How Reliable is Circulating Tumor DNA in Detecting Disease Progression and Regression of Non-colorectal Gastrointestinal Cancers?

<u>Bipin Ghimire¹</u>, Ujjwal Karki¹, Emma Herrman¹, Samiksha Pandey¹, Mohammad Muhsin Chisti² 1- Beaumont Health - Royal Oak, MI, Internal Medicine, Royal Oak, MI, United States of America; 2 - Beaumont Health - Royal Oak, MI, Hematology/Oncology, Royal Oak, MI, United States of America

Introduction

- Circulating tumor DNA is short DNA sequence shed by tumor cells to the circulation [1]
- CtDNA has a vast possible applications: tumor molecular profiling, tracking treatment response, detection of resistance, and detection of minimal residual disease [1]
- > Literature is limited for non-colorectal GI cancers, with a few studies available for pancreatic, hepato-biliary and gastric cancers [2-6]
- > Dynamic ctDNA changes during treatment and detection of progression or regression has not been well described in studies [2-6]

Methodology

- Study design and setting:
- Retrospective observational study of 18 patients with noncolorectal GI cancers at William Beaumont Hospital, MI

> Inclusion/Exclusion criteria:

• Included patients ≥ 18 years of age. Excluded patients without corresponding imaging to compare

Variables:

- Baseline characteristics: Demographics, BMI, tobacco/alcohol use, family history, stage of disease, treatment received
- Variables of interest:
 - Disease progression: increased size of known cancerous lesion or development of new lesion, noted in imaging
 - Disease regression: decreased or resolution size of known lesion
 - Presence of disease: Significant burden of disease noted on imaging
 - Absence of disease: no cancerous lesions on imaging

Results

Charao Age Sex Patien

> BMI Tobace Alcoho Family

> > Type of

Stage

Treatn

Methodology

> Statistical Analysis:

• With single ctDNA: Correlation of single ctDNA results with imaging to predict presence of disease

• With serial ctDNA: Analysis of pairs of consecutive ctDNA trend (either up-trending or down- trending or negative persistently) and correlation with imaging to predict disease

progression/regression

• Calculation of sensitivity, specificity, PPV, NPV for analyses of both single and serial ctDNA

Baseline characteristics of participants

Frequency
64 (31, 80)
Male 50% (9/18)
Caucasian 66.7% (12/18)
AA 16.7% (3/18)
Others 16.7% (3/18)
27 (20, 35)
77.8% (14/18)
16.7% (3/18)
0% (0/19)
Hepato-biliary carcinoma - 33.3% (6/18)
Pancreatic adenocarcinoma - 27.8% (5/18)
Anal squamous cell carcinoma - 11.1% (2/18)
Neuroendocrine tumor - 11.1% (2/18)
Gastric adenocarcinoma - 5.6% (1/18)
Small bowel adenocarcinoma - 5.6% (1/18)
GI cancer of unknown origin - 5.6% (1/18)
Stage I - 5.6% (1/18)
Stage II - 22.2% (4/18)
Stage III - 33.3% (6/18)
Stage IV - 38.9% (7/18)
Chemotherapy - 83.3% (15/18)
Surgery - 55.5% (10/18)
Targeted therapy - 44.4% (8/18)
Immunotherapy - 33.3% (6/18)
Radiation - 16.7% (3/18)

Results

> Analysis with single ctDNA: predicts presence of disease

CtDNA results (Single values)	Imaging finding		
	Presence of disease	Absence of disease	Total
Positive	12	0	12
Negative	8	13	21
Total	20	13	33

✓ Finding: Sensitivity - 60% ; Specificity - 100%; PPV - 100%; **NPV – 61.9%**

Analysis with serial ctDNA

□ All ctDNA values and disease trend

CtDNA trend (Pairs)	Imaging finding				
	Disease progression	Disease regression	Stable disease	Absence of disease	Total
Up trending	4	0	1	0	5
Down trending	0	4	0	0	4
Persistent Negative	0	0	1	5	6
Total	4	4	2	5	15

Up-trending ctDNA analysis: predicts disease progression

CtDNA trend	Imaging finding		
	Disease progression	Other than progression	Total
Up-trending	4	1	5
Non up-trending	0	10	10
Total	4	11	15

NPV – 100%

✓ Finding: Sensitivity – 100%; Specificity – 90.9%, PPV – 80%;

Results

Down trending ctDNA analysis: predicts disease regression

CtDNA trend	Imaging finding			
	Disease regression	Other than regression	Total	
Down-trending	4	0	14	
Non down-trending	0	11	11	
Total	4	11	15	

✓ Finding: Sensitivity- 100%; Specificity – 100%; PPV – 100%; **NPV – 100%**

> Median Lead time: Earlier detection of progression by ctDNA compared to imaging: 44 days

Discussion and Conclusion

- > We describe good sensitivity, specificity, PPV and NPV of serial ctDNA to detect either disease progression or regression. But lower than our separate analysis of colorectal cancers.
- > Above test results, and lead time of 44 days can assist physicians to make/change treatment plans prior to the imaging, and can reduce radiation exposure
- > Our sample size was small and we recommend larger prospective studies are required to describe impact of ctDNA – guided surveillance in clinical outcomes

References

- Corcoran RB, Chabner BA. Application of Cell-free DNA Analysis to Cancer Treatment. N Engl J Med. 2018 Nov;379(18):1754-65
- Azad TD, Chaudhuri AA, Fang P, et al. Circulating Tumor DNA Analysis for Detection of Minimal Residual Disease After Chemoradiotherapy for Localized Esophageal Cancer. Gastroenterology. 2020 Feb;158(3):494-505.e6
- Term Follow-Up Patients with Hepatocellular Carcinoma. Clinical Cancer Research. 2019 Sep 3;25(17):5284–94
- tumor DNA (ctDNA) and prognosis in pancreatic cancer. Critical Reviews in Oncology/Hematology. 2021 Dec;168:103528
- Lapin M, Huang HJ, Chagani S, et al. Monitoring of Dynamic Changes and Clonal Evolution in Circulating Tumor DNA From Patients With IDH-Mutated Cholangiocarcinoma Treated With Isocitrate Dehydrogenase Inhibitors. JCO Precis Oncol. 2022 Feb;6:e2100197
- Yang J, Gong Y, Lam VK, et al. Deep sequencing of circulating tumor DNA detects molecular residual disease and predicts recurrence in gastric cancer. Cell Death Dis. 2020 May 11;11(5):1–9



Beaumont

WR School of MEDICINE

Cai Z, Chen G, Zeng Y, et al. Comprehensive Liquid Profiling of Circulating Tumor DNA and Protein Biomarkers in Long-Guven DC, Sahin TK, Yildirim HC, et al. A systematic review and meta-analysis of the association between circulating