

# ctDNA (Liquid Biopsy for MRD) in the Curative Setting

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MATOS Breast Conference

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**HARVARD**  
MEDICAL SCHOOL



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# Objectives

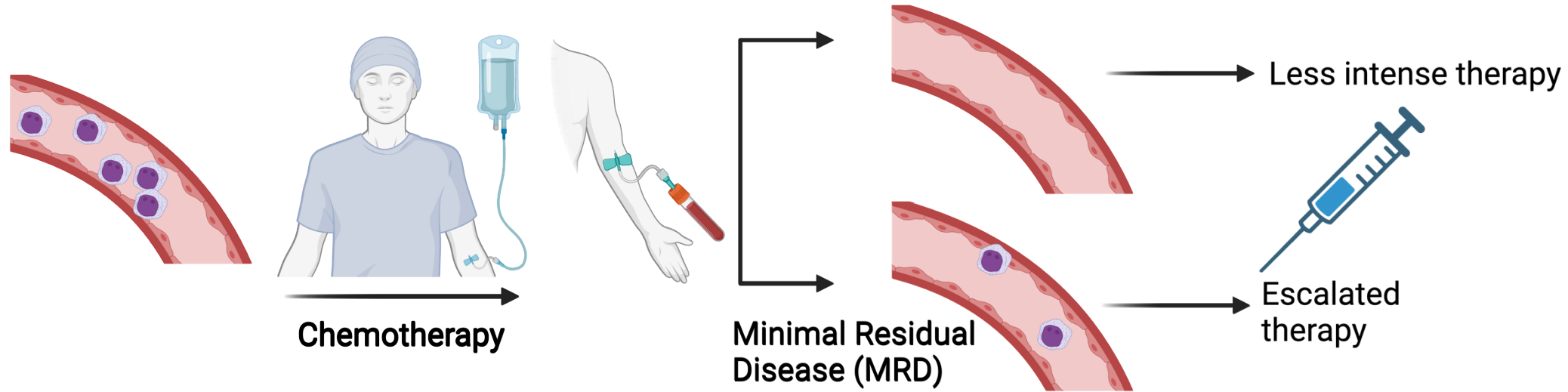
- What is molecular residual disease (MRD)?
- How is MRD detected?
- What is the prognostic significance of MRD?
- Can we intervene on MRD?

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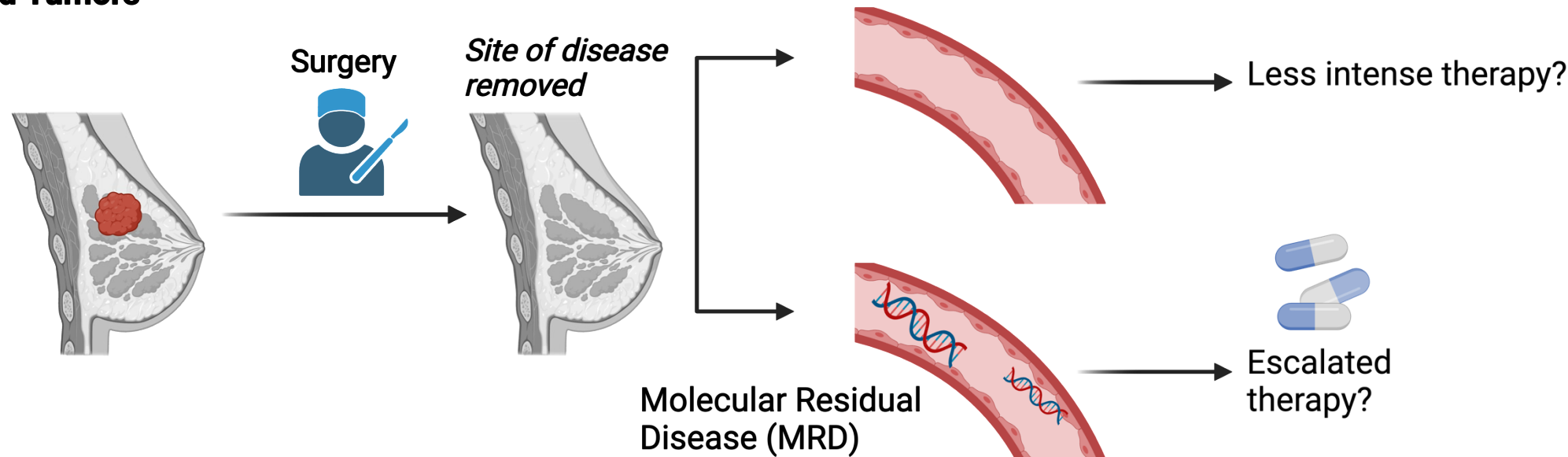
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# What is MRD?

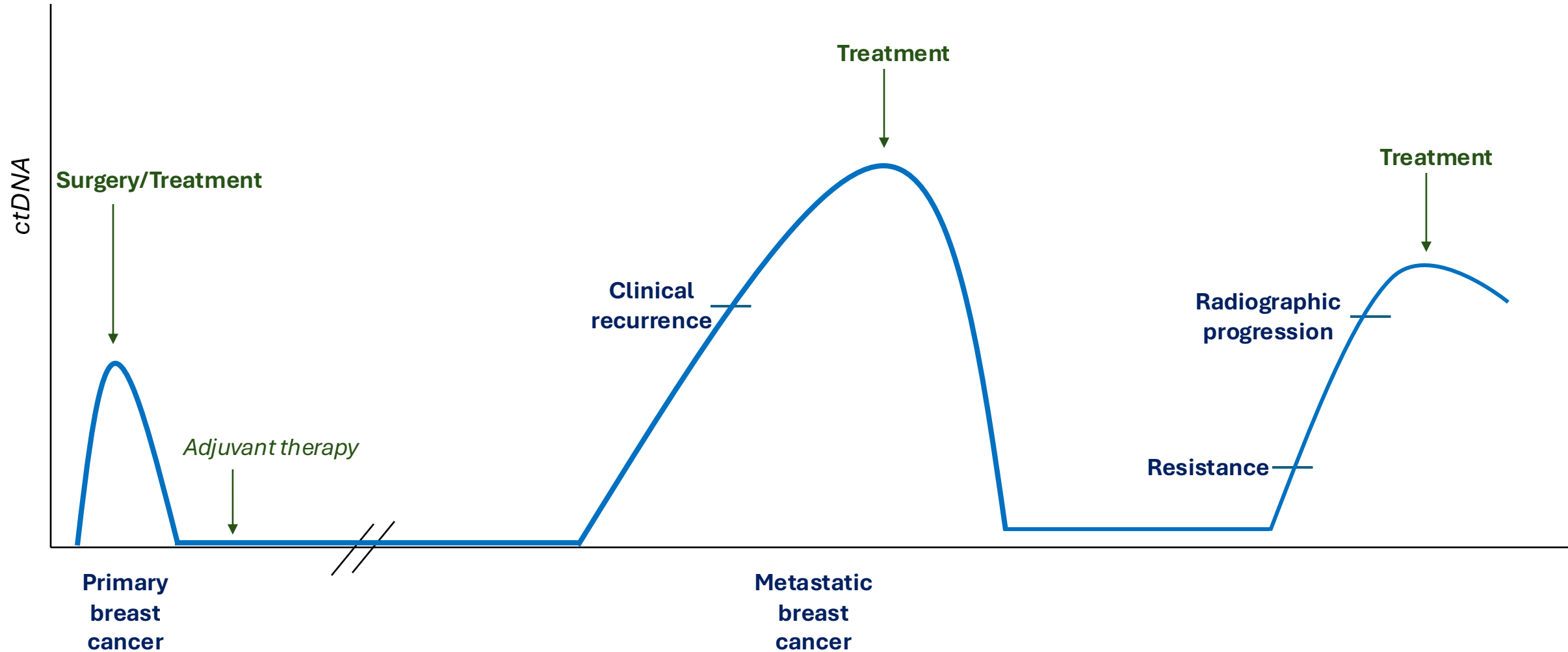
## Hematologic Malignancies

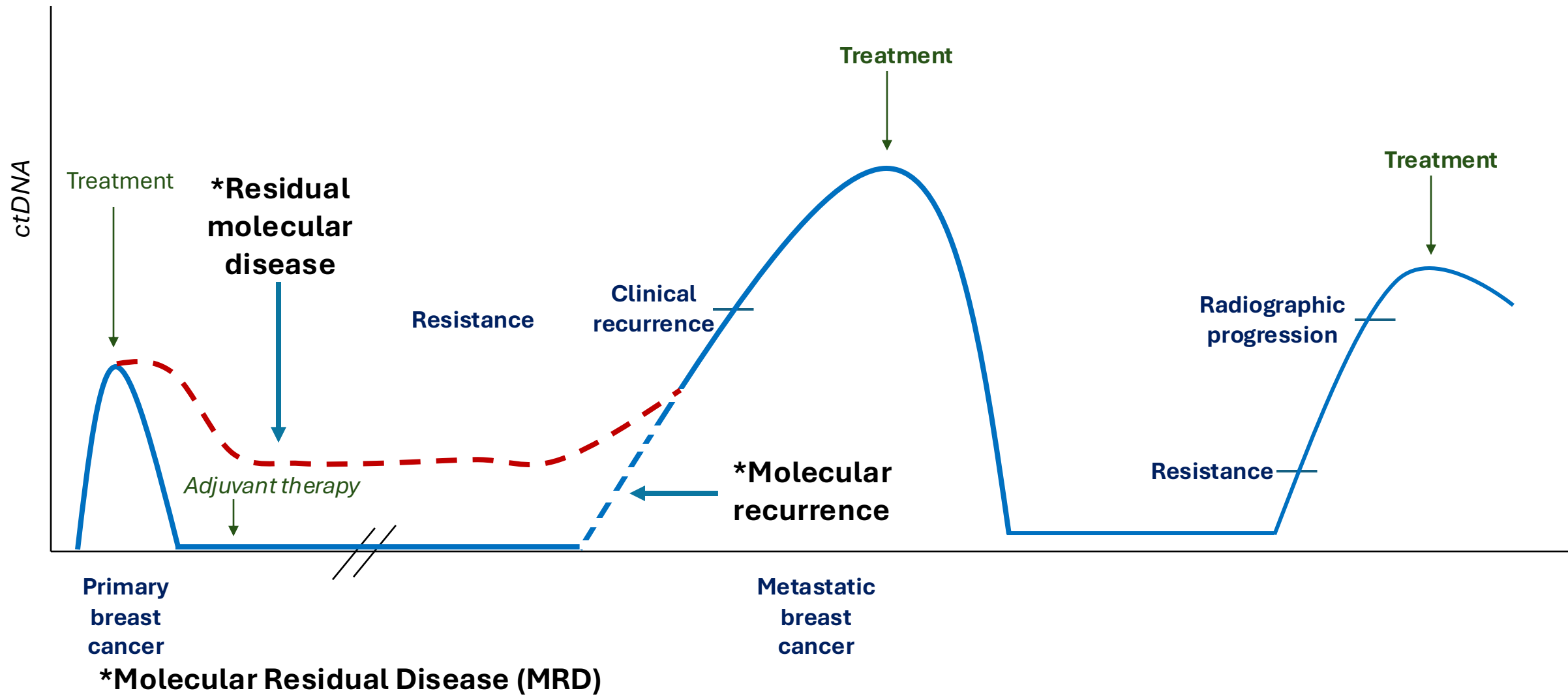


## Solid Tumors



# Circulating tumor DNA (ctDNA)



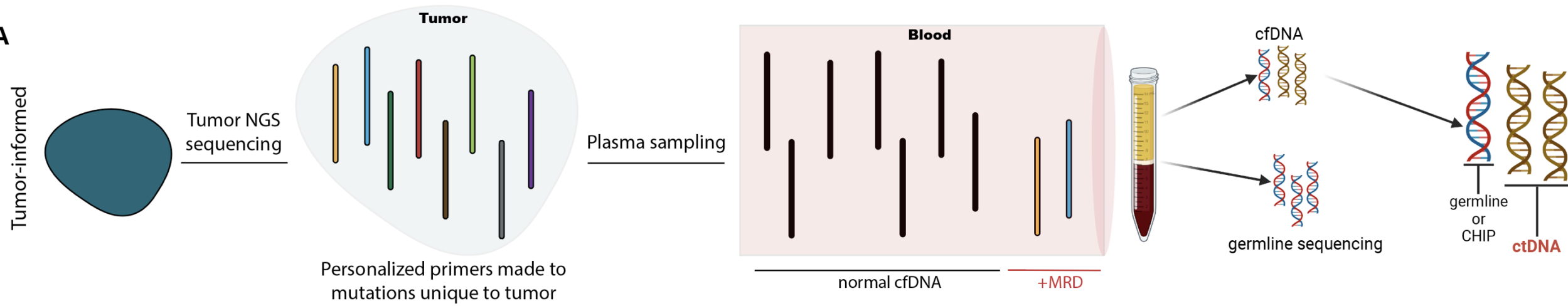


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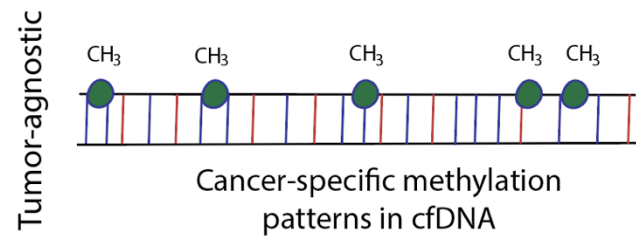
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# Types of MRD Assays

**A**



**B**

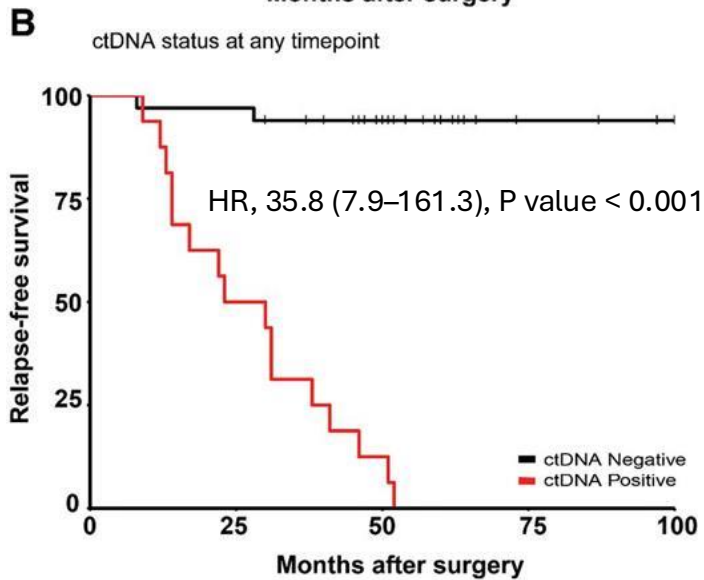
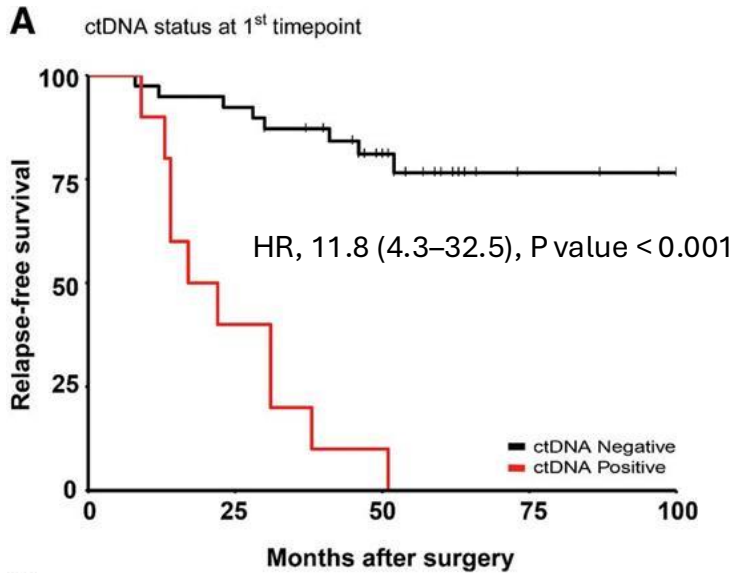




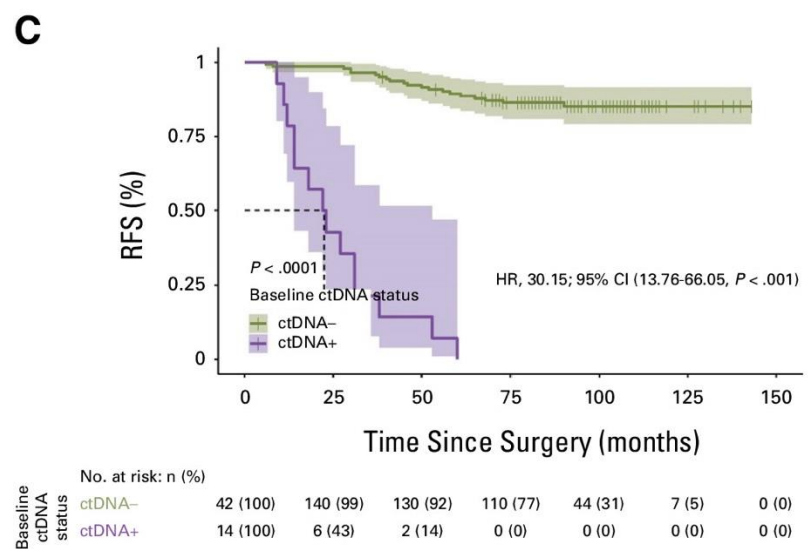
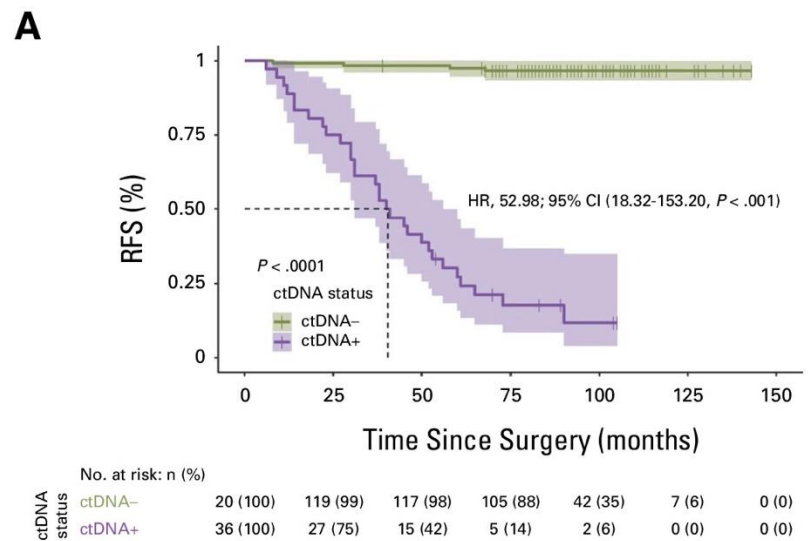
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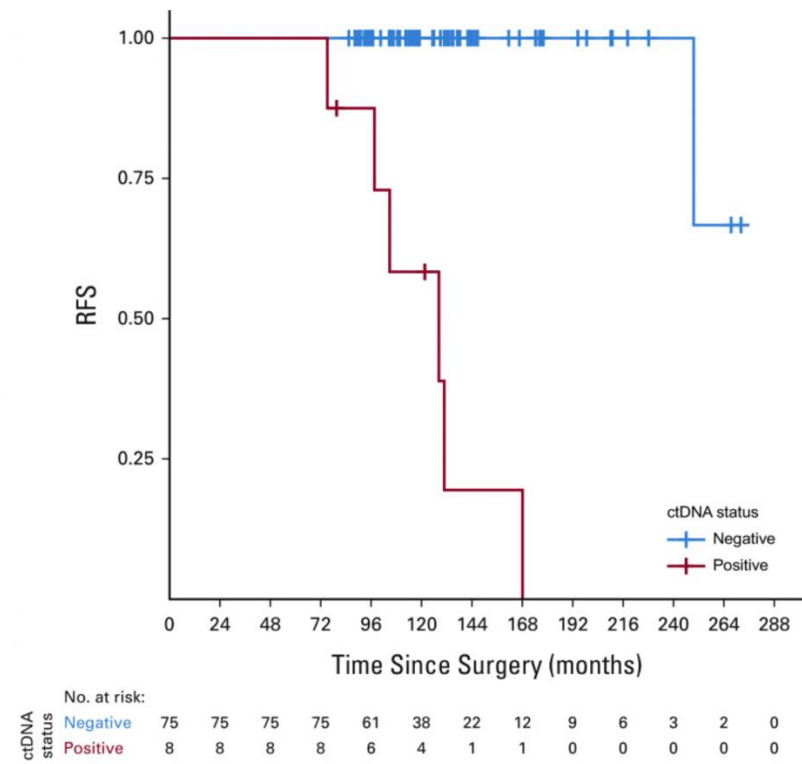
# MRD means a very poor prognosis – hazard ratio up to 50+, lead time to clinical recurrence 9-18 months



Coombes et al, CCR 2019



Shaw et al, JCO PO 2024



Lipsyc-Sharf, JCO 2022

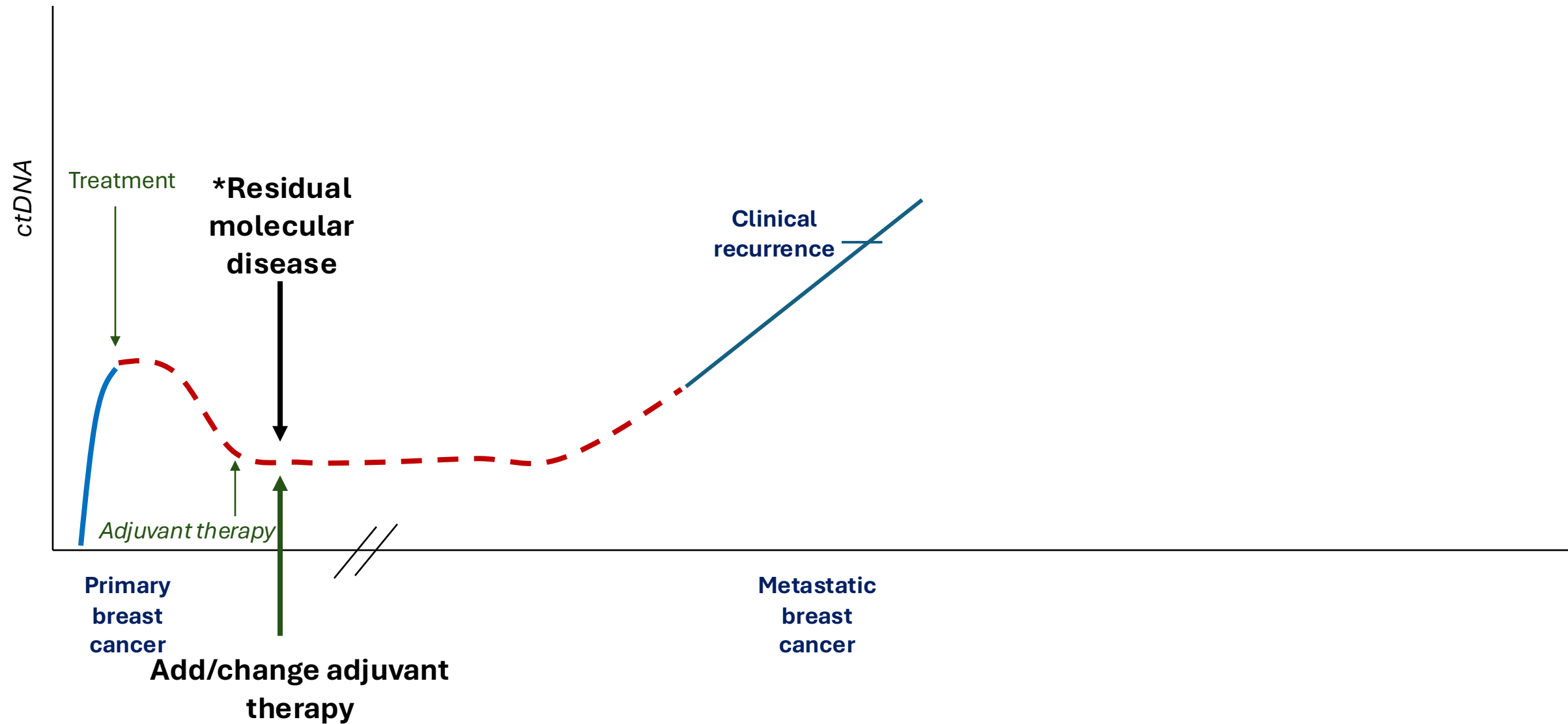
High-risk stage II-III ER+/HER2-  
≥5 years after surgery  
8 (10%) with +MRD  
6 distant recurrences at 2-years, all +MRD

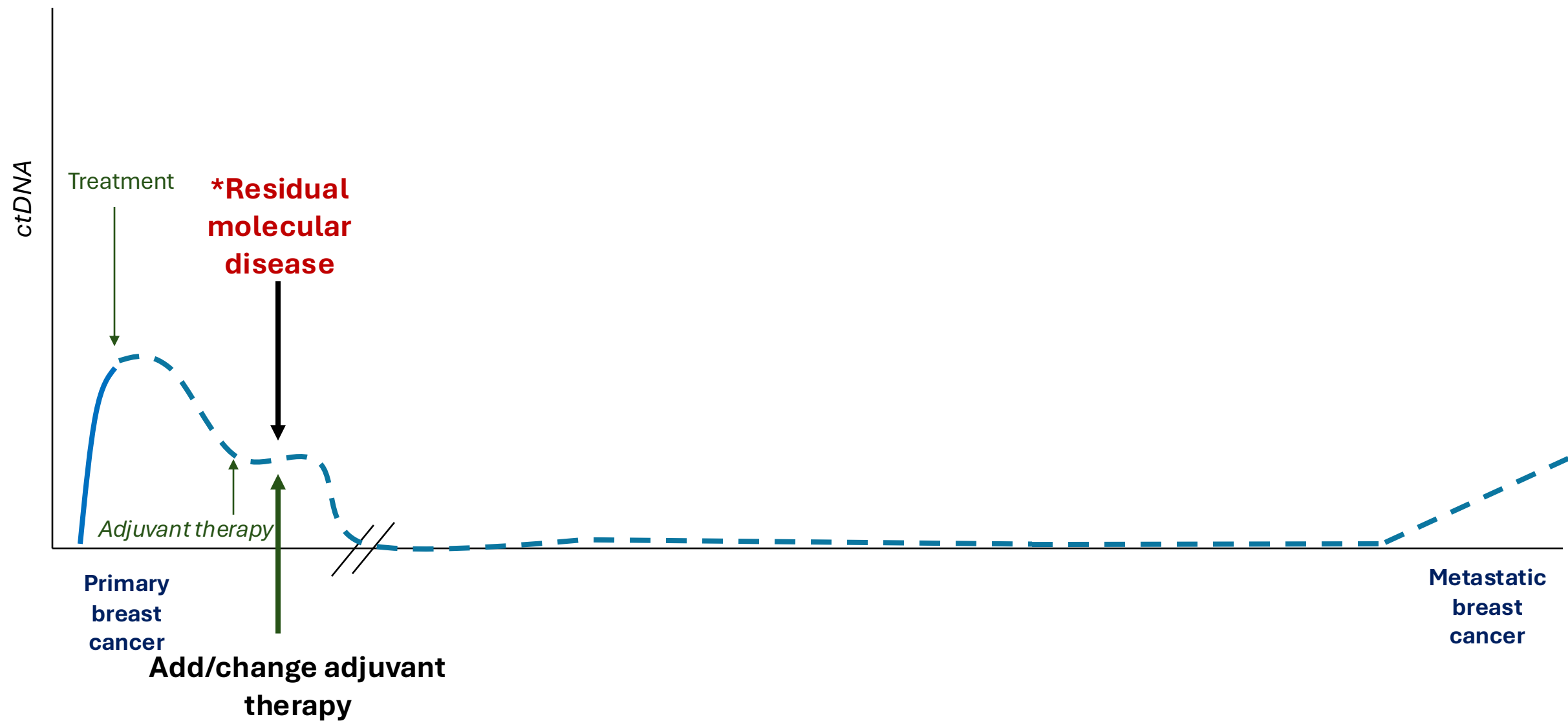
# MRD detection & lead times to BC recurrence

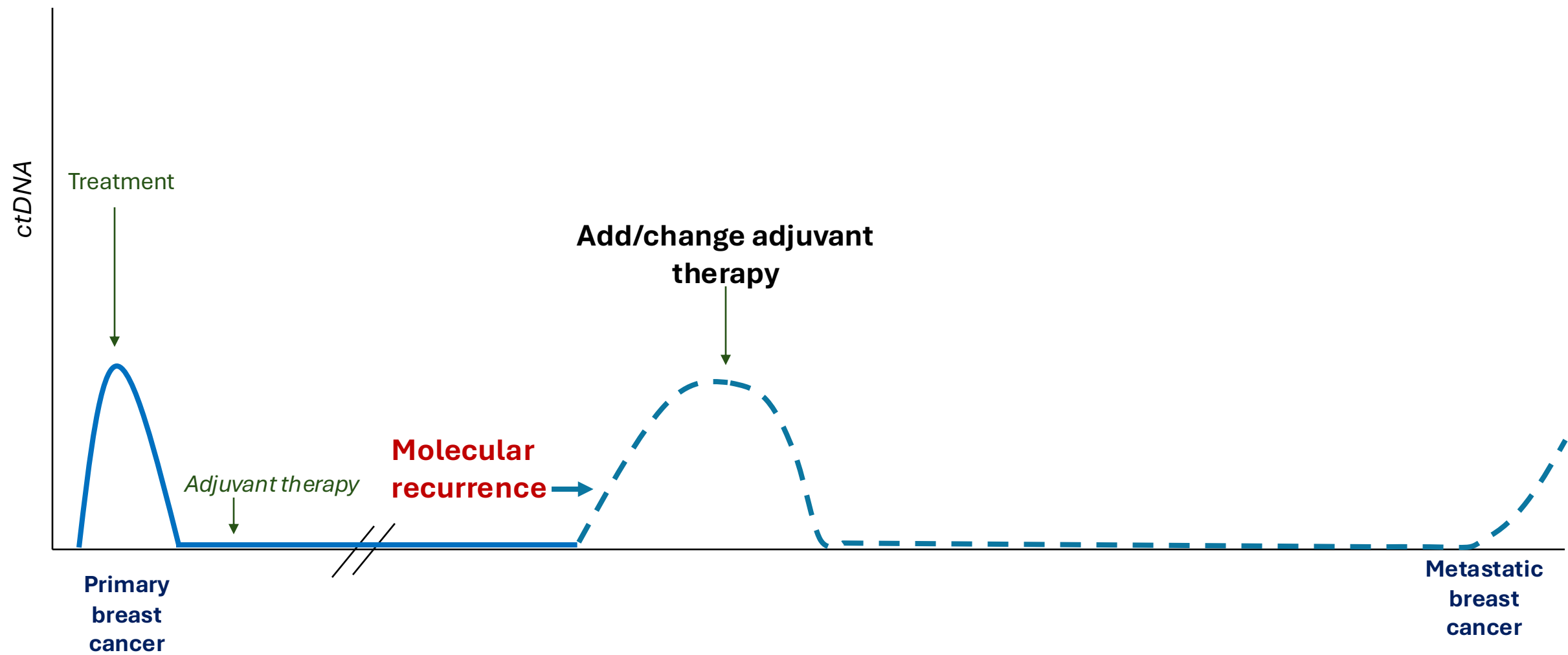
| Study                        | Assay                   | Cancer Type  | # patients                                  | Median Lead Time                              | Hazard Ratio (RFS)   | Median f/u                                    | Sensitivity                          | Specificity                       |
|------------------------------|-------------------------|--------------|---|---|--|---|--------------------------------------|-----------------------------------|
| Garcia-Murillas et al (2015) | dPCR                    | All subtypes | 55  | 7.9 months                                    | Single timepoint: 25.1 (95% CI 4.08-130.5)<br>Serial: 12.0 (95% CI 3.36-43.07)     | ~2 years                                      | Post-surgery: 50%<br><br>Serial: 80% | 96%                               |
| Olsson et al (2015)          | dPCR                    | All subtypes | 20  | 11 months                                     | N/A  | 9.2 years                                     | 93%                                  | 100%                              |
| Garcia-Murillas et al (2019) | dPCR                    | All subtypes | Primary cohort: 101<br>Combined cohort: 144 | Primary: 38.0 months<br>Combined: 10.7 months | Primary: 16.7 (95% CI 3.5-80.5, p < .001)<br>Combined: 17.4 (95% CI 6.3-47.8)      | Primary: 35.5 months<br>Combined: 36.3 months | 88.4%                                | 100%                              |
| Coombes et al (2019)         | Signatera               | All subtypes | 49  | 8.9   | Post-surgery: 11.8 (95% CI 4.3–32.5)<br><br>Serial: 35.8 (95% CI 8.0–161.3)        | ~2 years (study midpoint)                     | 89%                                  | 100%                              |
| Parsons et al (2020)         | Internal tumor-informed | HR+/HER2-    | 35  | 18.9 months                                   | Post-surgery: 5.1 (95% CI 2.0–12.7]<br>1 year post surgery: 20.8 (95% CI 7.3–58.9) | 7.1 years                                     | Post-surgery: 23%<br>1-year: 19%     | Post-surgery: 96%<br>1-year: 100% |
| Lipsyc-Sharf et al (2022)    | RaDaR                   | HR+/HER2-    | 85  | 12.4 months                                   | N/A  | 10.4  | 85.7%                                | 97.4%                             |

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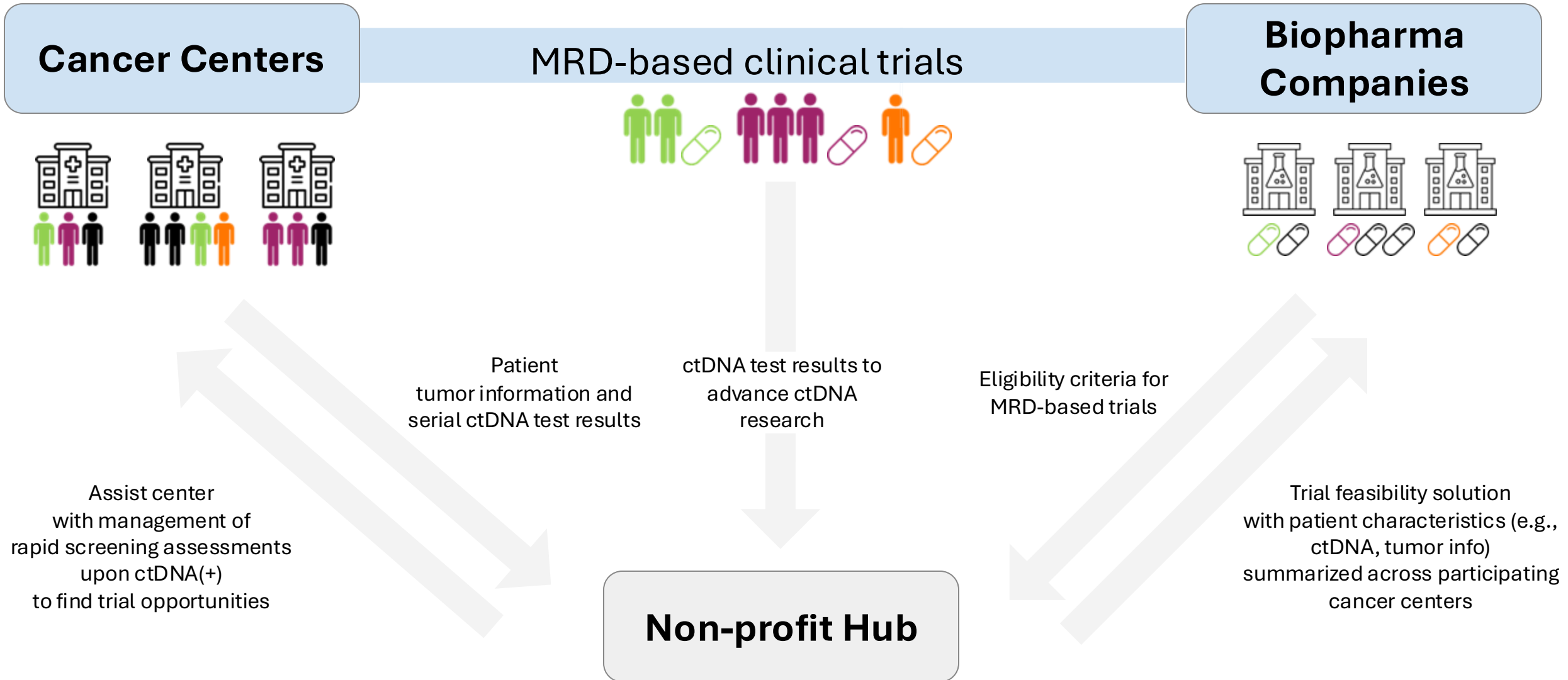


# Prospective interventional clinical trials in MRD and Breast Cancer

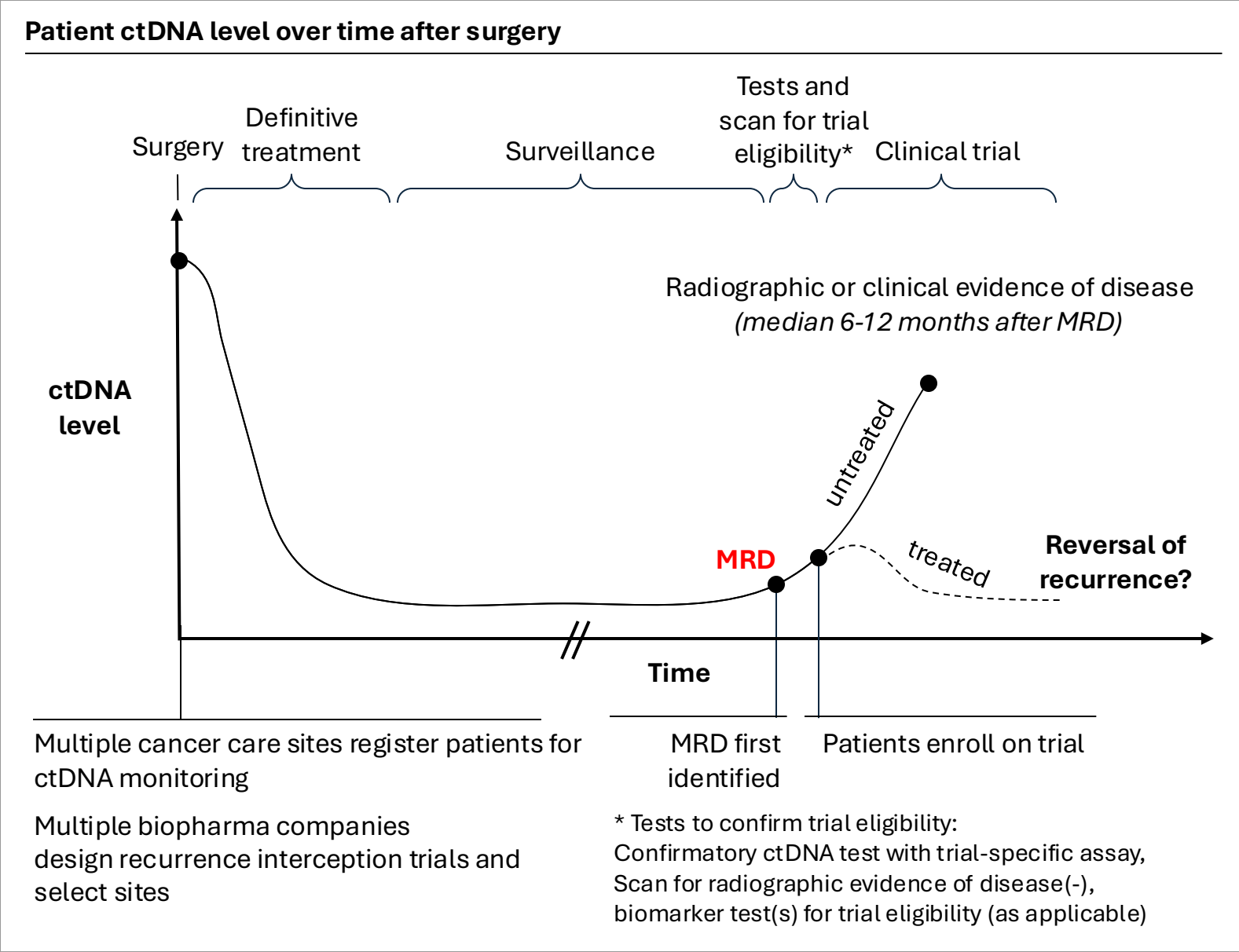
| Trial ID/name/ disease/phase   | Drug treatment                             | Primary end point  | Trial eligibility: MED <sup>a</sup> | Trial eligibility: scan required <sup>a</sup> | Sample size: ctDNA <sup>+</sup> patients in intervention arm <sup>b</sup> | Trial status              | Biopharma sponsored | Geography                               |
|--|--|--|-------------------------------------|---|---|---------------------------|---------------------|---|
| <b>Breast cancer</b>   |  |  |                                     |   |   |                           |                     |   |
| DARE NCT04567420 (ref. 73)<br>Stage II–III ER <sup>+</sup> /HER2 <sup>−</sup> breast<br>Phase II   | Palbociclib +<br>endocrine<br>therapy      | ctDNA positivity<br>during surveillance<br>Relapse-free<br>survival                | Landmark or<br>surveillance         | Yes, scan<br>required                         | 100 (target)  | Recruiting                | No                  | USA (13 states;<br>15 sites)            |
| LEADER NCT03285412 (ref. 74)<br>Localized ER <sup>+</sup> /HER2 <sup>−</sup> breast<br>Phase II  | Ribociclib +<br>endocrine<br>therapy       | ctDNA clearance  | Landmark or<br>surveillance         | No  | 30 (target) <sup>c</sup>  | Recruiting                | No                  | Massachusetts<br>(3 sites)              |
| TRAK-ER<br>NCT04985266 (ref. 75)<br>Early stage ER <sup>+</sup> /HER2 <sup>−</sup> breast<br>Phase II  | Palbociclib +<br>fulvestrant               | ctDNA positivity<br>during surveillance<br>Relapse-free<br>survival                | Landmark or<br>surveillance         | Yes, scan<br>required<br>after MED            | 1,100<br>(surveillance<br>target) <sup>d</sup>                            | Recruiting                | No                  | UK and France<br>(36 sites)             |
| TREAT ctDNA<br>NCT05512364 (ref. 76)<br>Early stage ER <sup>+</sup> /HER2 <sup>−</sup> breast<br>Phase III   | Elacestrant                                | Distant<br>metastasis-free<br>survival   | Landmark or<br>surveillance         | Yes, scan<br>required<br>after MED            | 220 (target)  | Recruiting                | No                  | Global<br>(>10 countries,<br>120 sites) |
| ZEST<br>NCT04915755 (ref. 77)<br>Stage I–III triple-negative breast<br>or HR <sup>+</sup> /HER2 <sup>−</sup> breast with BRCA<br>mutation<br>Phase III | Niraparib                                  | Treatment<br>emergent adverse<br>events<br>Change in ECOG<br>Performance<br>Status | Post-op or<br>landmark              | No  | 43 (actual)   | Discontinued              | Yes                 | Global<br>(>10 countries,<br>200 sites) |
| c-TRAK-TN<br>NCT03145961 (ref. 78)<br>Early stage triple-negative breast<br>Phase II   | Pembrolizumab                              | ctDNA positivity<br>during surveillance<br>ctDNA clearance                         | Landmark or<br>surveillance         | Yes, scan<br>required                         | 45 (actual) <sup>79</sup>   | Complete                  | No                  | UK (17 sites)                           |
| ASPRIA<br>NCT04434040 (ref. 80)<br>Early stage triple-negative breast<br>Phase II  | Atezolizumab +<br>sacituzumab<br>govitecan | ctDNA clearance  | Landmark                            | No  | 40 (actual)   | Active, not<br>recruiting | No                  | USA (5 states,<br>9 sites)              |



# A new approach to MRD-based clinical trials



# A standing platform for MRD-based clinical trials



# Summary

- MRD indicates breast cancer is going to recur.
- Patients are able to access MRD tests today, but no therapies are approved to intervene. **This is a key area of unmet clinical need.**
- Innovative translational research is required to understand mechanisms of therapy resistance and optimal MRD interventional clinical trials.
- An overhaul of the MRD-based clinical trial infrastructure is needed (and underway) to improve patient access and study opportunities.

# Thank you!

## **Mentorship**

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## **Science for America/Reversing Early Recurrence**

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Avanish Mishra

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Rachel Abelman  
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Michelle Trodella  
Lauren Scarpetti

## **LEADER Study Team**

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## **Getz Lab**

Team thanks!