



CIRCULATING TUMOR DNA TO GUIDE ADJUVANT THERAPY IN COLORECTAL CANCER IS READY FOR PRIME TIME: AGREE

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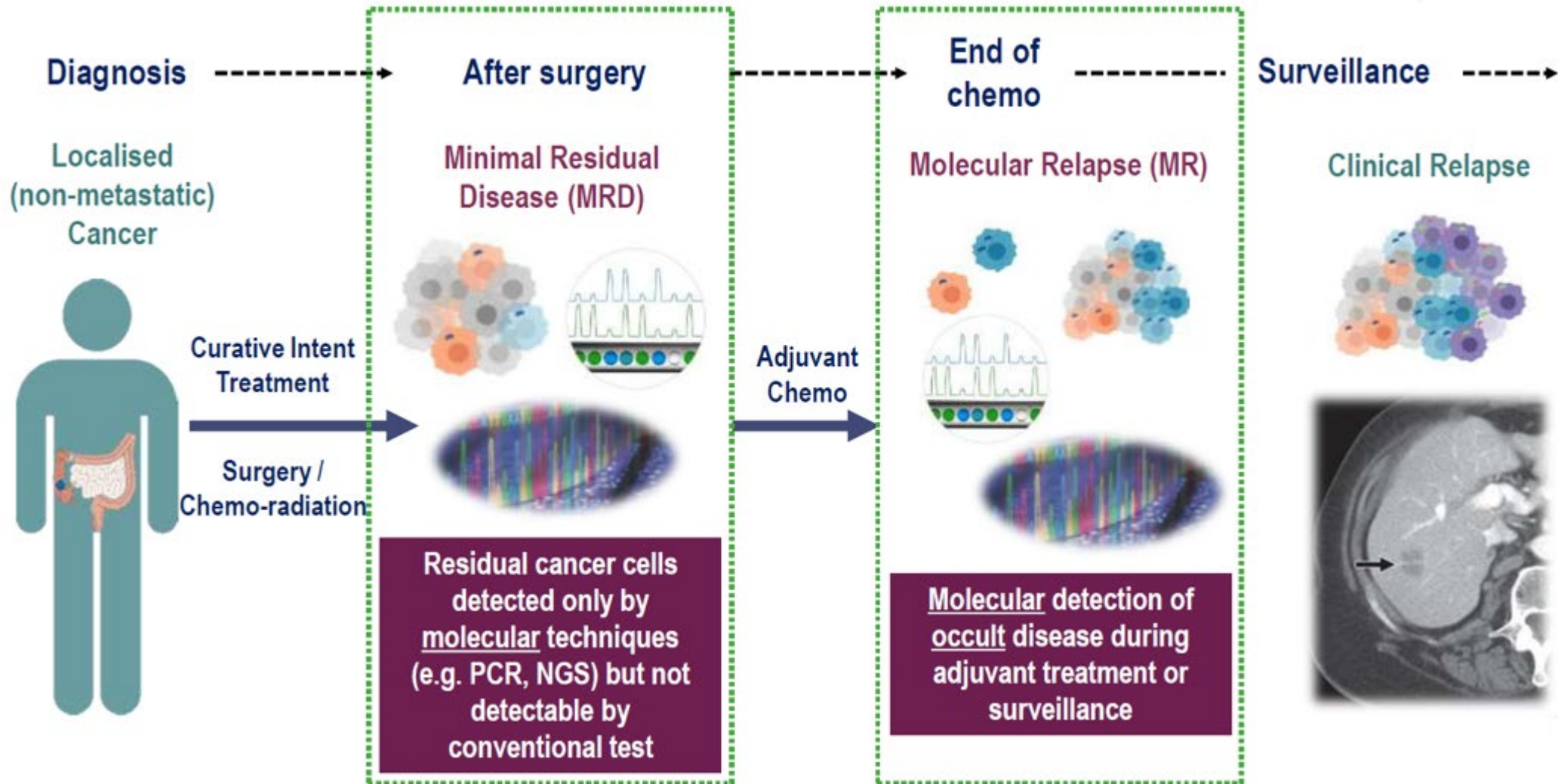
Winship Cancer Institute of Emory University



Disclosures

- Guardant, Novartis.

Minimal Residual Disease/Molecular Relapse- Definitions



Current challenges in adjuvant treatment

- **Stage-based approach is imprecise**

- Undertreating stage II
- Overtreating stage III

- **Lack of better treatments than FOLFOX**

- Irinotecan
- Bevacizumab
- Cetuximab

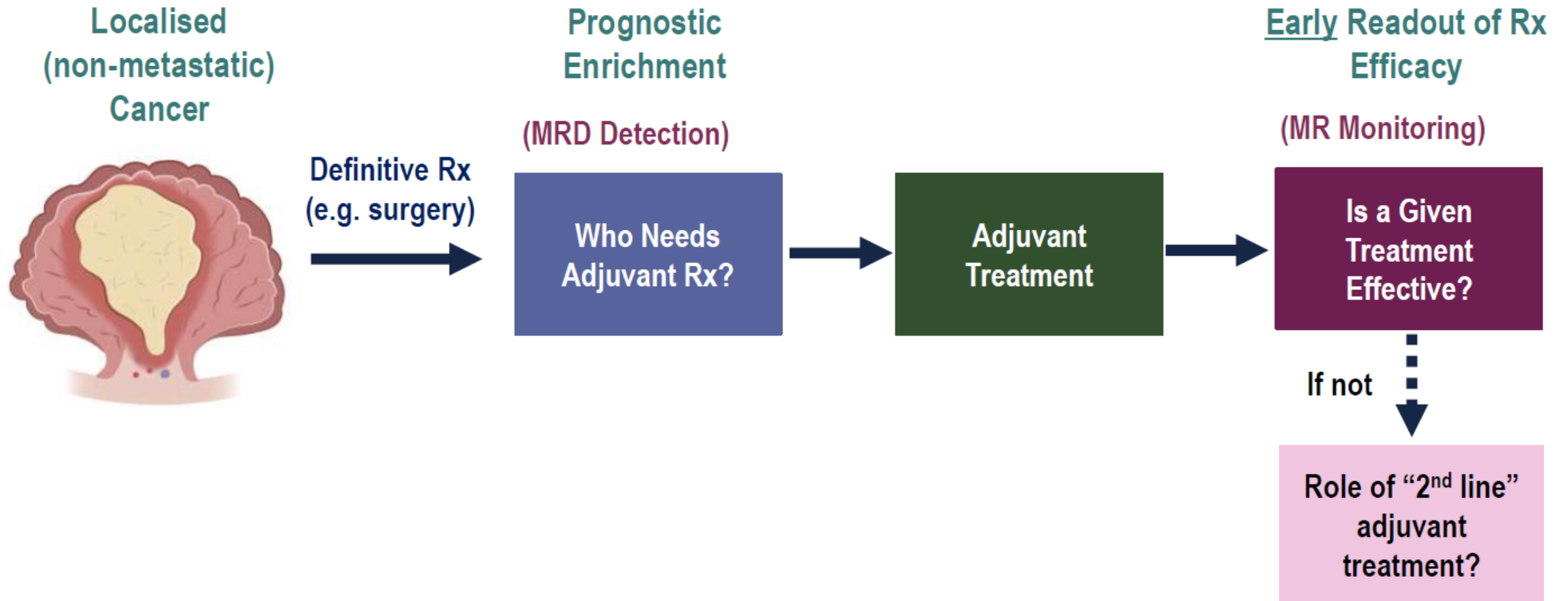


Negative
trials

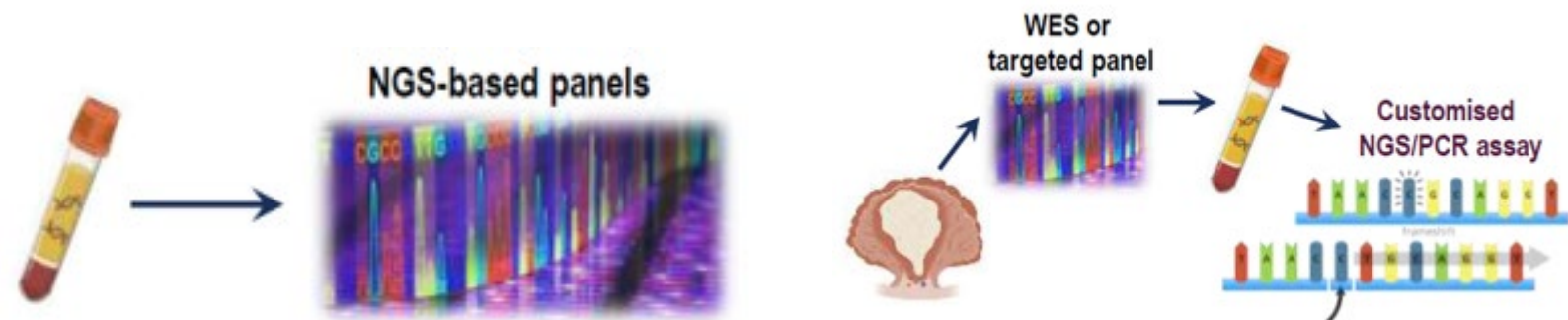
- **Adjuvant trials are inefficient**

- Lack of real-time marker to assess effectiveness of adjuvant therapy
- Requires large sample size, long follow-up
- Current trial infrastructure not suited for rapidly evolving novel therapies

Personalizing Adjuvant Therapy Biomarker

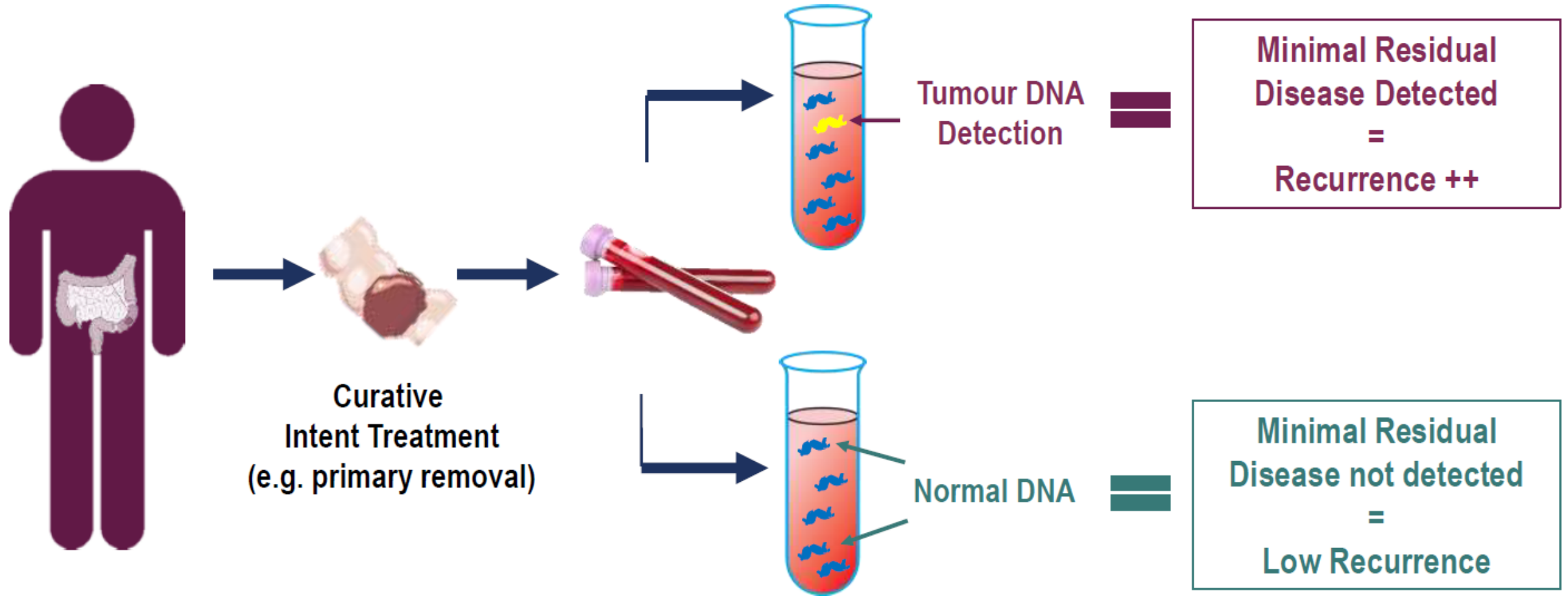


MRD- Two approaches to ctDNA analysis

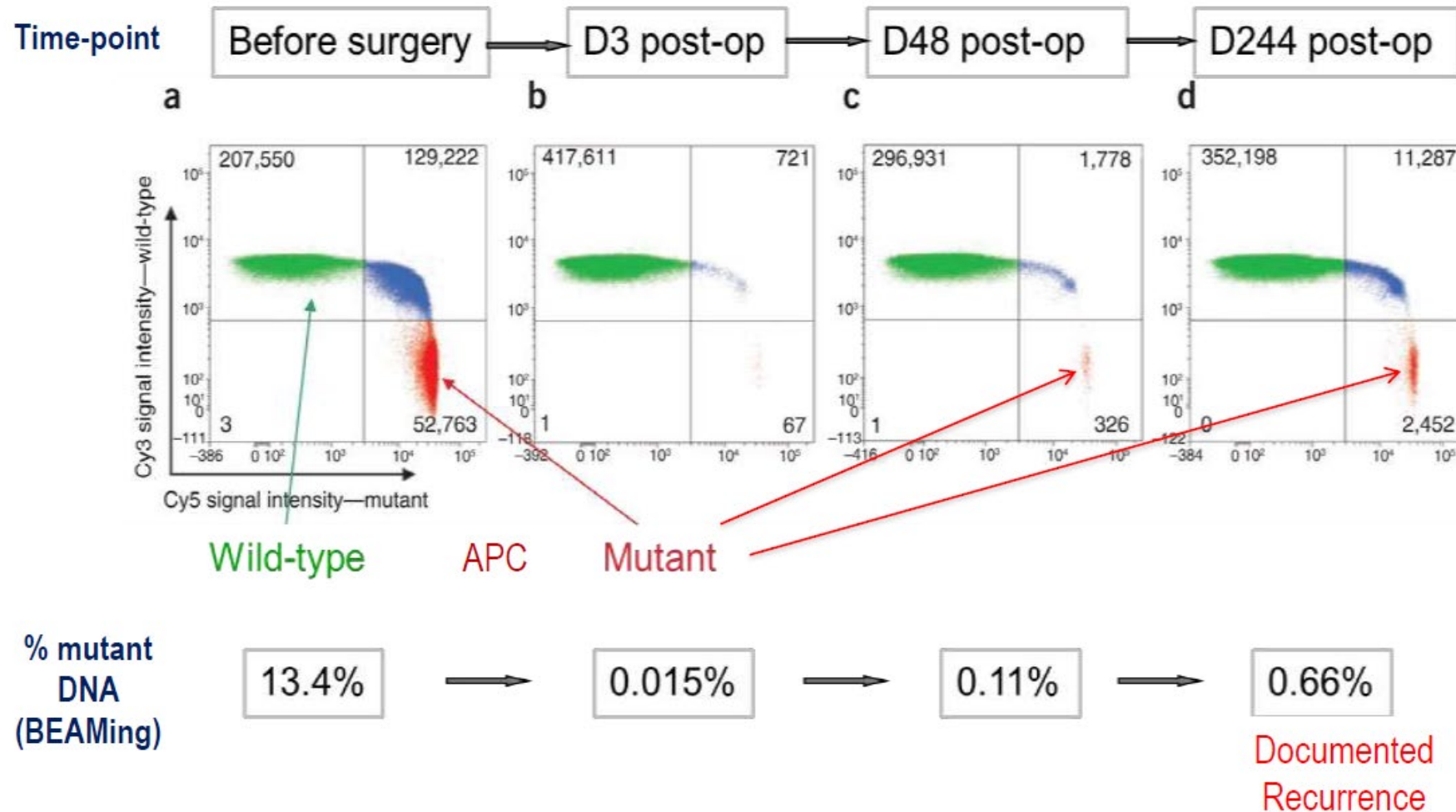


	Tumor-agnostic	Tumor-informed
Method	Detect mutation de novo from plasma (one assay for all)	Identify mutations in tumor tissue->track mutations in plasma (customized)
Key Advantage	Does not require tissue	Higher sensitivity
Key Disadvantage	Lower sensitivity due to multiple hypothesis testing	Requires tissue

ctDNA redefines recurrence risk assessment

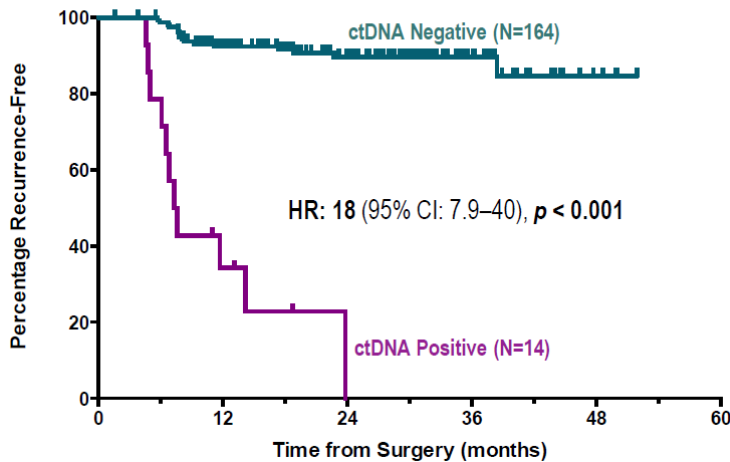


ctDNA reflects tumor burden in CRC

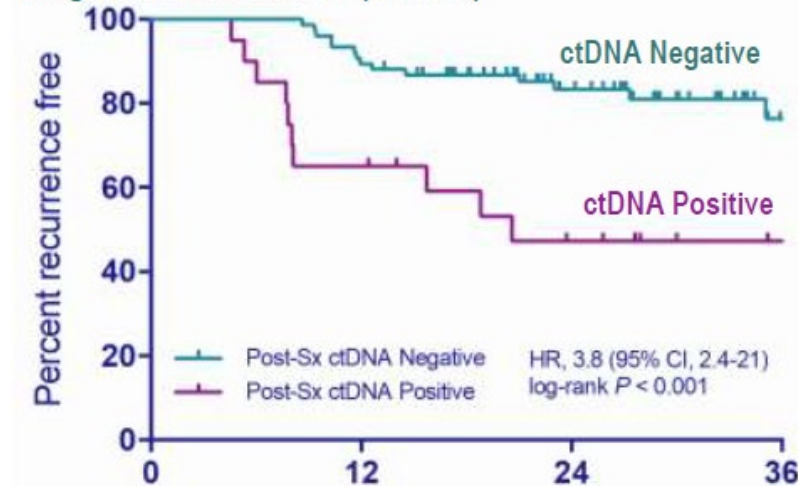


Post-op ctDNA Detection Predicts For Recurrence

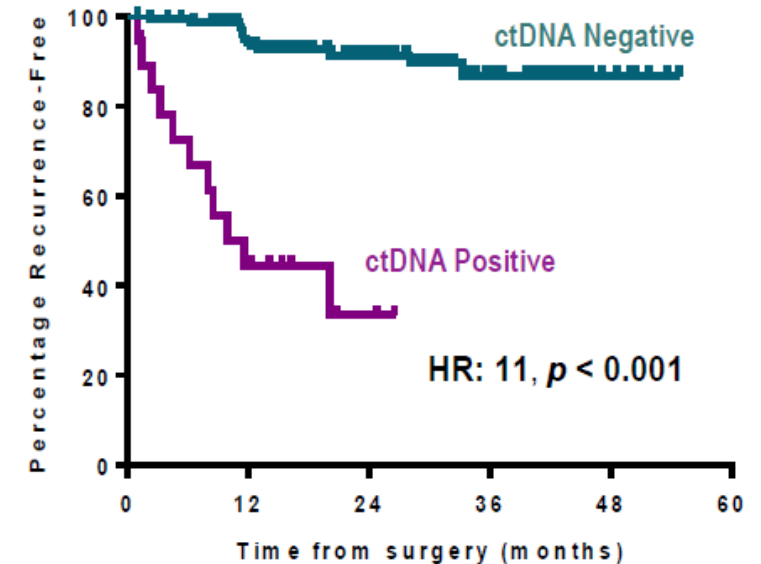
Stage II Colon Cancer (No Chemo)



Stage III Colon Cancer (Chemo)

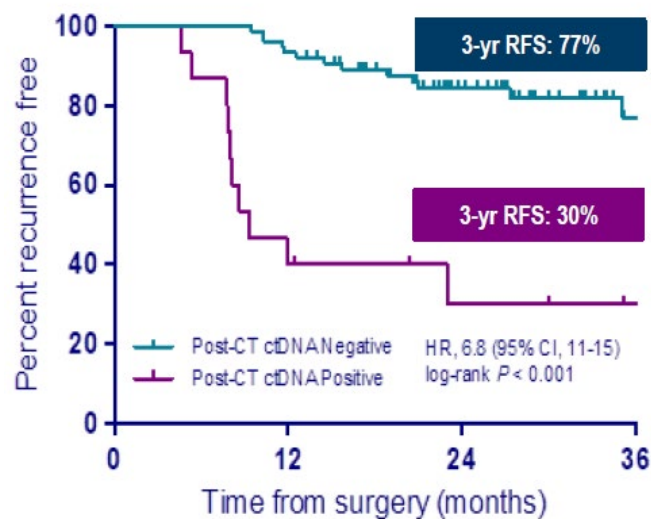


Locally Advanced Rectal Cancer

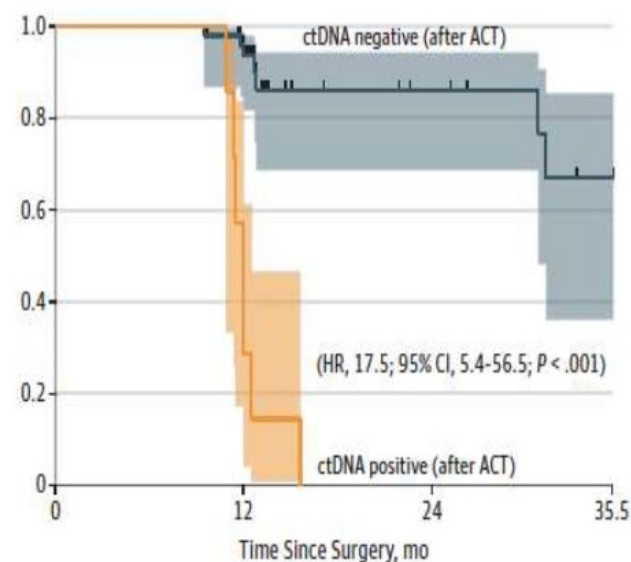


Post-treatment ctDNA is a surrogate for treatment efficacy

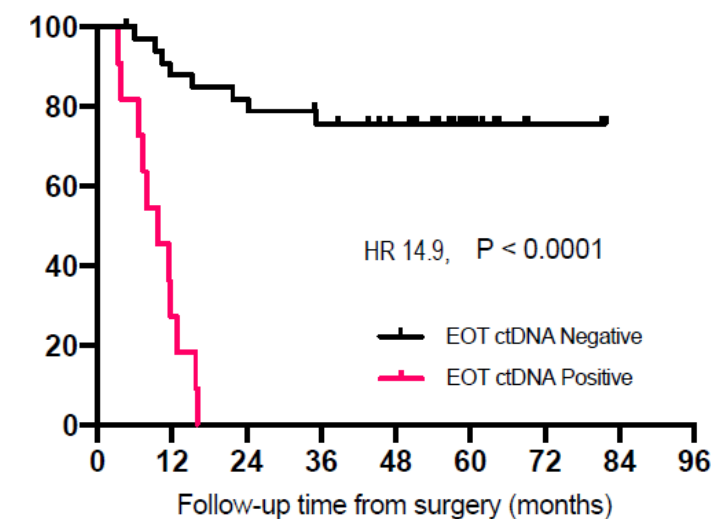
Stage III CC



Stage II-III CC

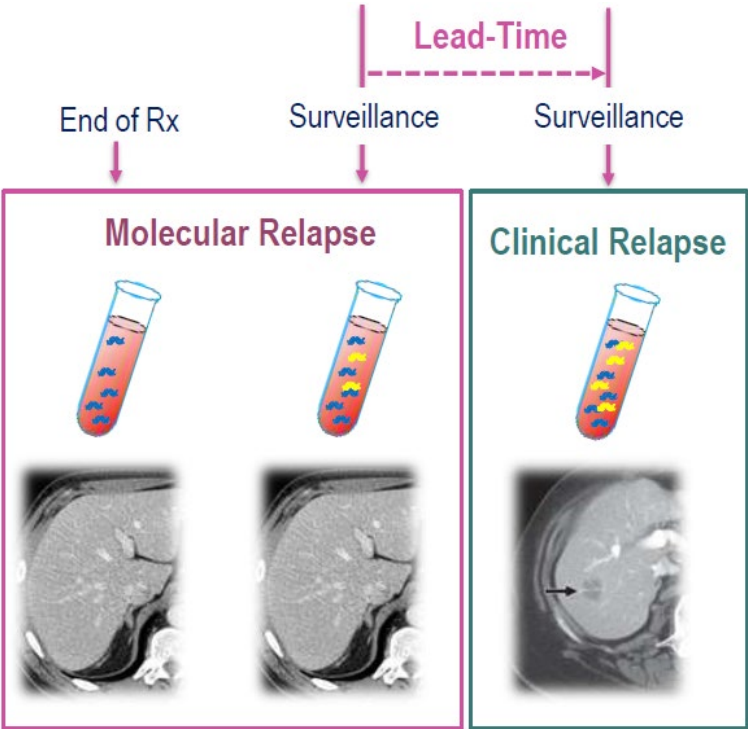


Resected CRC liver mets



ctDNA during surveillance- Lead-time to clinical recurrence

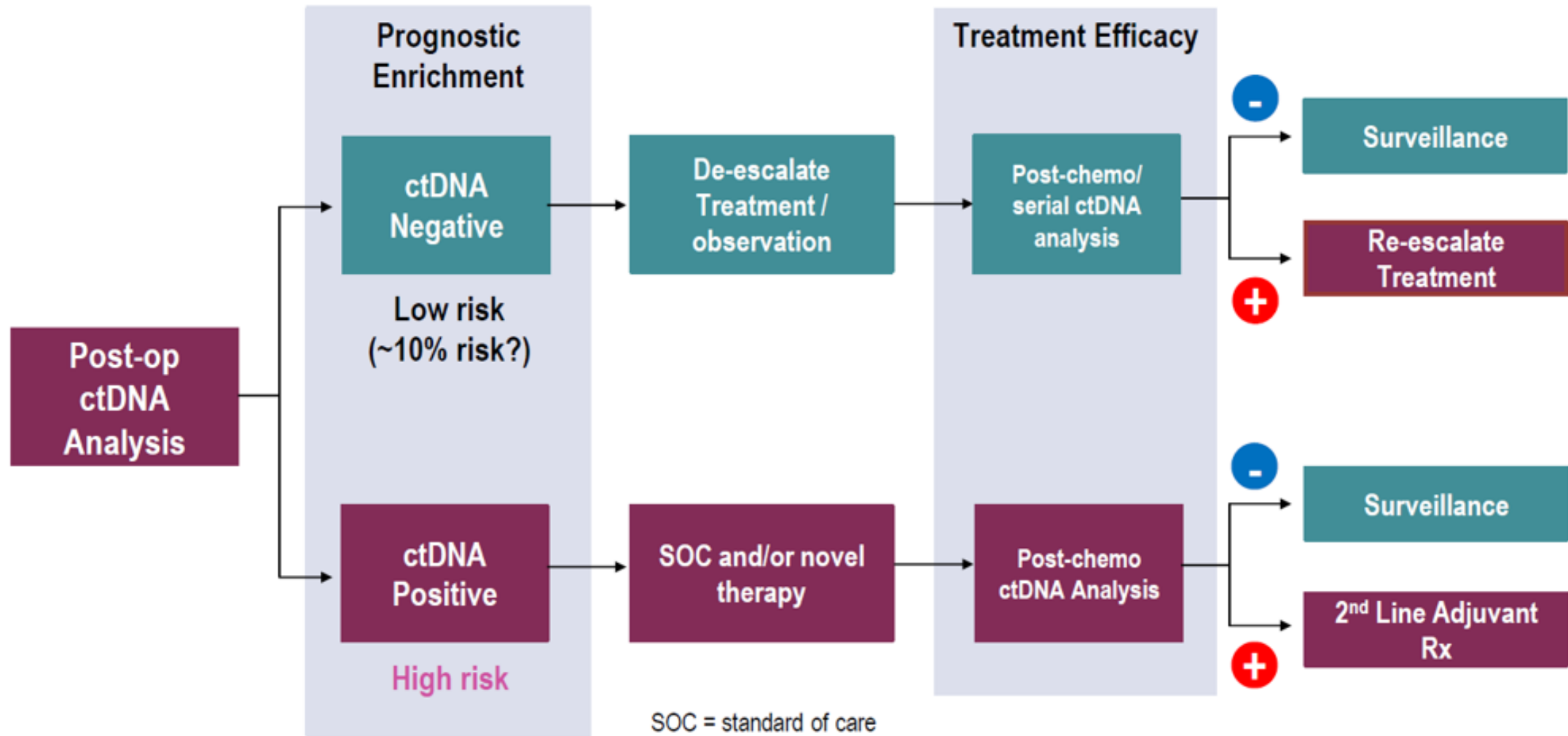
- **Lead-time**= Time between first detection of ctDNA from the end of definitive treatment and clinically detectable recurrence
- **Affected by:**
 - frequency of ctDNA testing and imaging
 - sensitivity of ctDNA assay and imaging modality
 - tumor biology / site of relapse
- **Increasing ctDNA detection rate with increasing time after surgery**



Study	Frequency of Testing		Lead-Time median (range)
	ctDNA	CT scan	
Tie J, <i>et al.</i> (SII)	Q3M	Q6M	5.6M (IQR 2.7-9.3)
Tie J, <i>et al.</i> (SIII)	Q3M	1M + Q12M	1.7M (0.3-15.7)
Tarazona N, <i>et al.</i>	Q4M	Q6M	11.5M (3-18)
Chen G, <i>et al.</i>	Q3M	Q12M	5.1M (not reported)
Wang Y, <i>et al.</i>	Q3-6M	not reported	4M (2-31)
Henriksen TV, <i>et al.</i>	Q3M	12M + 36M	8M (0.56-21.6)

Sci Transl Med 2016;8(346):346ra92; *JAMA Oncology* 2019;5(12):1710-1717; *Ann Oncol* 2019;30(11):1804–1812; *J Hematol Oncol* 2021;14:80; *JAMA Oncol* 2019;5(8):1118-1123; *JCO* 2021;39:3_suppl,11-11.

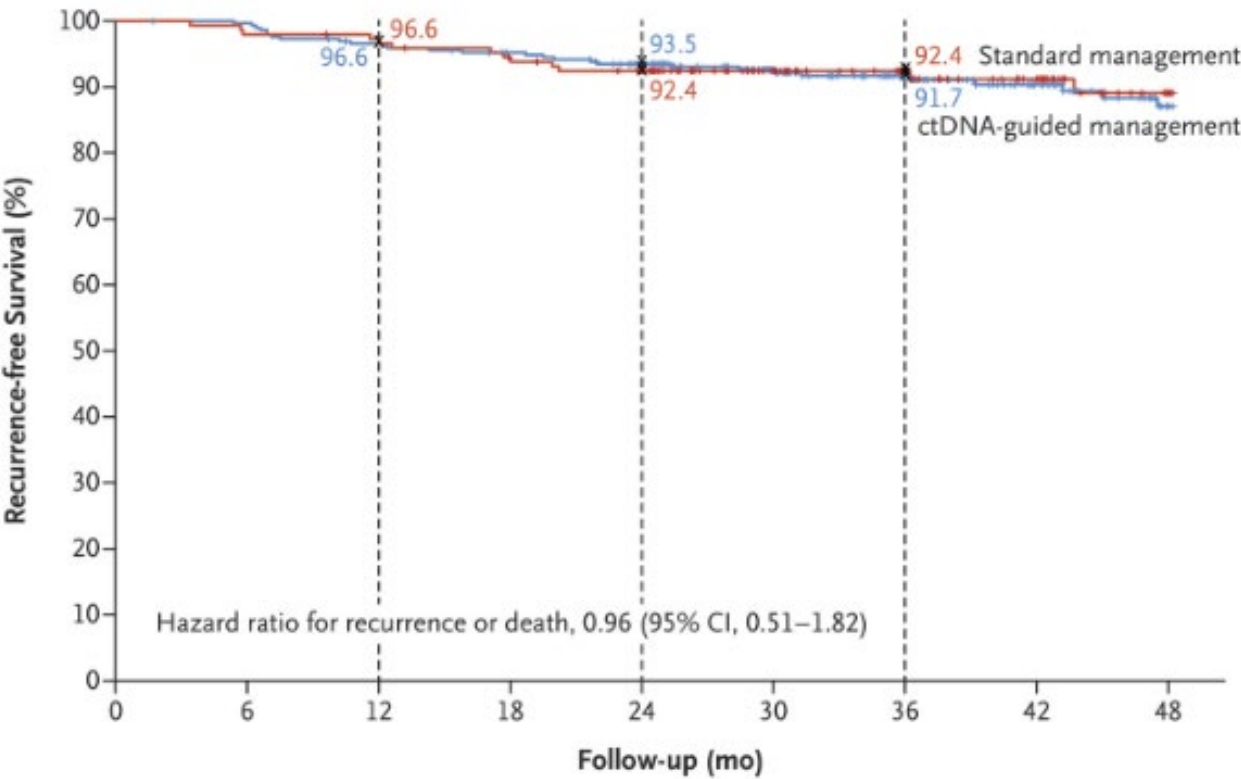
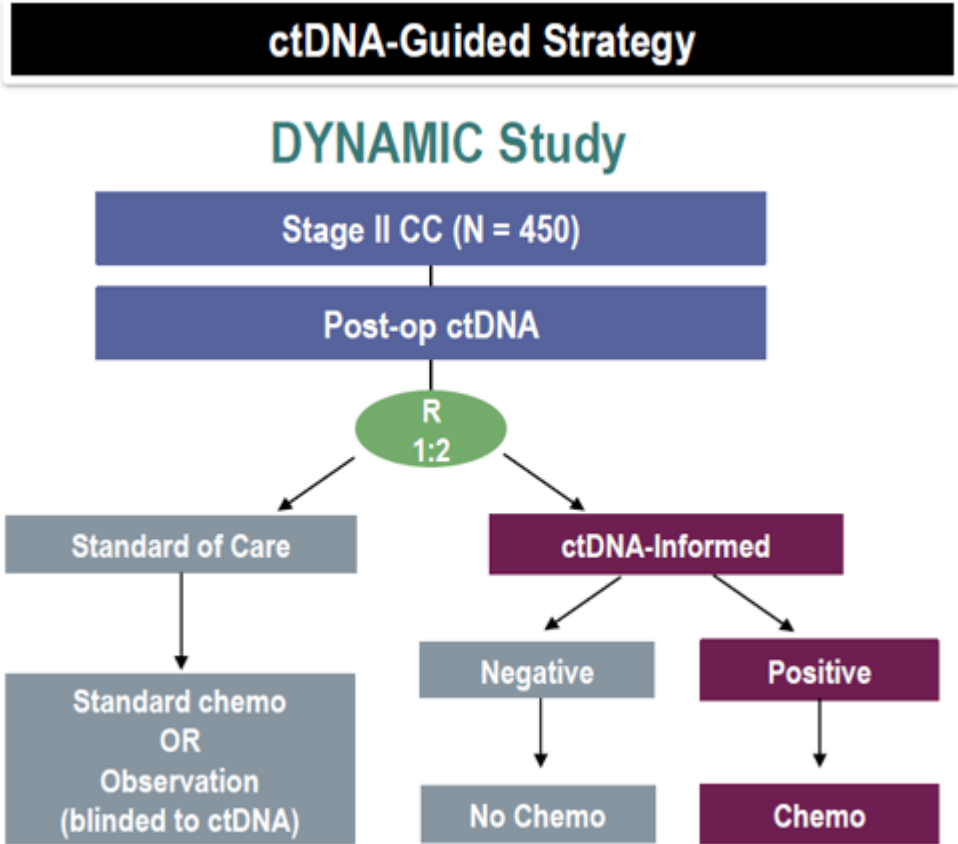
ctDNA-guided approach to adjuvant treatment



ctDNA-informed trials for colon cancer

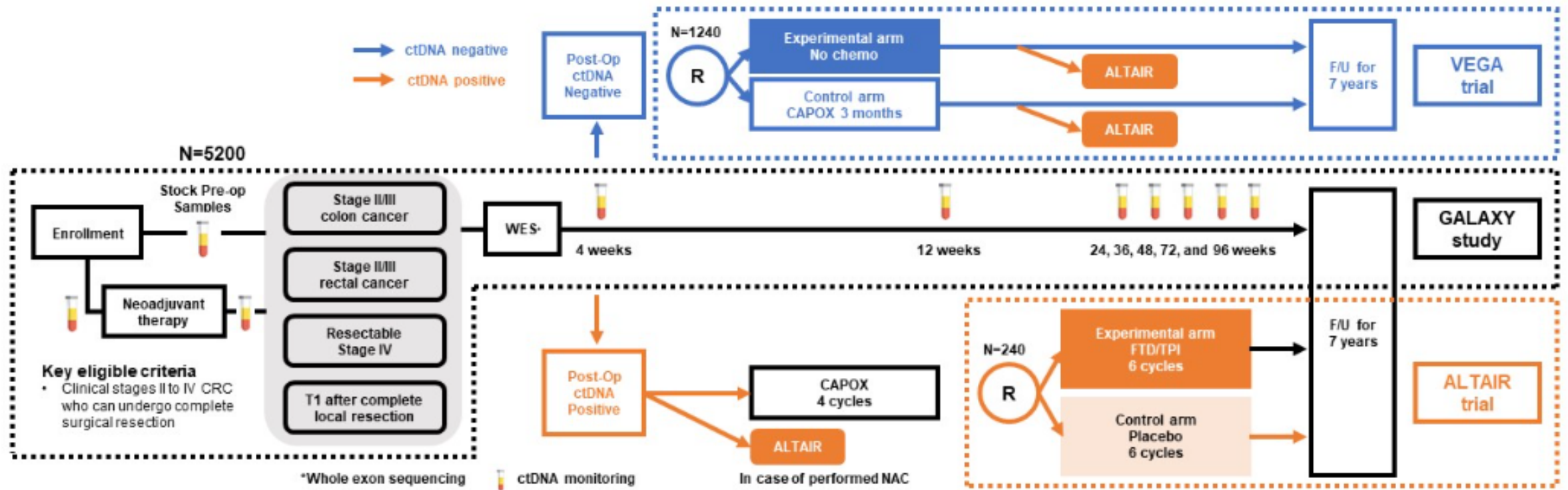
Stage 1	Stage 2	Stage 3	Stage 4
BESPOKE	BESPOKE	BESPOKE	BESPOKE
	COBRA		
	DYNAMIC	DYNAMIC-III	
	CIRCULATE-PRODIGE 70		
	CIRCULATE-JAPAN/US	CIRCULATE-JAPAN/US	CIRCULATE-JAPAN
		ACT3	

DYNAMIC trial



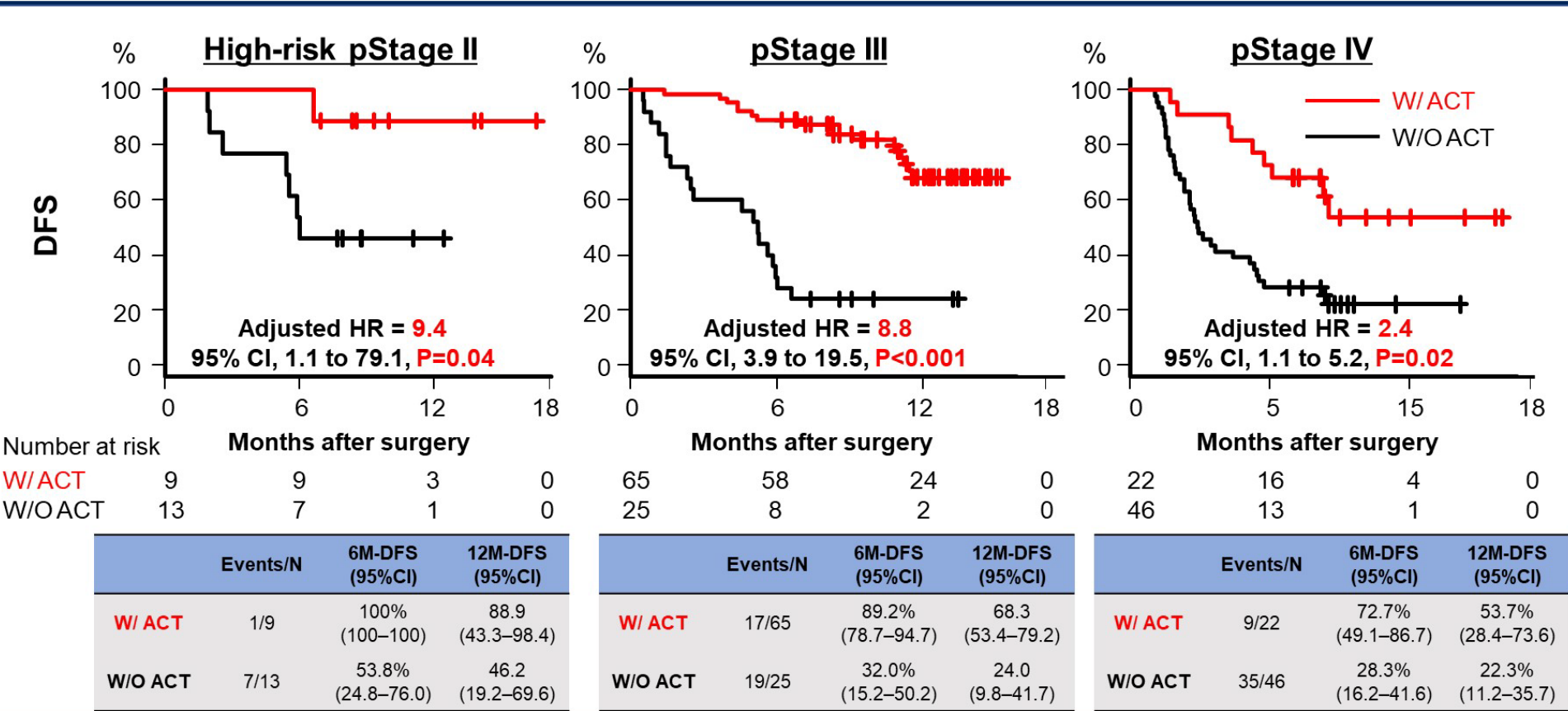
Tie J et al. N Engl J Med. 2022 Jun 16;386(24):2261-2272.

CIRCULATE-Japan



GALAXY-significant benefit from adjuvant chemo for MRD+

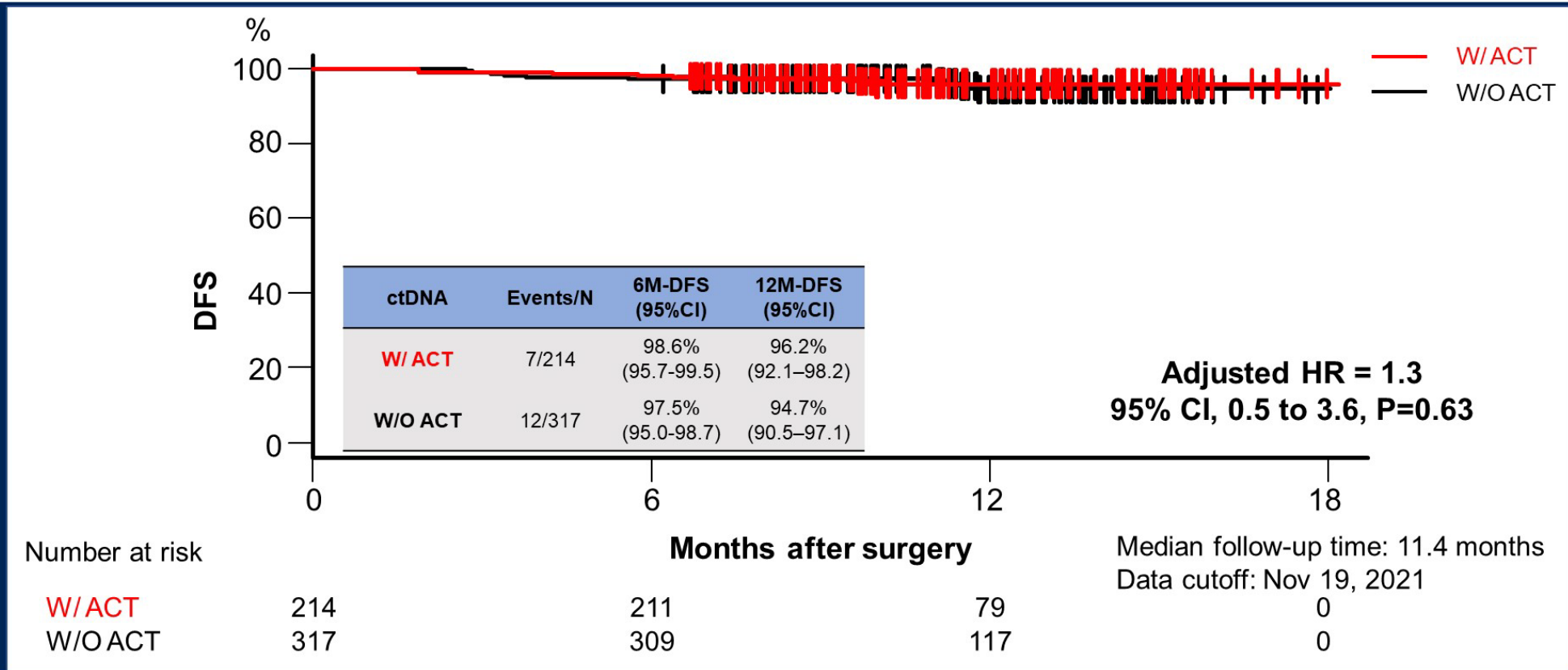
DFS by pStage in post-op-4w ctDNA positive population



HR was adjusted by sex, and performance status. ACT, adjuvant chemotherapy; DFS, disease-free survival; HR, hazard ratio; CI, confidential interval. DFS curve was estimated by the Kaplan-Meier method. HR and 95%CI were calculated by the Cox proportional hazard model.

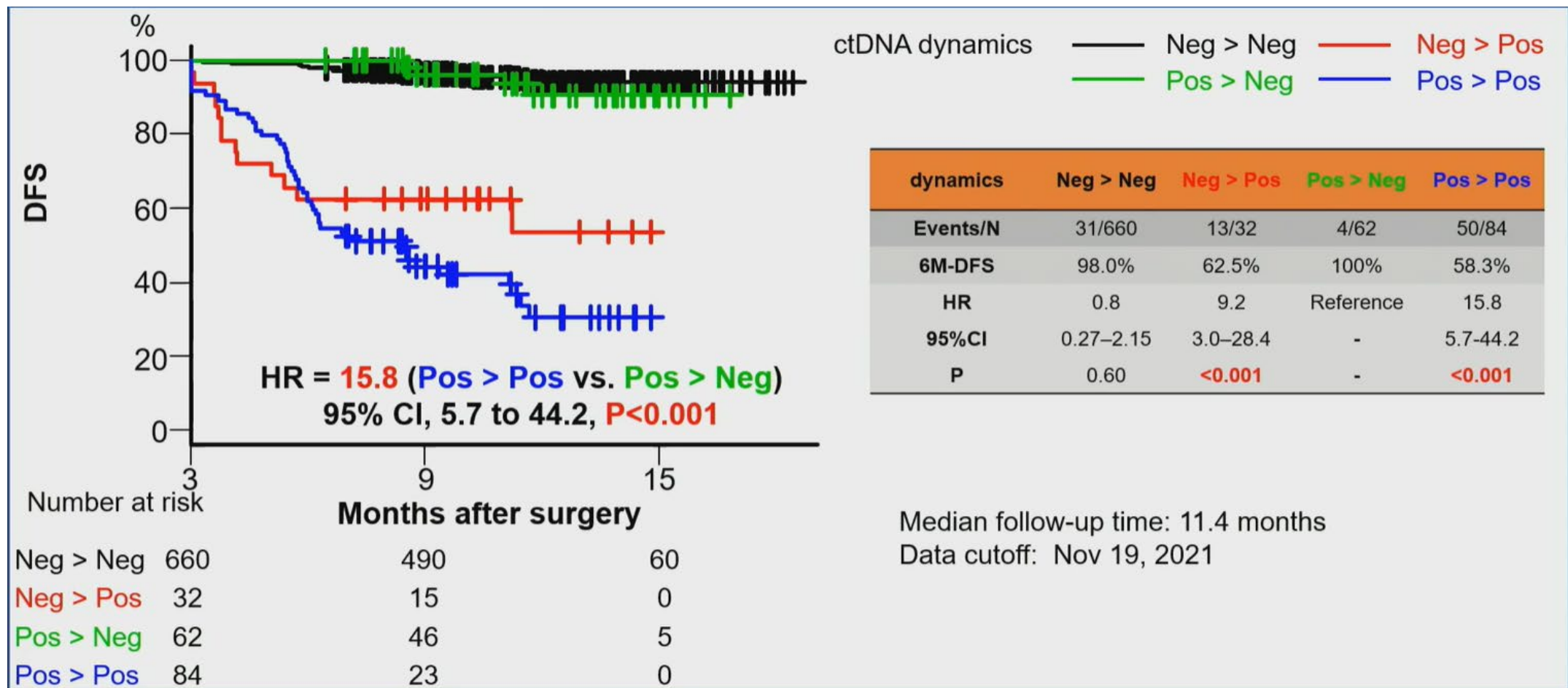
GALAXY -no benefit from adj chemo for MRD-

DFS by ACT in post-op-4w ctDNA negative population (High-risk pStage II-III)



HR was adjusted by age, performance status, pStage, and MSI status that are imbalanced between two groups.
ACT, adjuvant chemotherapy; DFS, disease-free survival; HR, hazard ratio; CI, confidential interval.
DFS curve was estimated by the Kaplan-Meier method. HR and 95%CI were calculated by the Cox proportional hazard model.

GALAXY - DFS by ctDNA dynamic from postop w4 to w12



Conclusion

**ctDNA is ready for
prime time!**

