# Liquid biopsies in clinical trials



#### Scott V. Bratman, MD PhD

Dr. Mariano Elia Chair in Head & Neck Cancer Research Associate Professor & Clinician-Scientist Princess Margaret Cancer Centre University Health Network University of Toronto

> CCTG NICTC August 22, 2024



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# Disclosures

#### **Employment Relationships:**

- University Health Network: Staff Radiation Oncologist Clinician-Scientist
- Adela: Co-Founder and Chief Innovation Officer

Consulting or Advisory Board/Speaker/Faculty/Research/Honoraria:

• EMD Serono: Advisory Board

Ownerships/Investments/Intellectual Property:

- Adela: Licensed patents, stock
- Roche: Licensed patent





# Learning Objectives

- To discuss the role for circulating tumour DNA as a key correlative component of clinical trials
- To recognize the distinct scenarios where liquid biopsy can inform on patient outcomes or trial endpoints
- To assess the limitations of current liquid biopsy tests, and opportunities for future improvements





# Lecture Overview

- Summary of liquid biopsy in oncology
- Types of ctDNA tests and practical considerations
- Evidence for ctDNA tests as tools to guide therapy





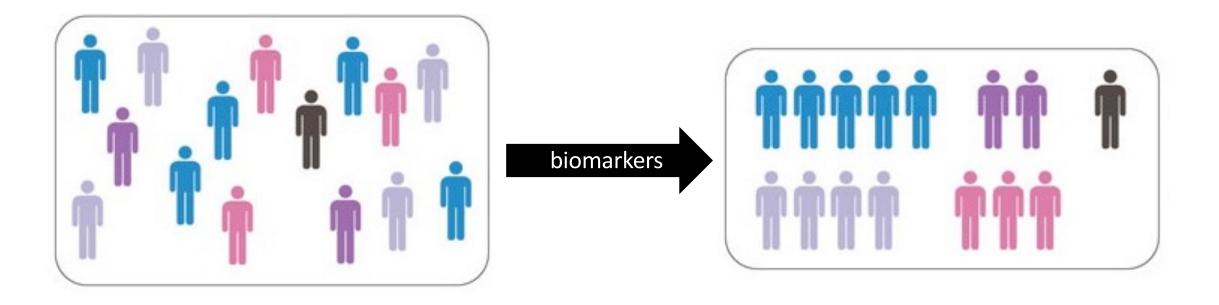
## Lecture Overview

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### How can biomarkers guide personalized treatment regimens?



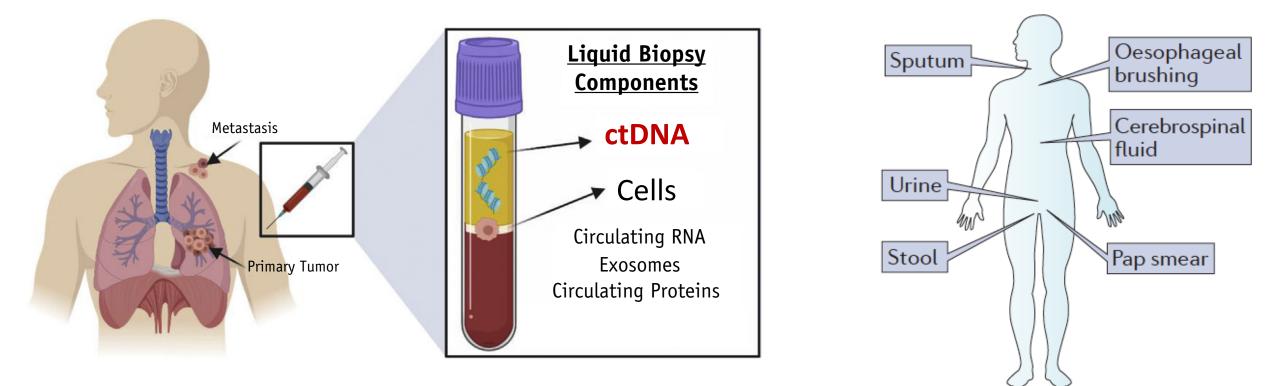
- Risk stratification / prognostication
- Prediction of response or toxicity
- Dynamic monitoring of response
- Surrogate endpoint for clinical trials

Baumann, Nature Rev Cancer 2016





# Circulating tumor DNA (ctDNA): Cancer DNA in the bloodstream

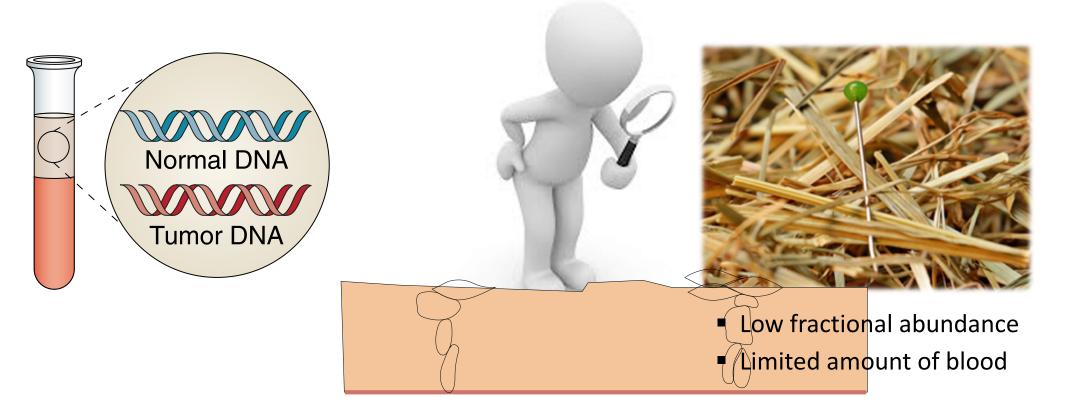


Wan...Rosenfeld et al. Nat Rev Cancer 2017



De Michino, Lok, Bratman et al. IJROBP 2020; 107(5),873-886 UNIVERSITY OF TORONTO FACULTY OF MEDICINE

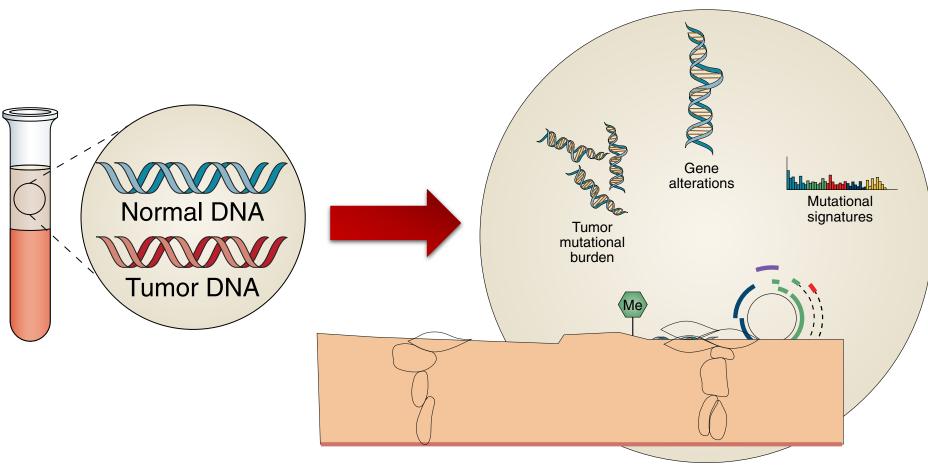
# ctDNA: Finding a needle in the haystack of normal non-cancer-derived cell-free DNA







## ctDNA: Distinguishing signal from noise



**Tumor-derived DNA markers** 

Cescon, Bratman, Chan, Siu. Nature Cancer 2020





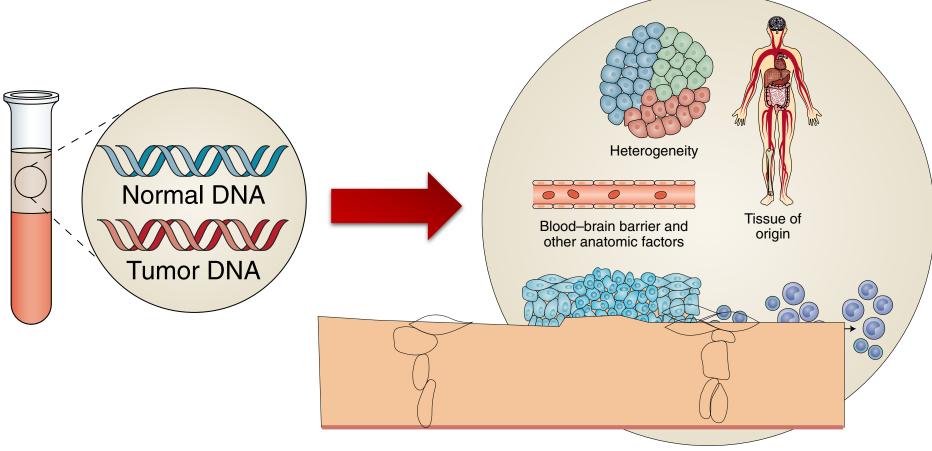
## ctDNA: Distinguishing signal from noise

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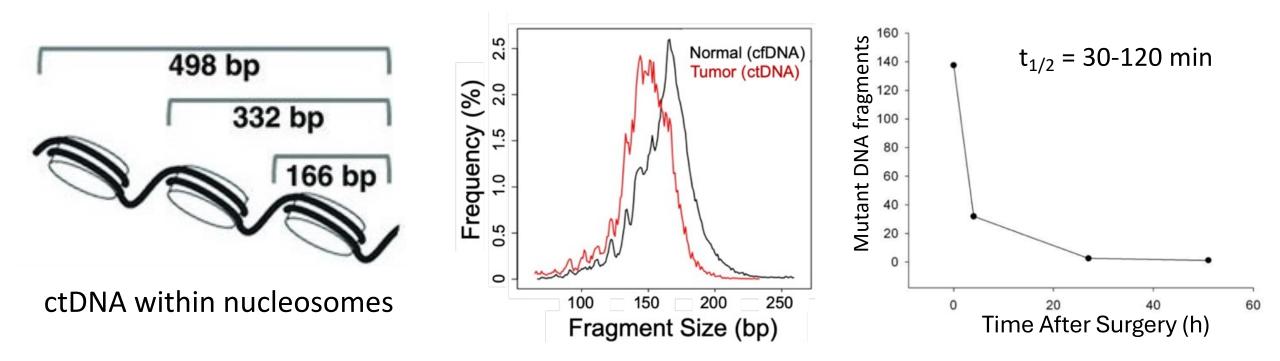
#### **Challenges to ctDNA detection**



Cescon, Bratman, Chan, Siu. Nature Cancer 2020



ctDNA is associated with nucleosome complexes and has a characteristic size distribution and short half life

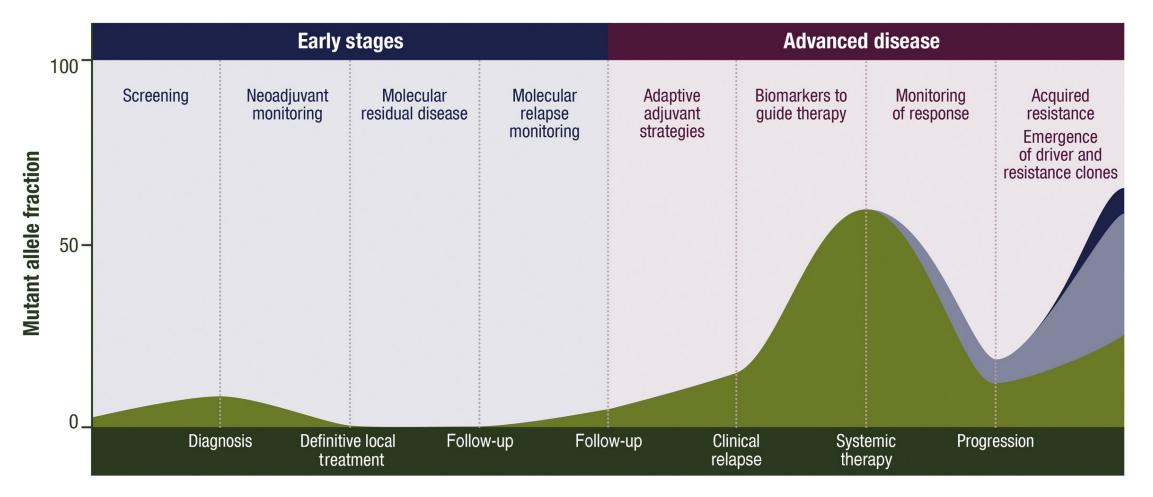


Heitzer, Ulz, Geigl, Clin Chem 2015 Leung, Han...Bratman, Clin Cancer Res 2021 Diehl...Diaz et al. Nat Med 2008





# Levels of ctDNA vary across distinct phases of the cancer care continuum



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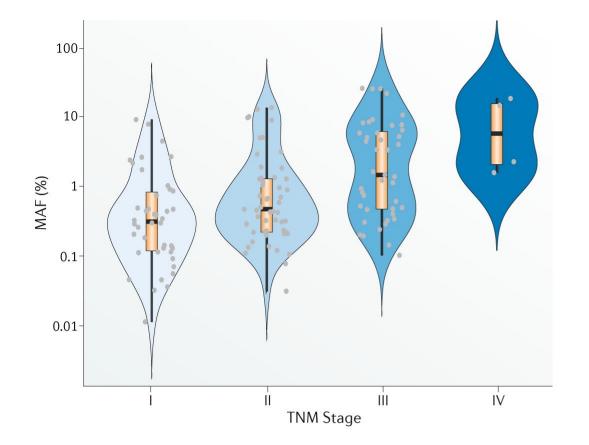
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Pascual, Turner et al. Annals Oncol 2022



#### Clinical factors influence ctDNA detectability & abundance



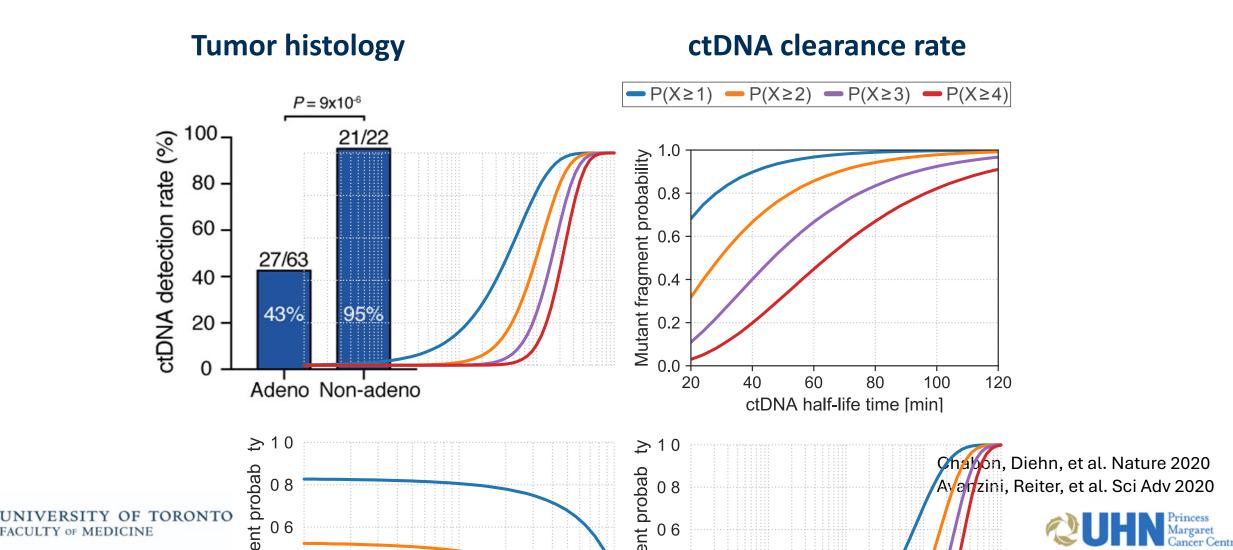
- Stage (M > N > T)
- Tumor volume (weak)
- Viability or necrosis
- Vascularization
- Blood brain barrier

Abbosh, Birkbak, Swanton. Nat Rev Clin Oncol 2018



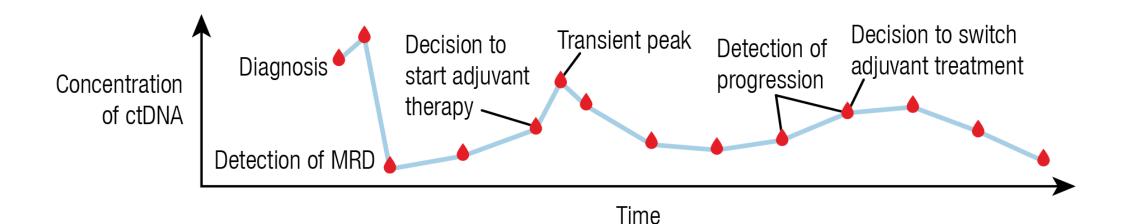


Impact of tumor biology & host factors on ctDNA detectability



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### Clinical actionability of on-treatment ctDNA kinetics



#### **Factors affecting ctDNA kinetics**



Degradation & Clearance Kinetics

Sanz Garcia, Zhao, Bratman, Siu, et al. Science Advances, 2022





# Lecture Overview

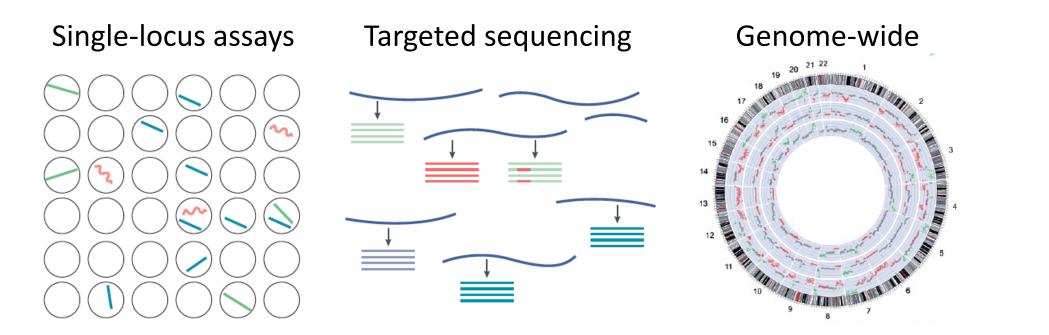
- Summary of liquid biopsy in oncology
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# Methods for ctDNA Detection & Analysis

- Targeting of genetic or epigenetic aberrations within cell-free DNA
- Accounting for tumor-derived **signal** as well as technical and biological **noise**



Wan...Rosenfeld et al. Nat Rev Cancer 2017



# Methods for ctDNA Detection & Analysis

### **Polymerase chain reaction (PCR)**

- Interrogates few targets at a time
- High analytical sensitivity
- Low cost & complexity
- Rapid procedures
- Mature technology

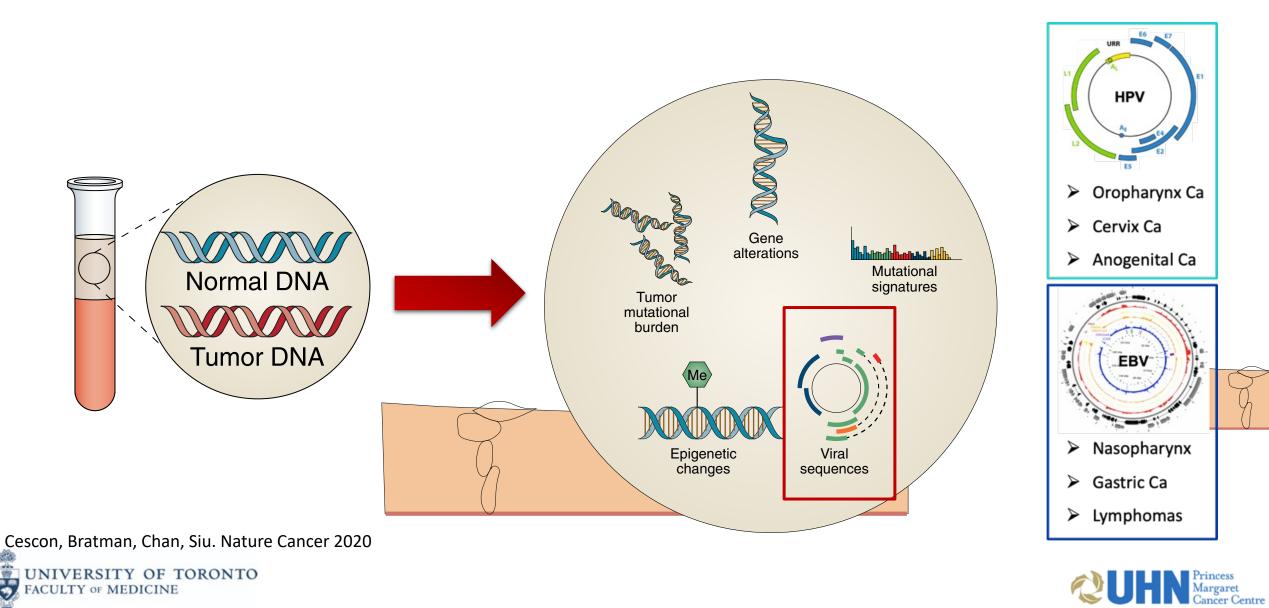
### Sequencing

- Can interrogates many targets simultaneously
- Analytical sensitivity is highly platform-dependent
- High cost & complexity
- Longer time-to-data
- Emerging technologies





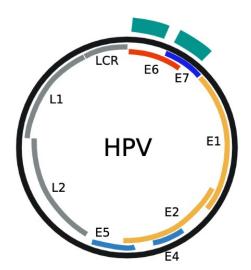
## Plasma viral DNA is the archetypal ctDNA biomarker



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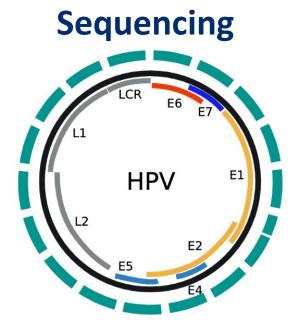
## Classes of viral ctDNA detection methodologies

PCR



Covers limited number of loci from one or multiple HPV genotypes

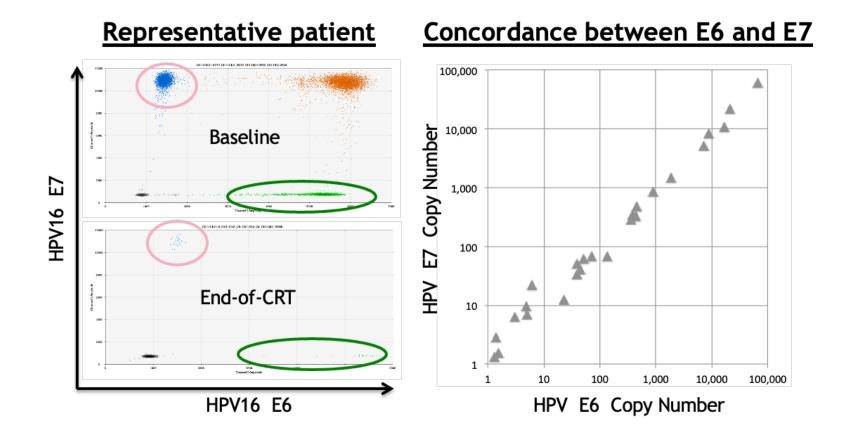
Han, Leung, Bratman et al. JCO Precis Oncol 2018 Leung, Han, Bratman et al. CCR 2021 UNIVERSITY OF TORONTO FACULTY OF MEDICINE



Allows full genome coverage of multiple HPV genotypes



### Digital PCR for HPV ctDNA is limited to 1-2 markers

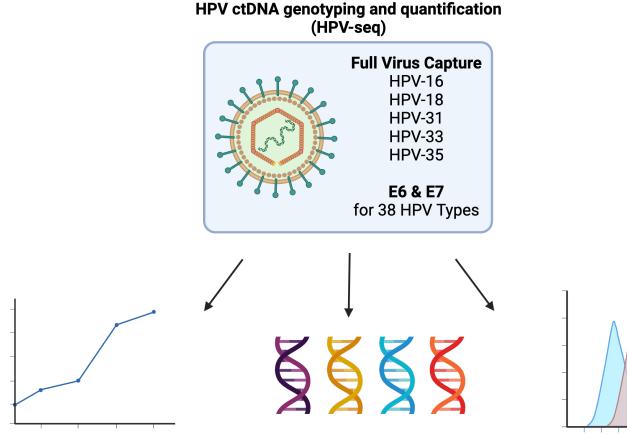




Han, Leung, Bratman, et al. JCO PO 2018



# HPV sequencing (HPV-seq) allows for deeper characterization of HPV ctDNA



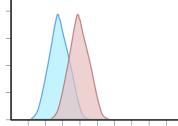
Quantification

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#### Genotyping

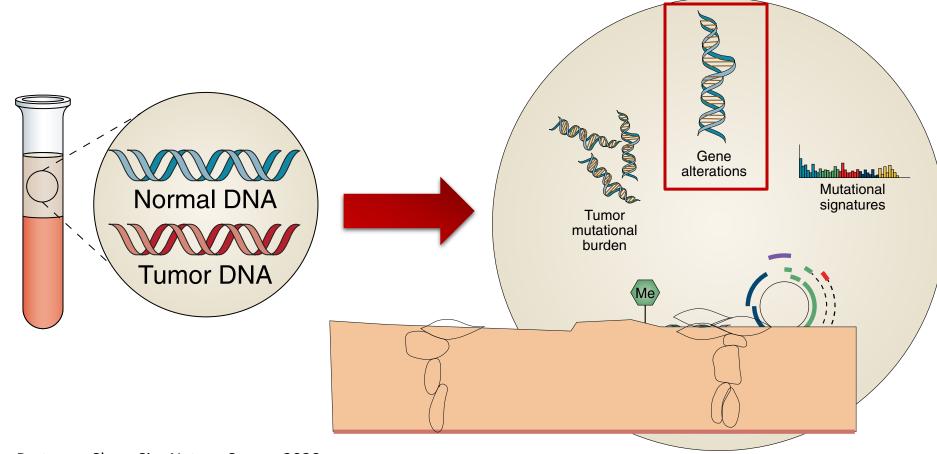
Leung, Han, Bratman, et al. CCR 2021 Han, Leung, Bratman et al. JCO 2024



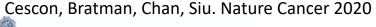
**Fragment Features** 



# Somatic genetic mutations represent the most common targets of commercial ctDNA assays

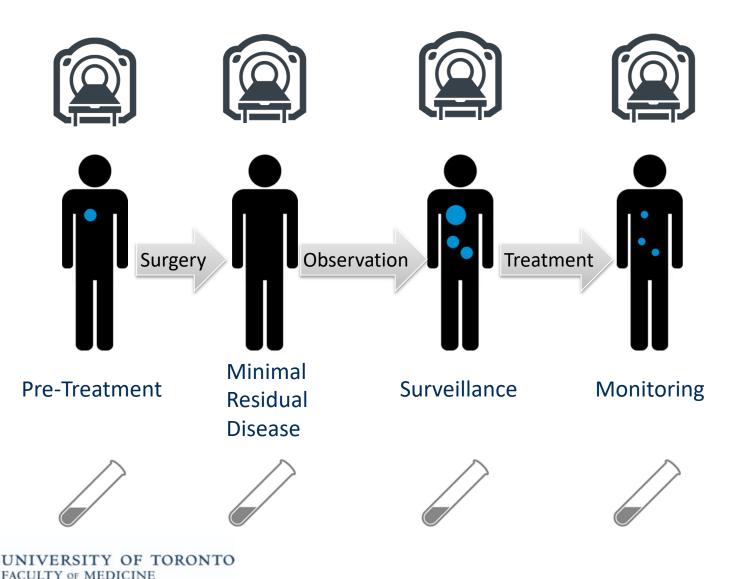






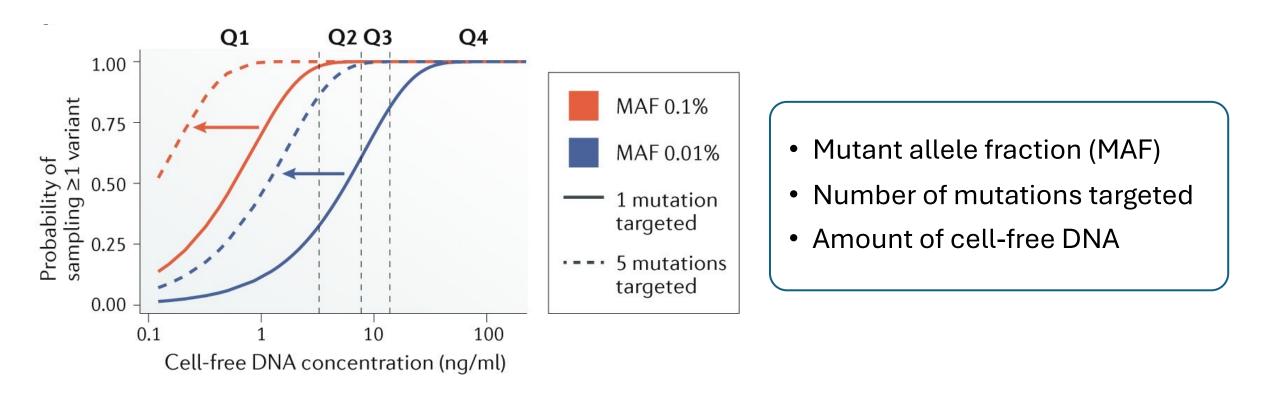
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- Sensitivity and specificity for cancer
- Quantitative ctDNA assessment
- Precedes detection by standard imaging
- Requirement of tumor tissue analysis
- Off-the-shelf vs personalized "bespoke"
- Turn-around-time to receive results

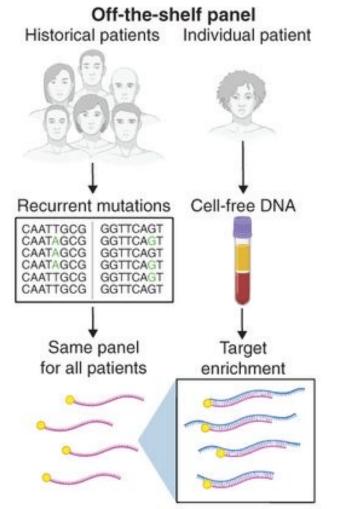




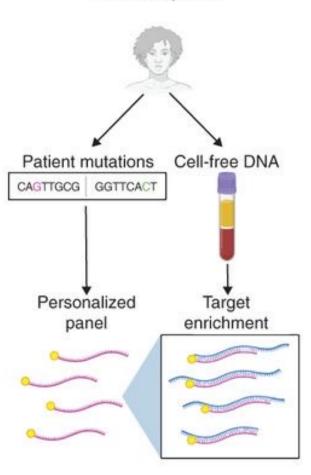


Abbosh, Birkbak, Swanton. Nat Rev Clin Oncol 2018





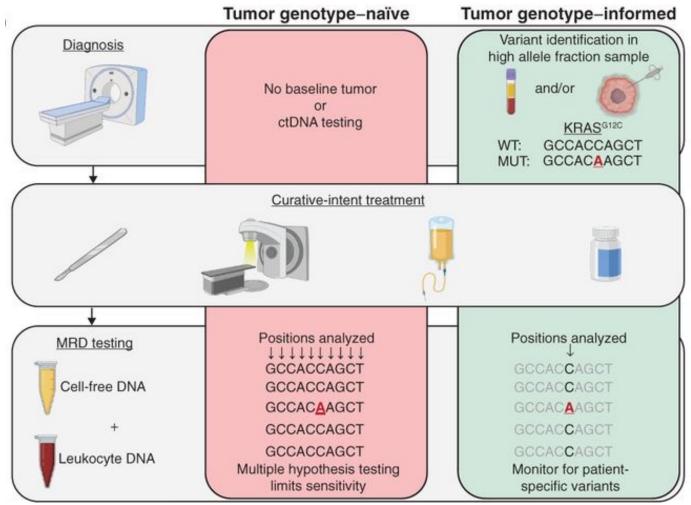
Personalized panel Individual patient



Moding, Diehn, et al. Cancer Discov. 2021;11(12):2968-2986.





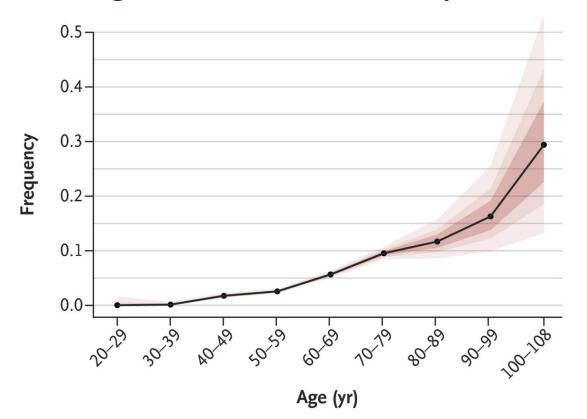




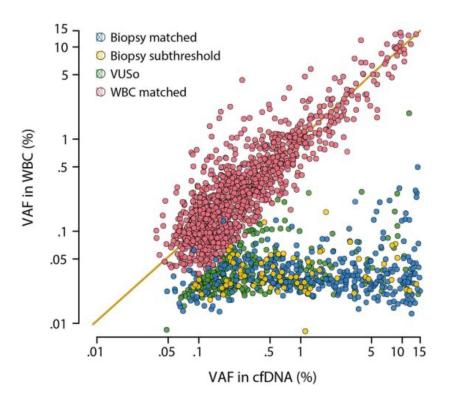




Need to account for biological sources of false-positive signal within circulating cell-free DNA



#### **Age-related clonal hematopoiesis**



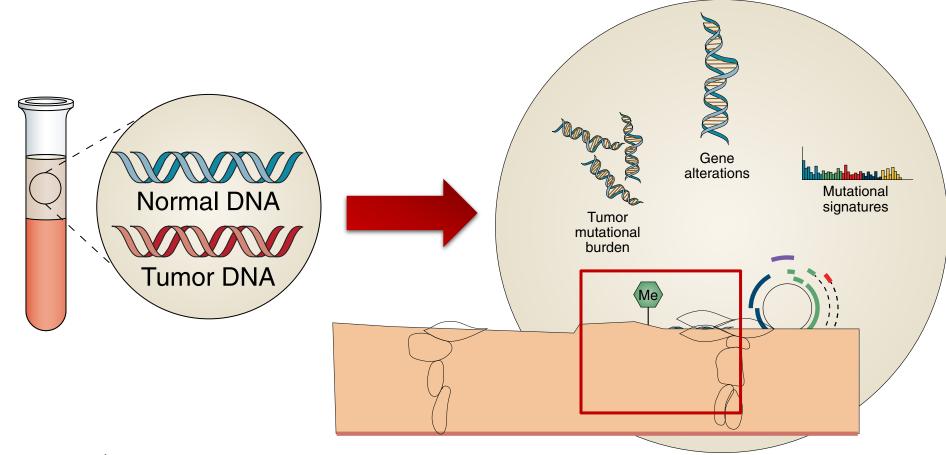
WBC mutations in cell-free DNA

Jaiswal et al. NEJM. 2014;371(26):2488-98 Razavi, et al. Nature Med 2019;25(12):1928-1937





Epigenetic changes represent another promising class of biomarkers detectable within cell-free DNA



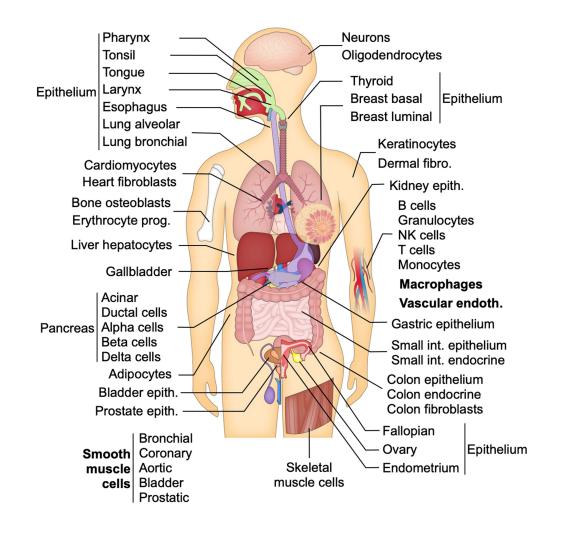




Cescon, Bratman, Chan, Siu. Nature Cancer 2020

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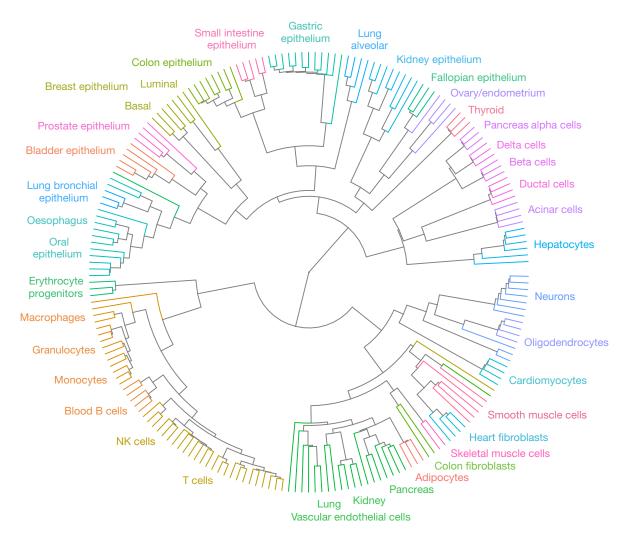
## DNA methylation patterns are specific to cell-of-origin



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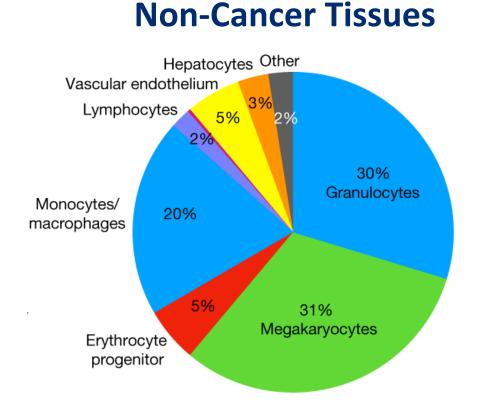
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Loyfer et al, Nature, 2023; doi: 10.1038/s41586-022-05580-6



## Cell-free DNA tissue-of-origin can be determined from analyzing DNA methylation patterns



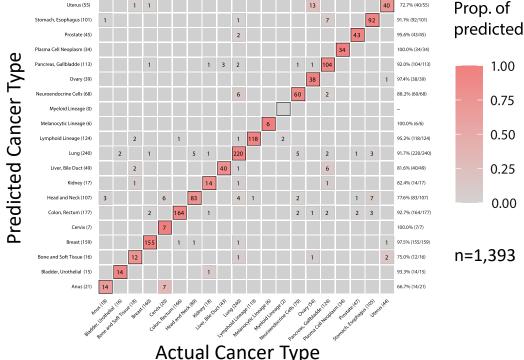
Lovfer et al. Nature (2023)

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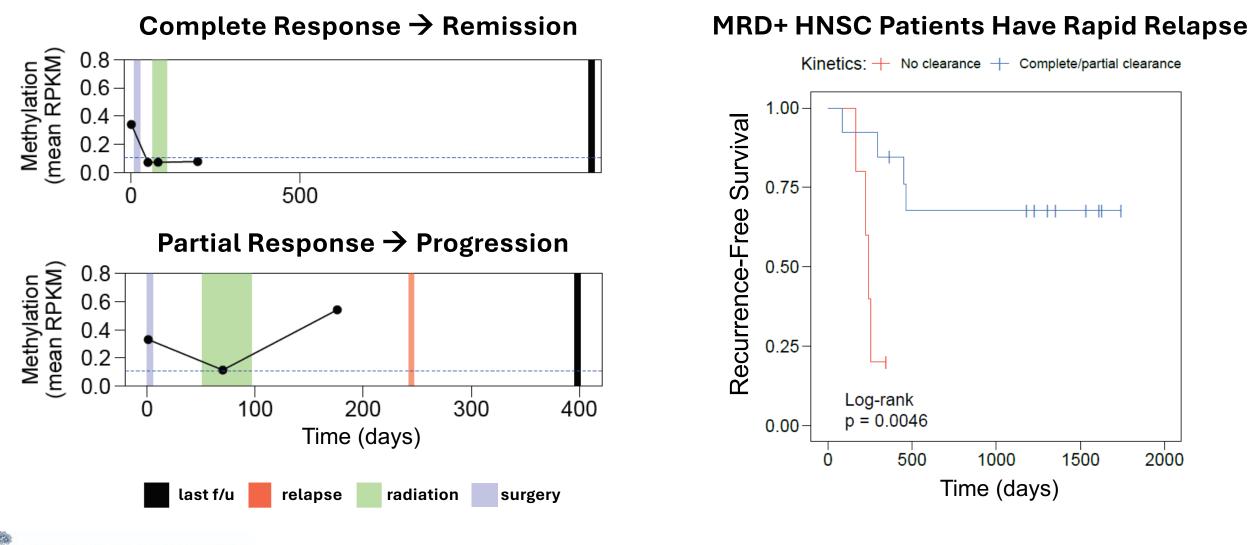
#### **Cancer Type**



Klein, et al. Annals Oncol 2021



## Methylated ctDNA for Tissue-agnostic MRD Detection



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Burgener, De Carvalho, Bratman, et al. Clin Cancer Res 2021



How could cell-free DNA *epigenetic* profiling potentially impact clinical decisions?

- Enable tissue-agnostic detection in screening or MRD settings
- Distinguish between <u>distinct cancer types</u> in unknown primary or early detection settings
- Uncover mechanisms of <u>therapeutic resistance</u> such as histologic transformation
- Reveal **tissue damage** that reflects treatment toxicity





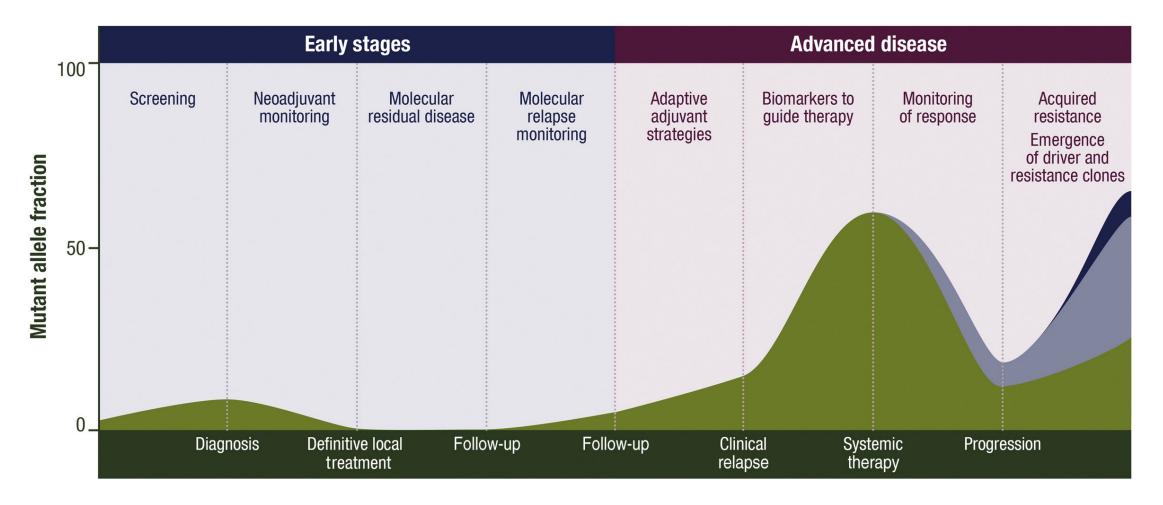
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### Could ctDNA have utility across the cancer care continuum?

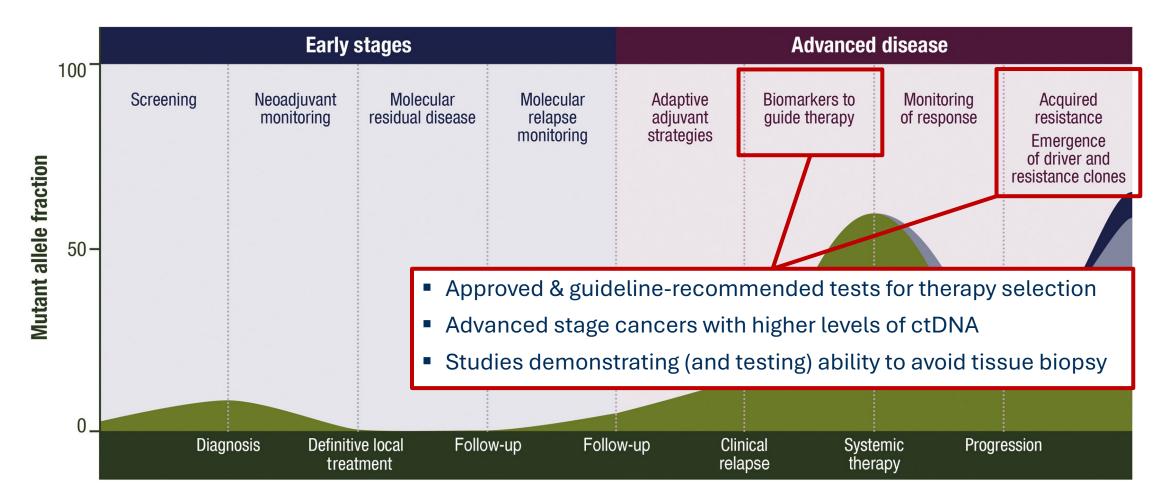




Pascual, Turner et al. Annals Oncol 2022



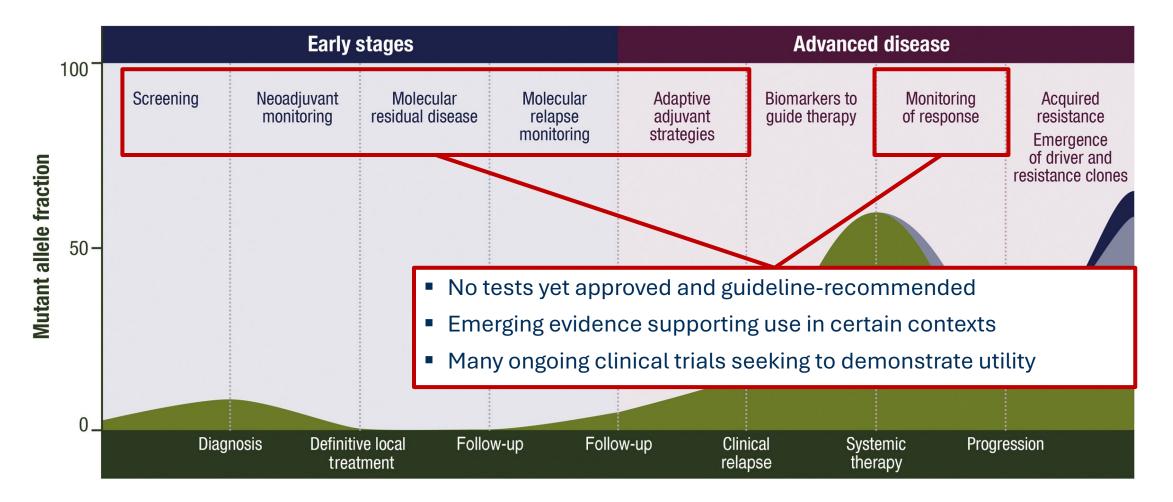
### Could ctDNA have utility across the cancer care continuum?







#### Could ctDNA have utility across the cancer care continuum?











#### **SPECIAL ARTICLE**

# ESMO recommendations on the use of circulating tumour DNA assays for patients with cancer: a report from the ESMO Precision Medicine Working Group

J. Pascual<sup>1</sup>, G. Attard<sup>2</sup>, F.-C. Bidard<sup>3,4</sup>, G. Curigliano<sup>5,6</sup>, L. De Mattos-Arruda<sup>7,8</sup>, M. Diehn<sup>9</sup>, A. Italiano<sup>10,11,12</sup>, J. Lindberg<sup>13</sup>, J. D. Merker<sup>14</sup>, C. Montagut<sup>15</sup>, N. Normanno<sup>16</sup>, K. Pantel<sup>17</sup>, G. Pentheroudakis<sup>18</sup>, S. Popat<sup>19,20</sup>, J. S. Reis-Filho<sup>21</sup>, J. Tie<sup>22,23</sup>, J. Seoane<sup>24,25</sup>, N. Tarazona<sup>26,27</sup>, T. Yoshino<sup>28</sup> & N. C. Turner<sup>19,20\*</sup>





### ESMO recommendations on the use of ctDNA

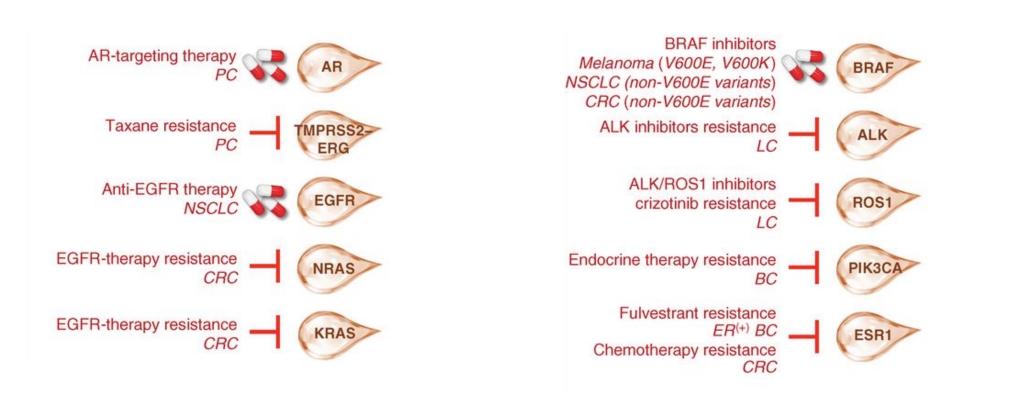
- For advanced cancers, validated and adequately sensitive ctDNA assays have utility in identifying actionable mutations to direct targeted therapy, and may be used in routine clinical practice, provided the limitations of the assays are taken into account
- For early-stage cancers, detection of MRD has high evidence of clinical validity in anticipating future relapse, but MRD detection cannot be recommended in routine clinical practice due to lack of clinical utility studies
- Additional potential applications of ctDNA assays are not recommended for routine practice



Pascual, Turner et al. Annals Oncol 2022



## Therapeutic targets and resistance mechanisms potentially detectable within ctDNA

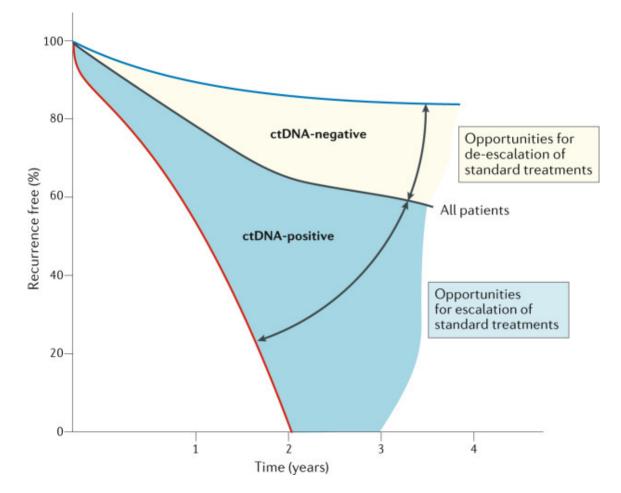


Alix-Panabières and Pantel, Cancer Discov. 2021;11(4):858-873.





# Potential applications of post-treatment ctDNA-based MRD in tailoring the aggressiveness of adjuvant therapy

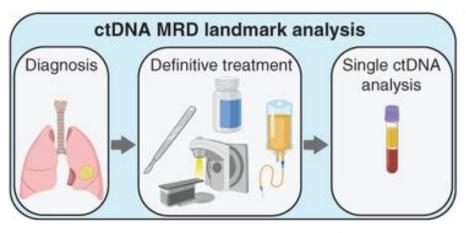


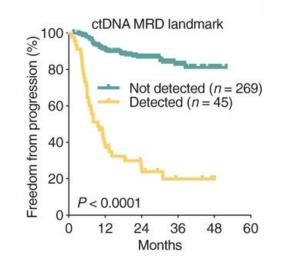


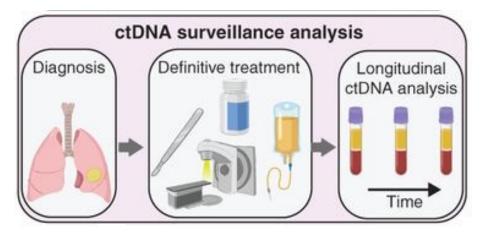
Dasari et al. Nat Rev Clin Oncol 2020

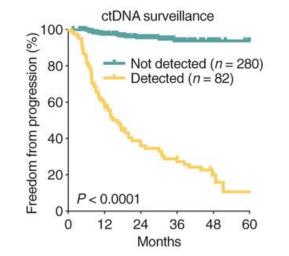


### Assay characteristics impact accuracy for MRD detection; Longitudinal ctDNA analysis boosts sensitivity & NPV









Moding, Diehn, et al. Cancer Discov. 2021;11(12):2968-2986.





Emerging clinical utility of ctDNA-based MRD-guided cancer therapy

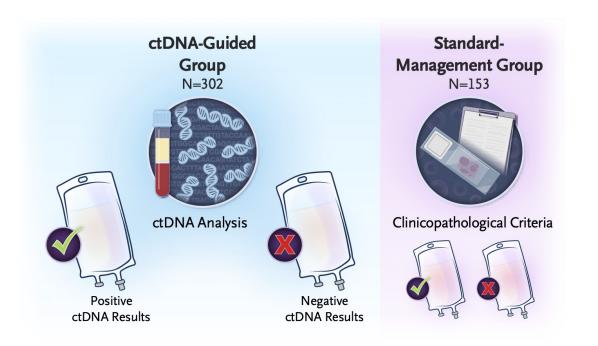
The NEW ENGLAND JOURNAL of MEDICINE						
ESTABLISHED IN 1812	JUNE 16, 2022	VOL. 386 NO. 24				
Circulating Tumor DNA Analysis Guiding Adjuvant Therapy in Stage II Colon Cancer						



Tie, et al. NEJM 2022. doi: 10.1056/NEJMoa2200075



## ctDNA-guided approach in stage II colon cancer reduced adjuvant chemotherapy use without compromising RFS

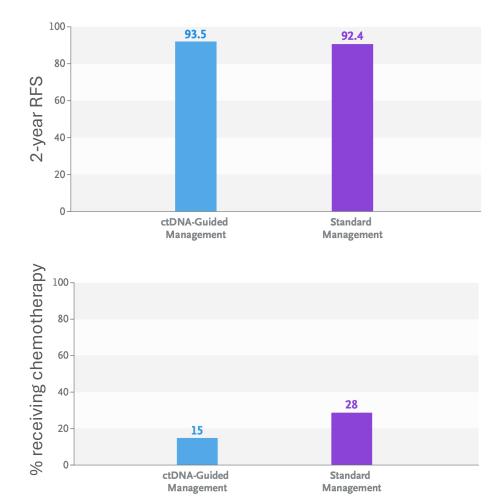


- Phase 2 RCT with noninferiority design
- Primary endpoint: 2-yr RFS

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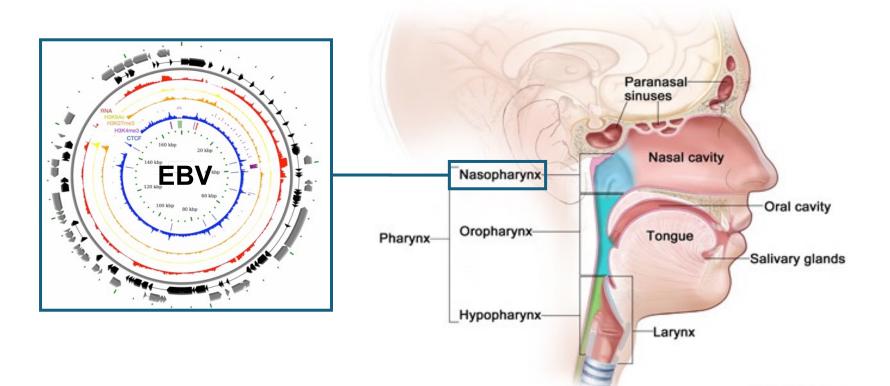
• Noninferiority margin: -8.5%



Tie, et al. NEJM 2022. doi: 10.1056/NEJMoa2200075



# Lessons from studies on plasma Epstein-Barr Virus (EBV) in endemic nasopharynx cancer patients







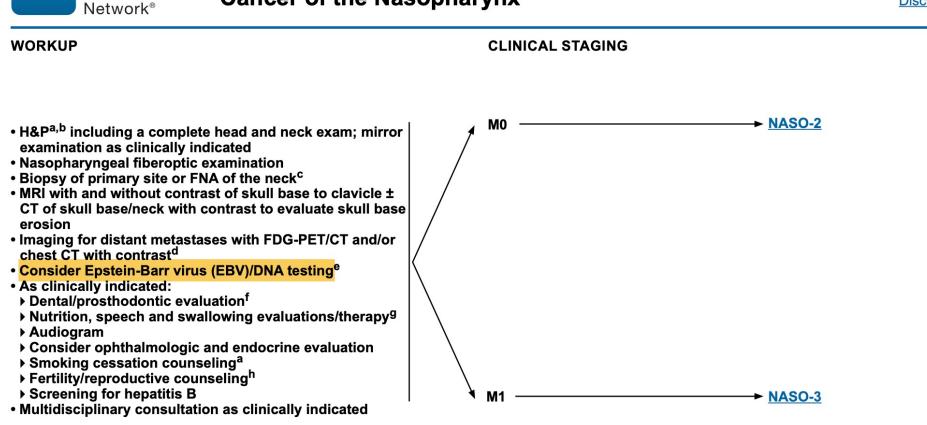
National Comprehensive N

NCCN Cancer

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### NCCN Guidelines Version 3.2024 Cancer of the Nasopharynx



<sup>a</sup> H&P should include documentation and quantification (pack years smoked) of tobacco use history, as well as alcohol use and counseling. All patients who currently smoke should be advised to quit smoking, and those who formerly smoked should be advised to remain abstinent from smoking. For additional cessation support, refer to the Patient/Provider Smoking Cessation and Treatment Resources in the <u>NCCN Guidelines for Smoking Cessation</u>.

<sup>b</sup> Screen for depression (NCCN Guidelines for Distress Management).

<sup>c</sup> Image-guided (US or CT) needle biopsy of cystic neck nodes may offer better diagnostic yield than FNA by palpation alone for initial diagnosis in this setting. For unresectable or metastatic disease where there is a plan for systemic therapy, a core biopsy would allow for ancillary immune-genomic testing.
<sup>d</sup> Principles of Imaging (IMG-A).

<sup>e</sup> For nonkeratinizing or undifferentiated histology, consider testing for EBV in tumor and blood. Common means for detecting EBV in pathologic specimens include ISH for EBV-encoded RNA (EBER) or immunohistochemical staining for latent membrane protein (LMP). The EBV DNA load within the serum or plasma may be quantified using PCR targeting genomic sequences of the EBV DNA such as BamHI-W, Epstein-Barr virus nuclear antigen (EBNA), or LMP; these tests vary in their sensitivity. The EBV DNA load may reflect prognosis and change in response to therapy.

<sup>f</sup> Principles of Dental Evaluation and Management (DENT-A).

- 9 Principles of Nutrition: Management and Supportive Care (NUTR-A).
- <sup>h</sup> See fertility and reproductive endocrine considerations in the <u>NCCN Guidelines for</u> <u>Adolescent and Young Adult (AYA) Oncology</u>.



## Plasma EBV DNA detectability and levels are prognostic in patients with EBV-associated nasopharynx cancer

Outcomes	Studies No.	Hazard radio, 95%CI	Z-value	P-value
Pre-DNA				
OS	13	2.81, 2.44-3.24	14.31	< 0.00001
PFS	6	2.74, 2.37-3.18	13.33	< 0.00001
DMFS	6	3.89, 3.39-4.47	19.27	< 0.00001
LRFS	6	2.02, 1.52-2.70	4.82	< 0.00001
Post-DNA				
OS	6	4.26, 3.26-5.57	10.58	< 0.00001
PFS	2	5.21, 3.29-8.27	7.01	< 0.00001
DMFS	2	7.54, 3.39-16.77	4.95	< 0.00001
LRFS	4	7.51, 5.11-11.02	10.30	< 0.00001
Mid-DNA				
OS	1	3.29, 1.37-7.89	2.67	0.0077
PFS	1	4.05, 1.89-8.67	3.60	0.0003
DMFS	1	12.02, 2.78-51.93	3.33	0.0009
LRFS	1	2.05, 0.79-5.31	1.48	0.1378
LKF5	1	2.05, 0.79-5.51	1.48	0.13/8

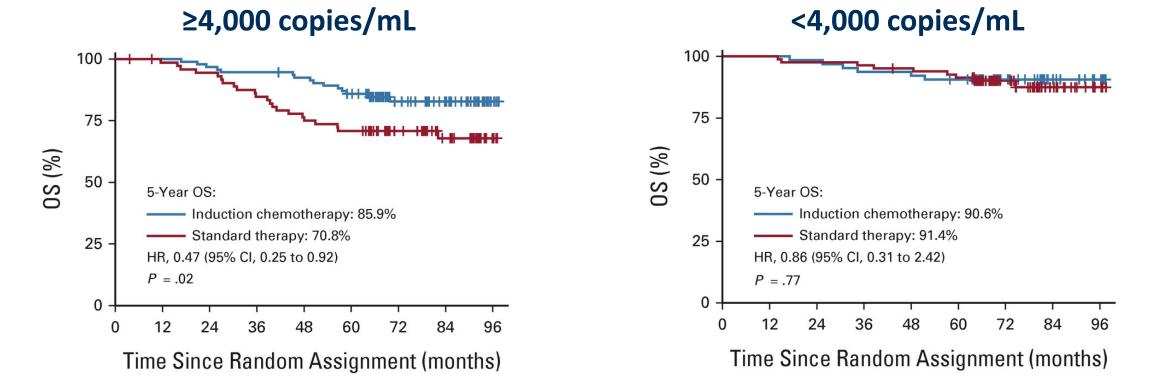
- EBV DNA levels in plasma pre-, mid-, and post-treatment are prognostic
- Stronger association with distant metastasis than locoregional recurrence
- Results become more strongly prognostic during and after treatment

Zhang W, Chen Y, Chen L, et al. Medicine (Baltimore) 2015;94:e845.





*Potential utility:* Benefit of neoadjuvant chemotherapy restricted to patients with high pre-treatment plasma EBV DNA levels



### This exploratory analysis of a phase 3 trial should be confirmed in future prospective studies

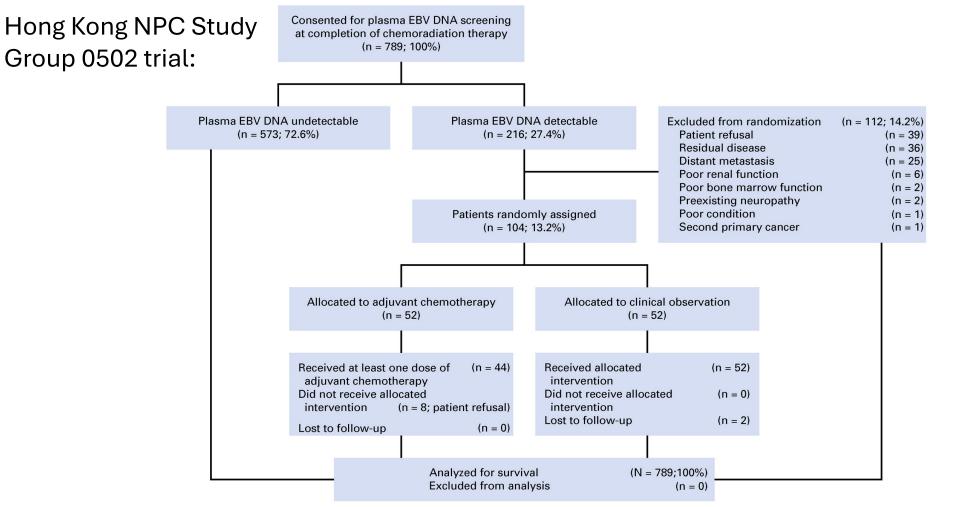
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Zhang Y, et al. J Clin Oncol 2022. 40(22):2420-2425



### A cautionary tale: Randomized trial testing utility of post-treatment detectable plasma EBV DNA to escalate adjuvant chemotherapy

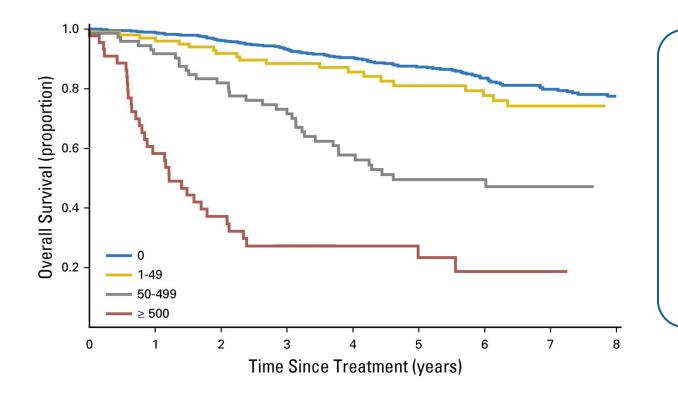


Chan ATC et al. J Clin Oncol 2018 doi: 10.1200/JCO.2018.77.7847





A cautionary tale: Randomized trial testing utility of post-treatment detectable plasma EBV DNA to escalate adjuvant chemotherapy



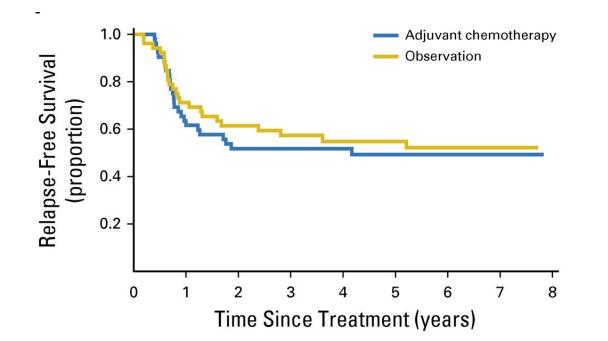
 Validated prognostic value of plasma EBV DNA following chemoRT

Chan ATC et al. J Clin Oncol 2018 doi: 10.1200/JCO.2018.77.7847





A cautionary tale: Randomized trial testing utility of post-treatment detectable plasma EBV DNA to escalate adjuvant chemotherapy



- Validated prognostic value of plasma EBV DNA following chemoRT
- No benefit to adjuvant chemotherapy in MRD+ patients following chemoRT
- Role of resistance to prior agents?
- Need new treatment options for MRD+ patients

#### NRG-HN001 randomized phase 2 study is current testing alternative adjuvant chemotherapy in MRD+ patients

Chan ATC et al. J Clin Oncol 2018 doi: 10.1200/JCO.2018.77.7847





Plasma EBV DNA surveillance detects distant but not locoregional recurrence with an average of ~6 months lead time

1.0 1.0 0.9 0.9 0.8 0.8 Proportion with recurrence Proportion with recurrence 0.7 0.7 0.6 0.6 0.5 0.5 0.4 0.4 0.3 0.3 0.2 0.2 EBV surveillance 0.1 0.1 0.0 0.0 0 100 200 300 400 500 600 0 100 200 300 400 500 600 700 700 800 900 1000 Days from completion of RT to the first sign of local failure Days from completion of RT to the first sign of distant failure

**Distant Recurrence** 

#### **Locoregional Recurrence**

Hong et al. Cancer; 2004, 100(7):1429-1437

1000

1100

FBV surveillance

900

800





### Take Home Messages

- ctDNA is increasingly utilized in clinical practice and trials, particularly for noninvasive genotyping of advanced disease
- Many early-stage cancers and post-treatment recurrences are below detection limits of available assays
- ctDNA testing can enable serially updated prognostication
- ctDNA-based MRD detection has potential clinical utility, but rigorous prospective studies are needed



