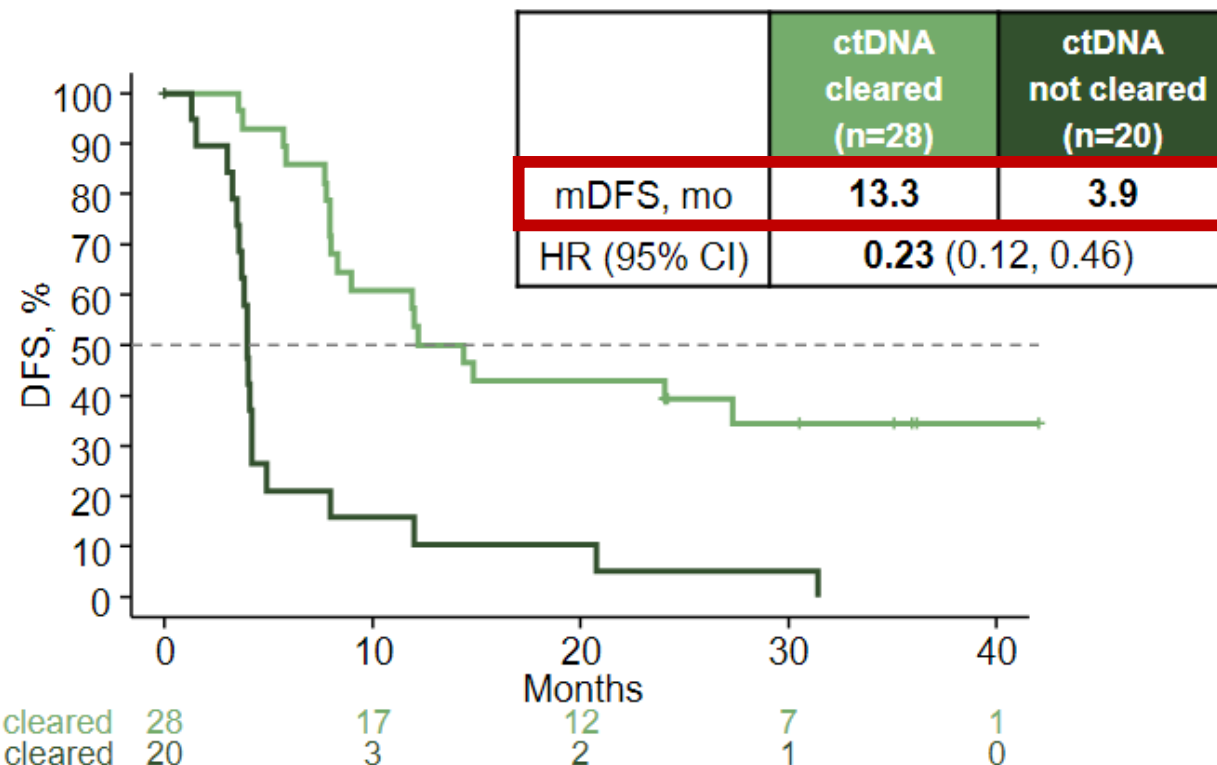
IMpower-010: patients with detectable ctDNA MRD after adjuvant chemotherapy have worse prognosis



Impact of chemo on ctDNA clearance status

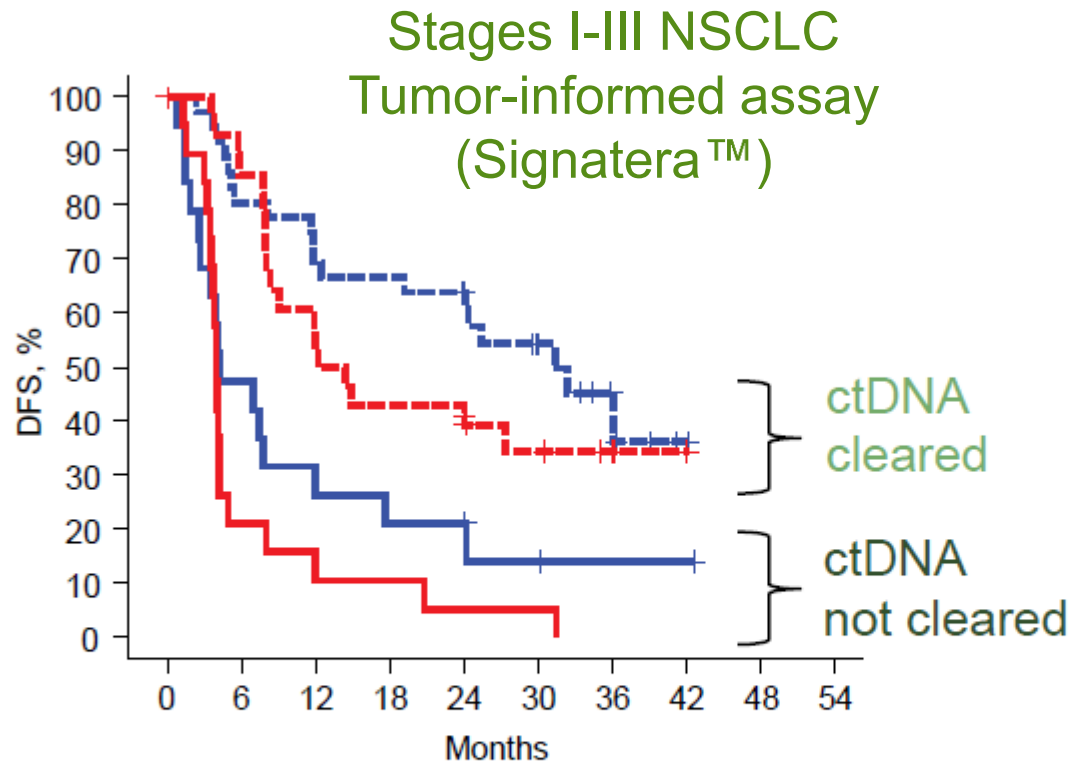


DFS by ctDNA clearance status in the BSC arm





IMpower-010: data suggests adjuvant atezolizumab delays conversion to ctDNA +



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

ctDNA status
post- adjuvant
chemotherapy

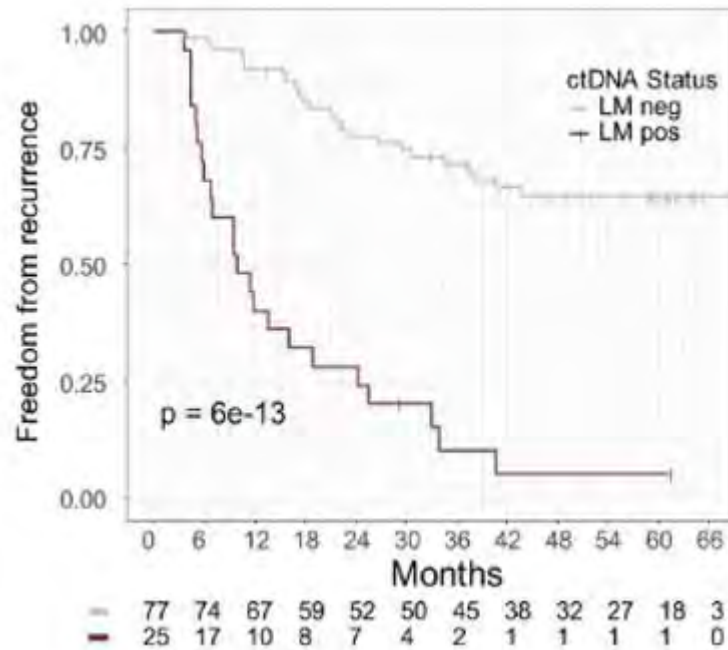
ctDNA cleared	Atezo (n=36)	BSC (n=28)
mDFS, mo	31.3	13.3
HR (95% CI)	0.7 (0.37, 1.34)	

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

Presented by Enriquera Felip. ESMO Immuno-Oncology Congress 2022, Abstract 10 (<https://bit.ly/3sZVgye>)



Limited clinical sensitivity of current ctDNA assays to detect disease recurrence remains a challenge for clinical practice implementation



TRACERx
LOD= 0.008%
(0.05%)

Stages I-III NSCLC
Invitae PCM™
(Tumor-informed assay)

- Stage I – 50%, stage II – 31%, stage III – 19%
- 25% pts had + ctDNA on post-operative samples
- 51/108 patients (47%) developed disease recurrence
- Clinical Sensitivity = 49% and Clinical Specificity = 96% to detect disease recurrence

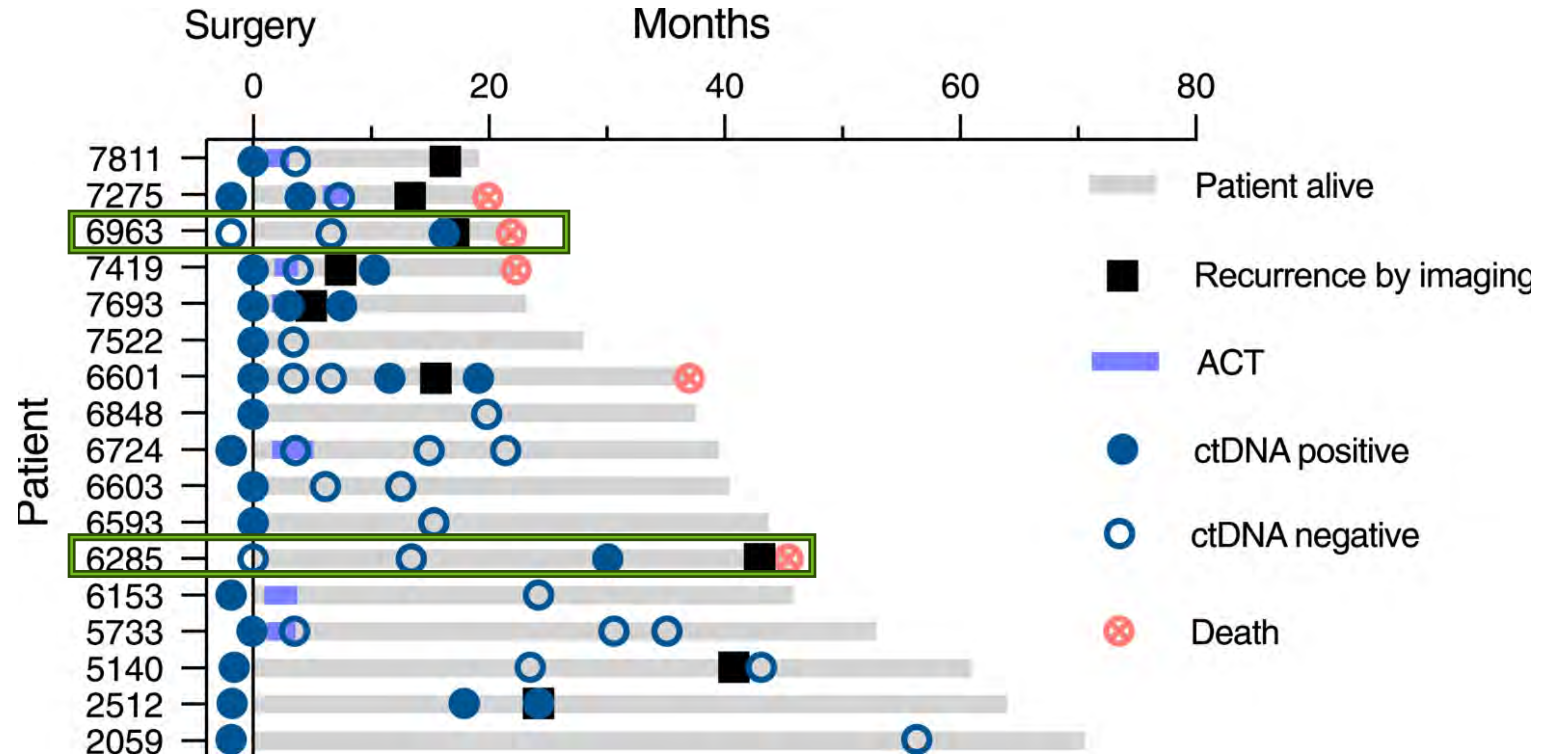


Data suggests one ctDNA MRD timepoint may not be enough to achieve meaningful clinical sensitivity to detect disease recurrence

15 patients developed disease recurrence

Patient	ctDNA positive pre-surgery	ctDNA positive post-surgery	Site(s) of recurrence
5597	No	No	Lung
6285	No	Yes	Lung, nodal
7811	Yes	No	Brain, lung
5944	No	No	Lung, nodal
7246	No	No	Brain
7487	No	No	Lung
7275	Yes	Yes	Bone, liver, pleura
7693	Yes	Yes	Brain, lung, nodal
6963	No	Yes	Liver, lung
5140	Yes	No	Lung, nodal
7419	Yes	Yes	Lung, nodal
6601	Yes	Yes	Lung
5470	No	No	Lung
2512	Yes	Yes	Lung, nodal
2493	No	No	Lung

Stages I-III NSCLC
Tumor-informed assay
(Signatera™)



Clinical sensitivity with one MRD timepoint=
46%
(post-surgery)

The sensitivity to detect disease recurrence is heterogenous across different platforms and may be too low 

Author & Year	No. ^a	Clinical Stage	Treatment	ctDNA Assay	Sensitivity (%) ^b	Specificity (%) ^b
Chaudhuri et al (2017) ^c	32	IB-III B	CRT or RT and/or surgery +/- chemo	CAPP-Seq	94	100
Abbosh et al (2017) ^d	24	IA-III B	Surgery +/- chemo +/- PORT	Signatera	36	90
Chen et al (2019) ^e	25	IIB-III B	Surgery +/- chemo	cSMART	44	88
Zviran et al (2020) ^f	22	IA-III	Surgery +/- chemo and RT	MRDetect	100	71

Pellini B & Chaudhuri A. *J Clin Oncol*. 2022



Interrogating ctDNA in multiple timepoints after curative-intent treatment improves the sensitivity to detect disease recurrence

Author & Year	No. ^a	Clinical Stage	Treatment	ctDNA Assay	Sensitivity (%) ^b	Specificity (%) ^b
Chaudhuri et al (2017) ^c	37	IB-IIIB	CRT or RT and/or surgery +/- chemo	CAPP-Seq	100	100
Abbosh et al (2017) ^d	24	IA-IIIB	Surgery +/- chemo	Signatera	93	70
Abbosh et al (2020) ^e	78	I-III	Surgery +/- chemo	ArcherDx	82	96

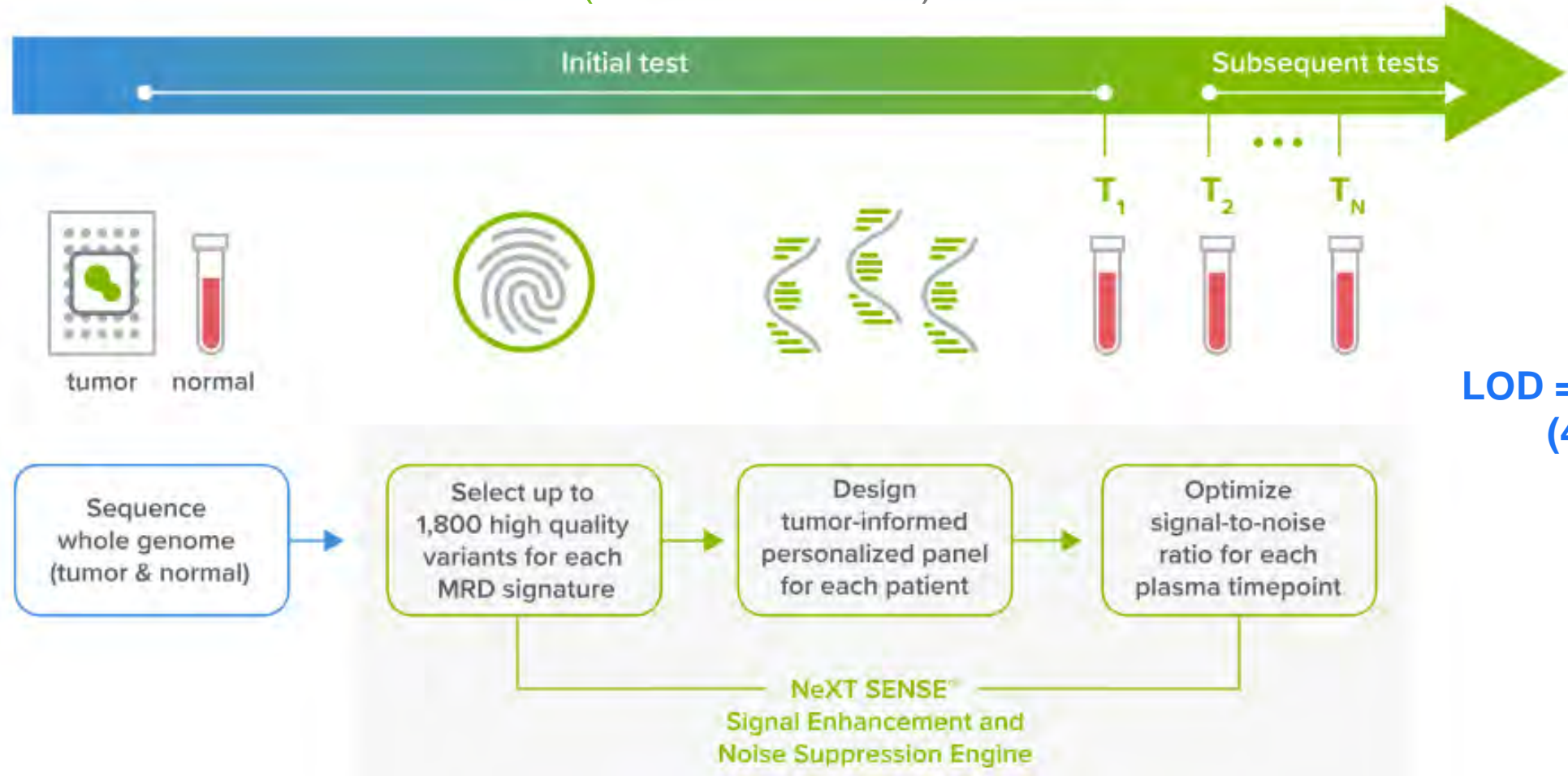
ctDNA post-treatment surveillance studies in NSCLC

Pellini B & Chaudhuri A. *J Clin Oncol*. 2022

Will WGS solve the issue of clinical sensitivity?

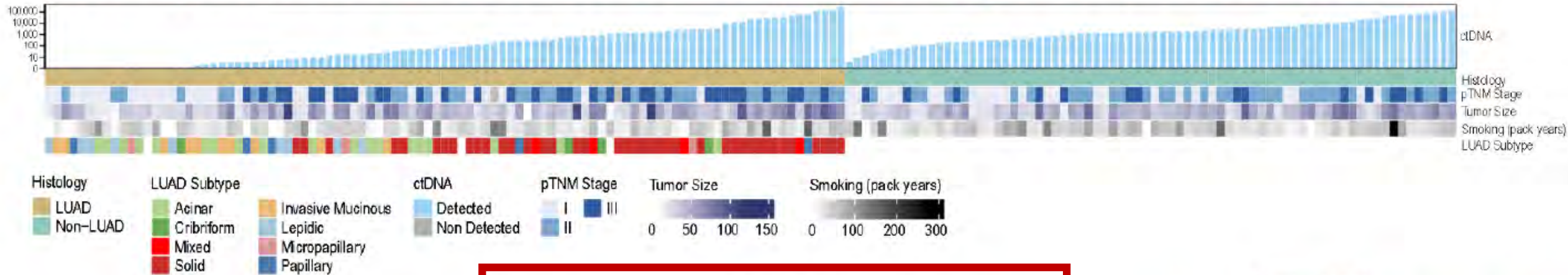


Stages I-III NSCLC
Tumor-informed assay
(NeXT Personal®)

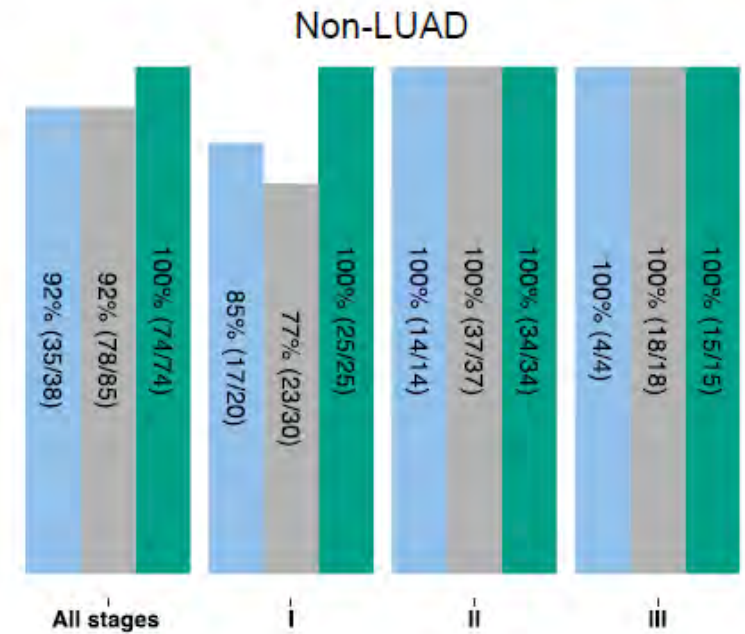
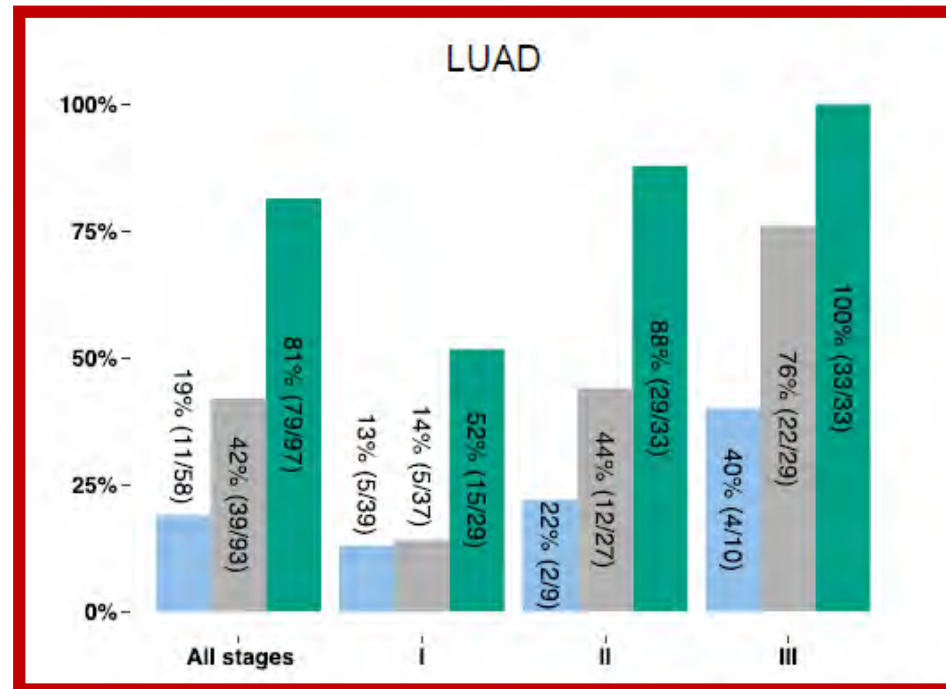


Source <https://www.personalis.com/products/next-personal/>

Pre-operative detection of ctDNA using NeXT Personal



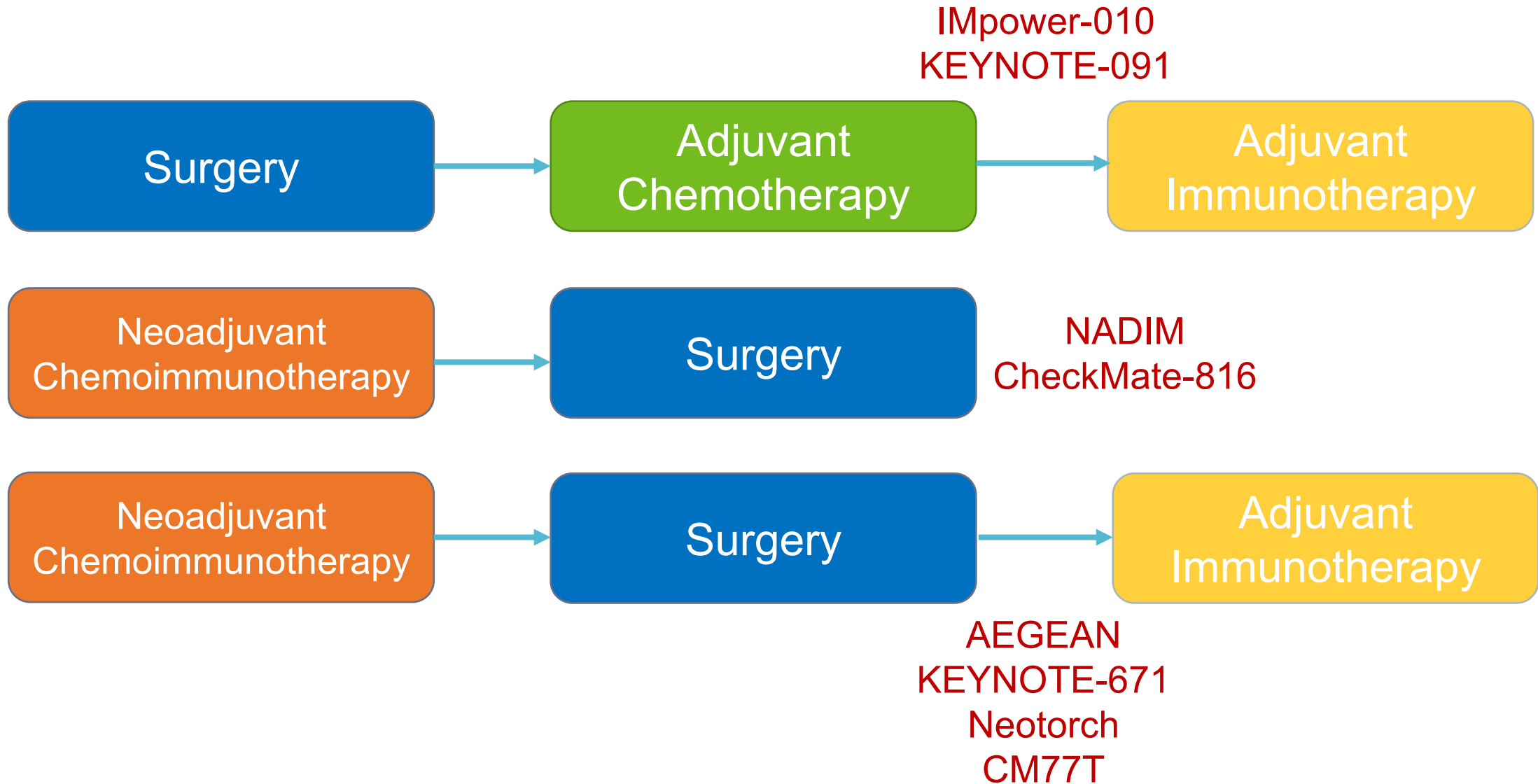
- ctDNA detected from 1.7 to 253,826 PPM
- Pre-operative ctDNA detected in **81%** of LUAD and **100%** of non-LUAD patients
- Includes **52%** of pTNM stage I LUAD patients



Abbosh, 2017 Abbosh, 2023 Current study

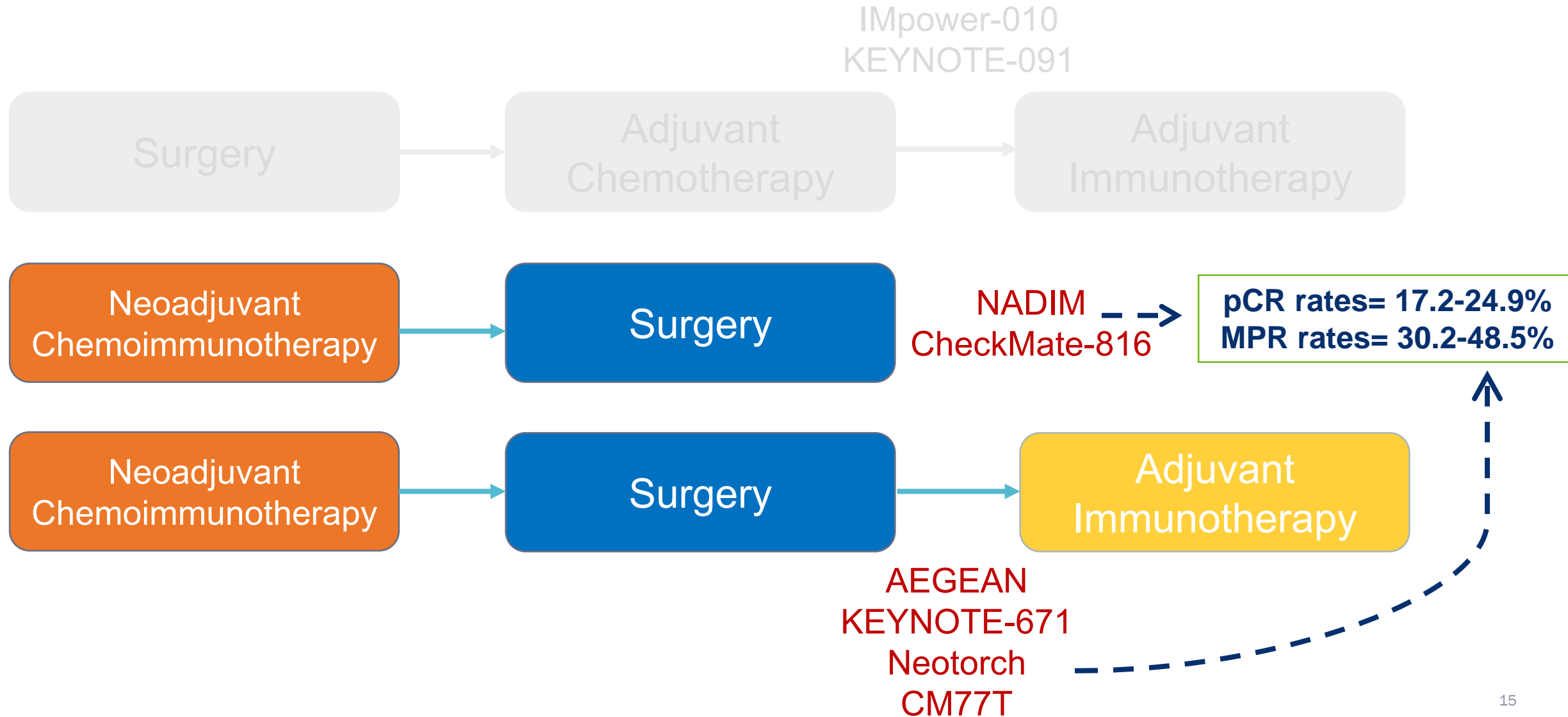


Current treatment landscape for resectable NSCLC





Current treatment landscape for resectable NSCLC



Unanswered questions



- Tumor-informed vs. tumor-naïve assays– which one is better?
- How many timepoints should we use to guide treatment escalation or de-escalation in trials and clinical practice?
- Will treatment de-escalation based on MRD status be equal to SOC?
- Will treatment escalation based MRD status improve DFS and OS?

How do I treat ctDNA positive after a complete resection?



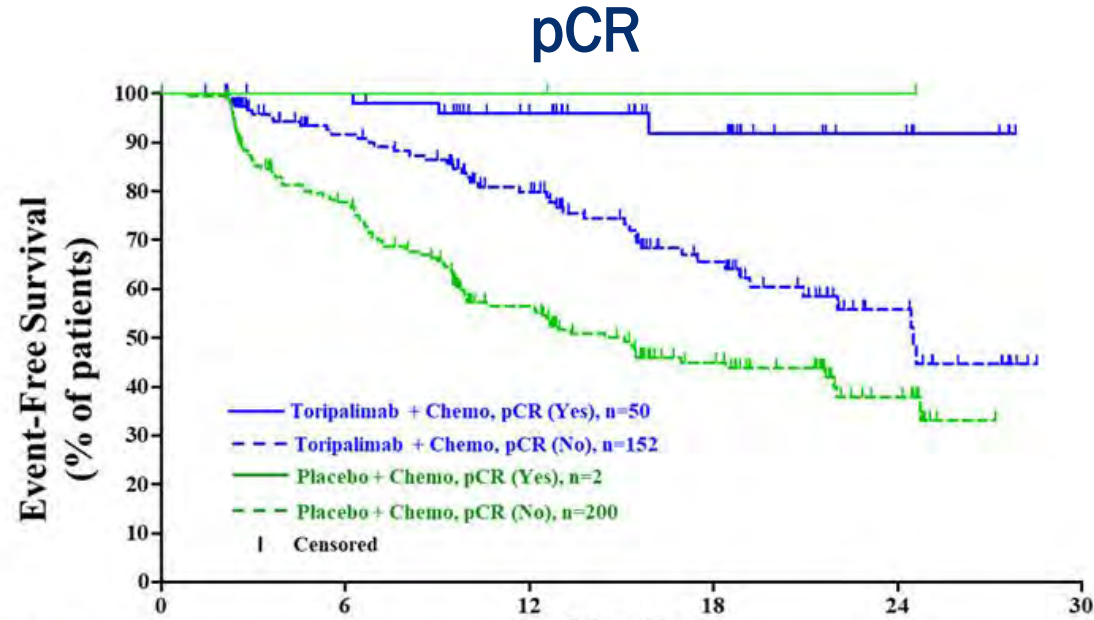
I do not order ctDNA in the MRD setting given assay limitations and lack of proven clinical utility

If you choose to order ctDNA in the post-op setting, what would I recommend?

- Obtain a **pre-neoadjuvant or pre-surgical** blood specimen for **ctDNA** analysis → **ctDNA dynamics matter** for risk stratification
- Assess **ctDNA after surgery** if you have given neoadjuvant therapy or **after chemotherapy** if you have **opted for the surgery first approach** → your **MRD timepoint** should be **prior to adjuvant immunotherapy** selection
- If clinically feasible, obtain **blood samples in 2 MRD timepoints** → **2 weeks** post-op & **4 weeks** post-op
- Analyze your **ctDNA MRD** results **together** with your **pathological response** findings

You need to analyze **ctDNA** in at least **3 timepoints** to increase the odds of obtaining clinically meaningful information

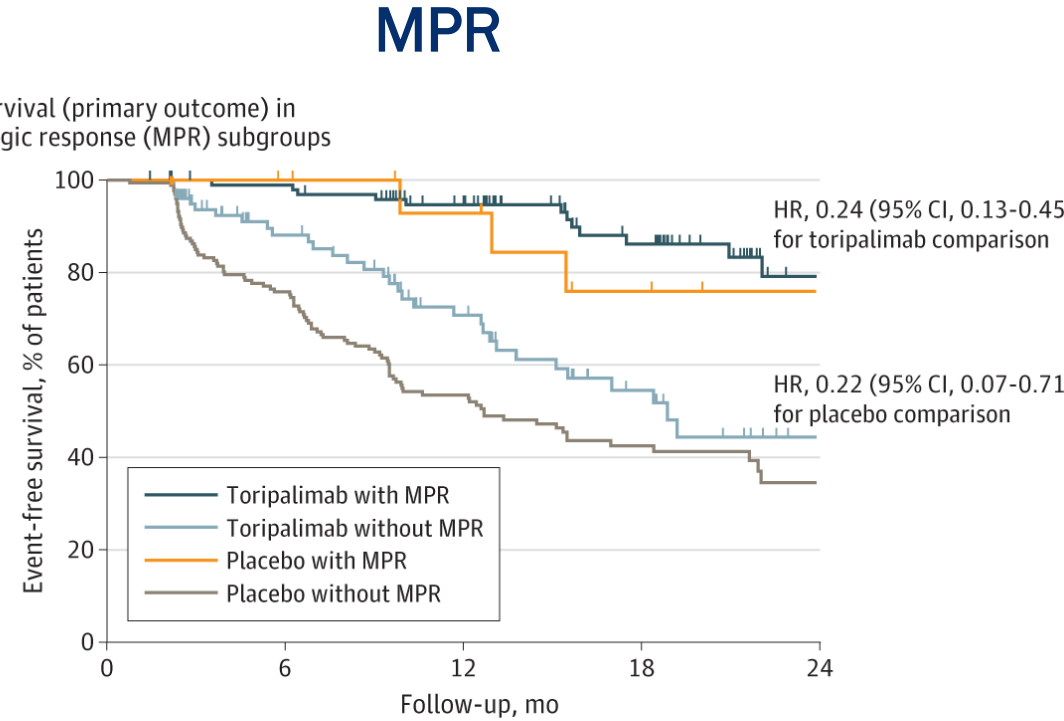
Neotorch: EFS based on pCR and MPR



No. at Risk		Months					
		0	6	12	18	24	30
Toripalimab + chemo, pCR	Yes	50	49	36	21	7	0
	No	152	107	80	45	16	0
Placebo + chemo, pCR	Yes	2	2	2	1	1	0
	No	200	137	84	42	14	0

Toripalimab arm		Placebo arm	
<input type="checkbox"/>	Median EFS NE versus 24.5 months	<input type="checkbox"/>	Median EFS NE versus 15.1 months
<input type="checkbox"/>	HR= 0.16 (95%CI 0.05-0.51)	<input type="checkbox"/>	HR= NE

B Event-free survival (primary outcome) in major pathologic response (MPR) subgroups

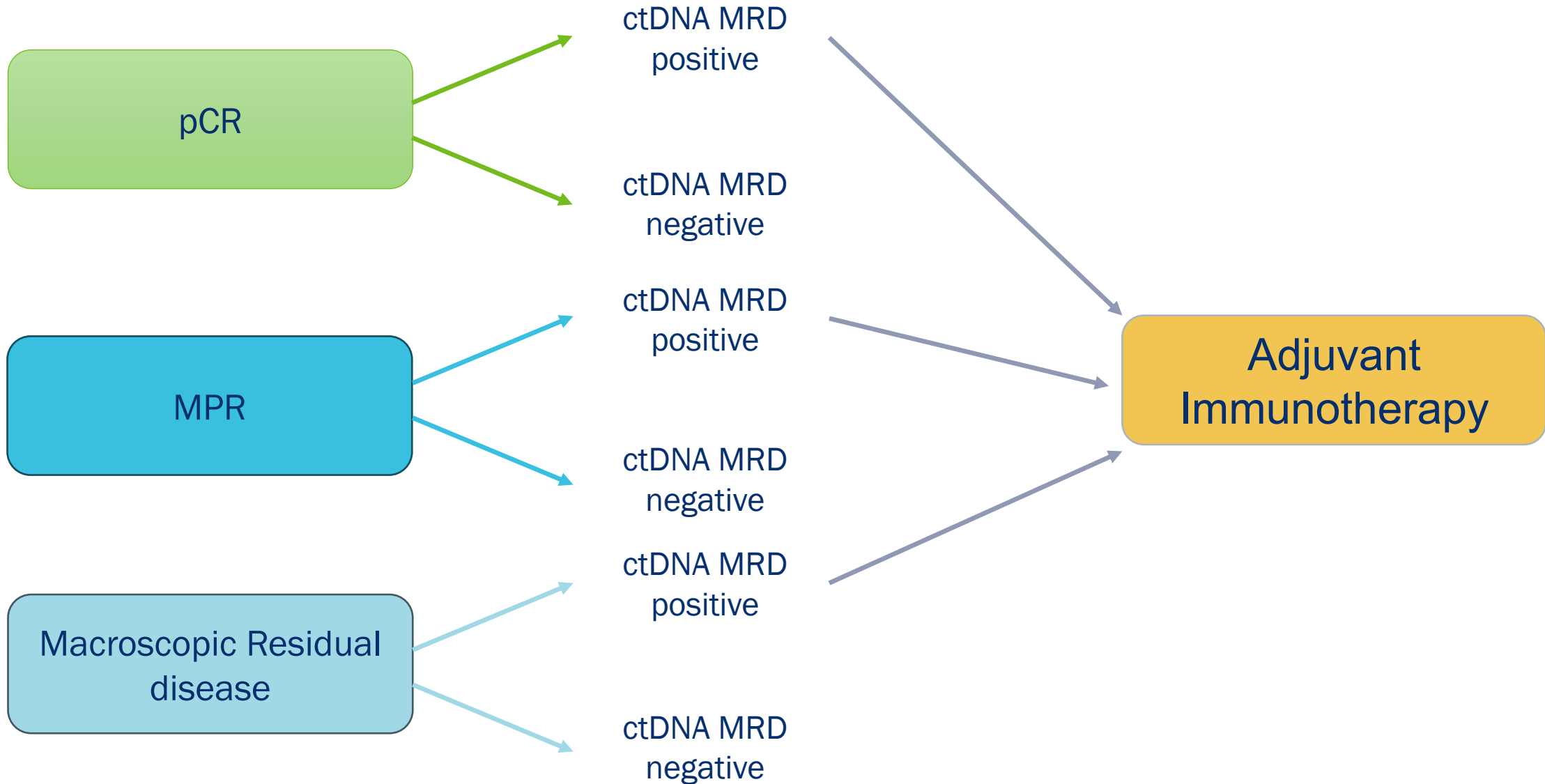


No. at risk		Follow-up, mo				
		0	6	12	18	24
Toripalimab with MPR		98	95	77	46	17
Toripalimab without MPR		104	61	39	20	6
Placebo with MPR		17	16	13	7	5
Placebo without MPR		185	123	73	36	10

How do I treat ctDNA positive after a complete resection?



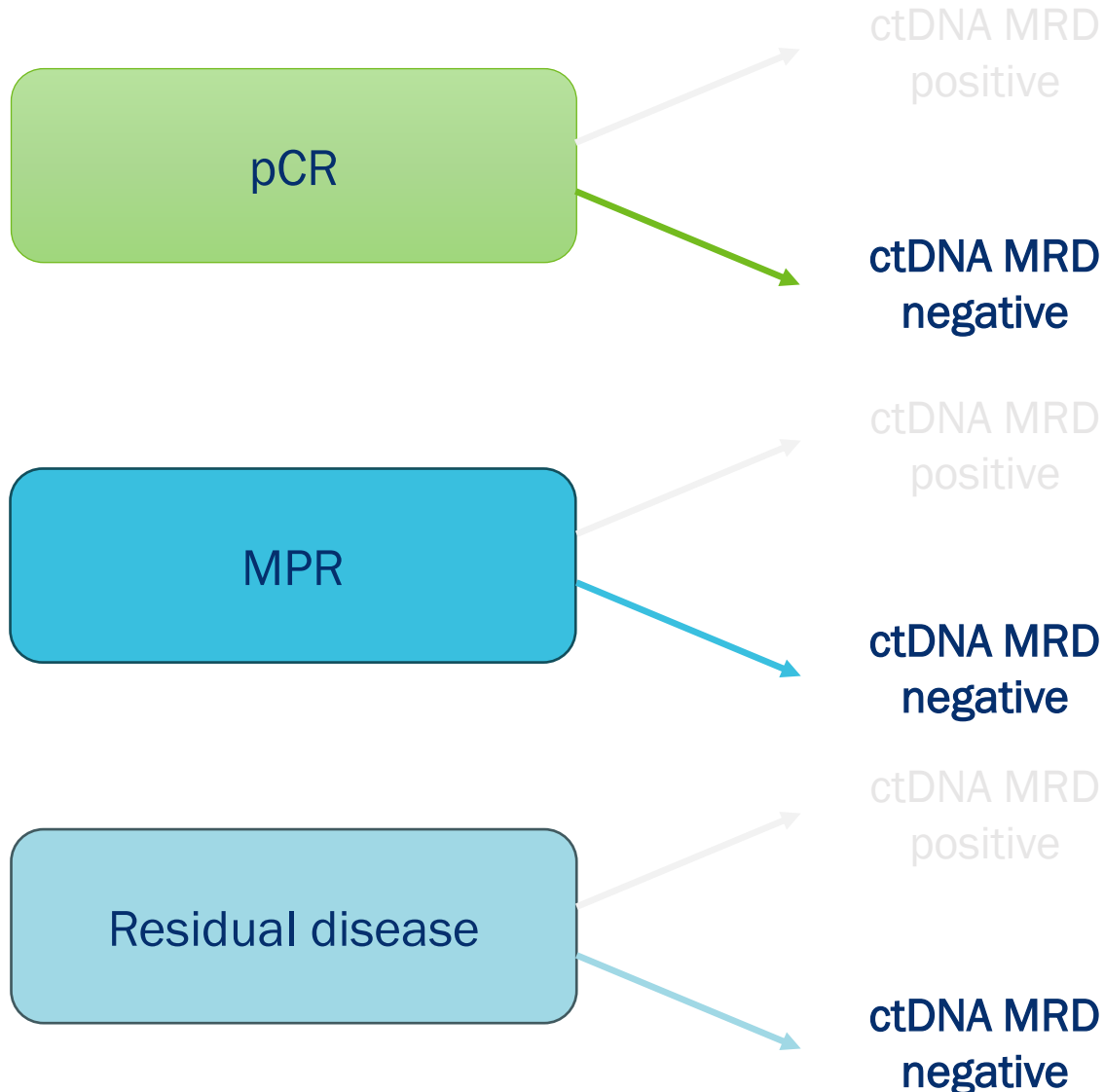
My recommended approach as of 01/2025



How do I treat ctDNA positive after a complete resection?



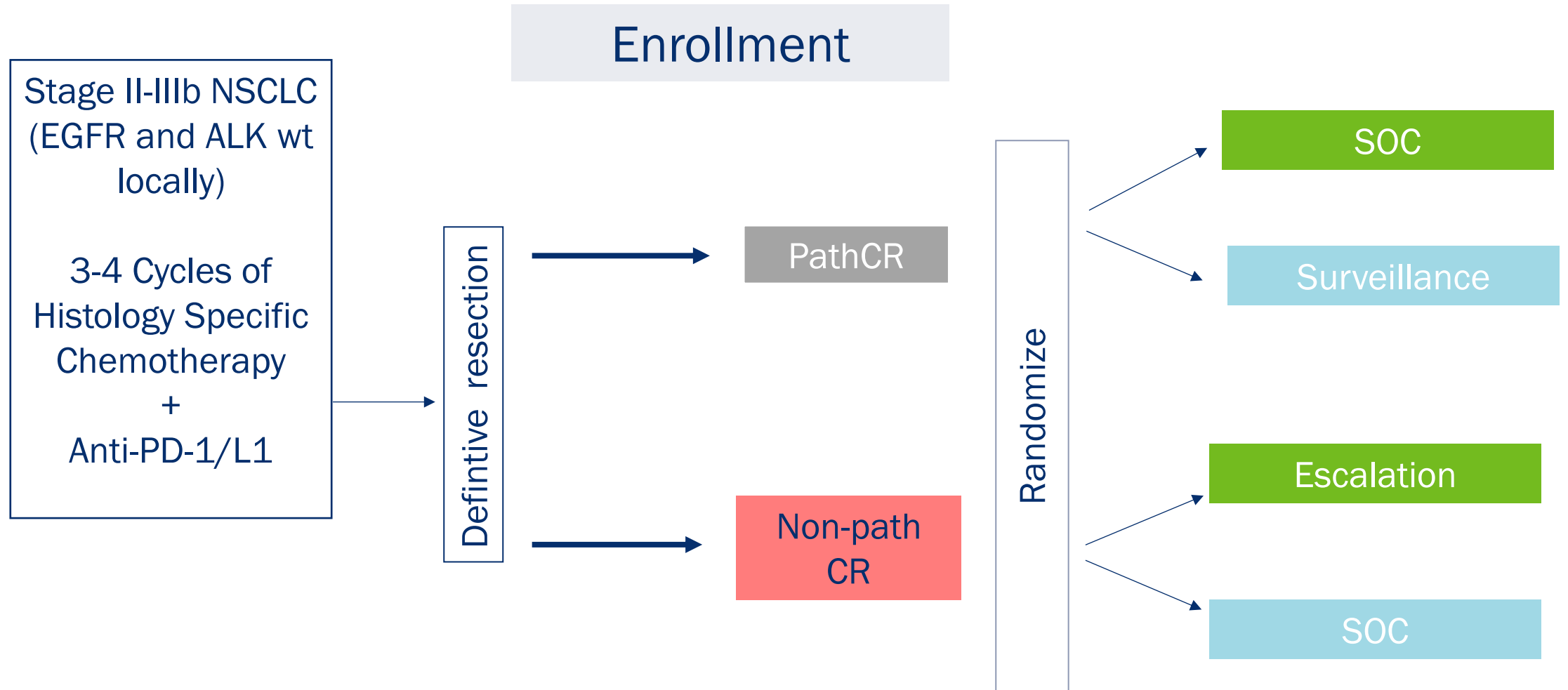
My recommended approach as of 01/2025



It depends...

- If **ctDNA undetectable prior to curative-intent treatment** start in the setting of **pCR and MPR and ctDNA MRD (-)**, pts likely do not need additional treatment
- If **ctDNA (+) pre-treatment and cleared (ctDNA MRD -)**, I would recommend **adjuvant immunotherapy**

We need clinical trials that will investigate escalation and de-escalation based on MRD status to then change our clinical practice





Take home points

- ctDNA can detect **MRD** and it is a strong **prognostic biomarker**
- **One timepoint MRD status may not be enough** to inform clinical-decision making
- **Tumor-informed MRD assays will not be feasible** for patients with **pCR** and **MPR**
- Ongoing trials will inform **if clinical decision-making can be guided by ctDNA** and if that improves patients' outcomes

Thank you!



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