# Multimodal circulating tumor DNA blood-based colorectal cancer screening test demonstrates clinically meaningful sensitivity across multiple clinical parameters

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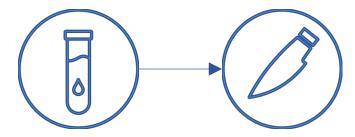
#### Introduction

- Blood-based colorectal cancer (CRC) screening holds potential to improve screening compliance due to:
  - Ability to seamlessly integrate into standard of care clinical pathways
  - High patient and provider acceptability
- A blood-based screening test must detect CRC across multiple clinical parameters in order to prove clinically meaningful in a screening population
- We aimed to describe the performance of a multimodal ctDNA blood-based CRC screening test in a cohort of patients with newly diagnosed CRC

## Cohort Description

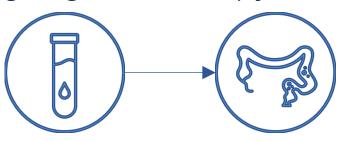
#### Validation cohort:

Cases: Consented individuals with newly diagnosed CRC (N = 699)



- Whole blood collected prior to surgical resection
- Analysis restricted to Stage I III CRC diagnoses
- Median age: 63 years (20-89)

Controls: Consented individuals undergoing colonoscopy (N = 297)



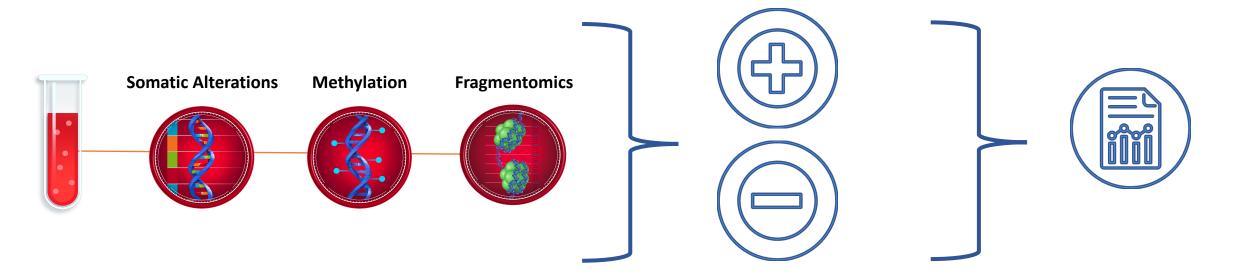
- Whole blood collected prior to colonoscopy procedure
- Analysis restricted to those confirmed negative for advanced colorectal neoplasia
- Median age: 57 years (20 91)

#### **Training cohort:**

Separate cohort of individuals with CRC (N = 850) and colonoscopy confirmed negative controls (N = 541)



# Blood-based CRC detection cfDNA analysis



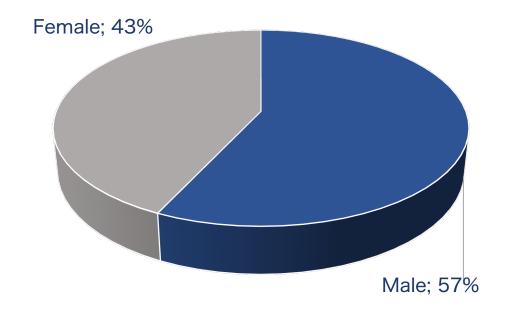
Plasma analyzed on a 500kb NGS based panel and bioinformatic platform incorporating cfDNA methylation-based partitioning to identify cancer related genomic alterations and epigenomic modifications

The output is a "ctDNA detected" or "ctDNA not detected" result

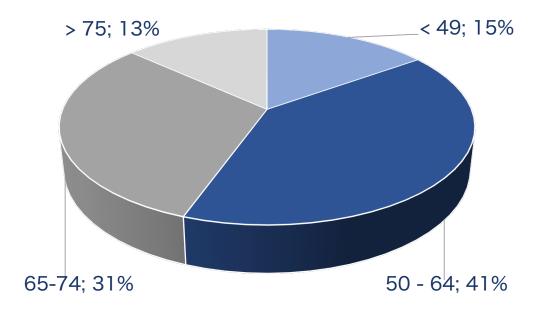
Final results were correlated with key clinical parameters known to influence clinical decision making in CRC

### Case Cohort Demographics

Biological Sex



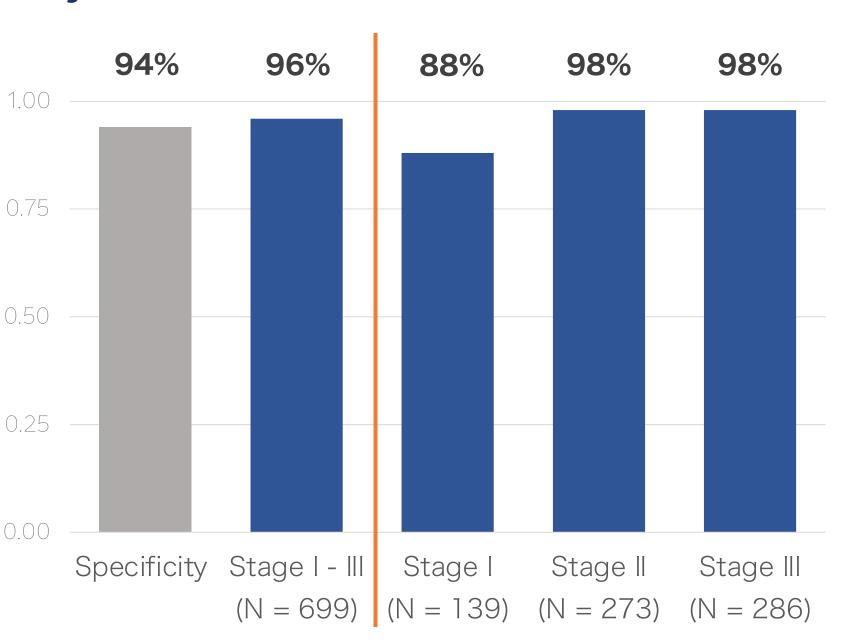
Age at Cancer Diagnosis (in years)



## Overall sensitivity for CRC detection: 96%

In this cohort of individuals with CRC, overall sensitivity was 96% at 94% specificity 0.75

Across stage I, II, and III disease, observed sensitivity was clinically meaningful, on par with currently available non-invasive screening modalities (i.e. stoolbased screening)



Clinically meaningful sensitivity across multiple

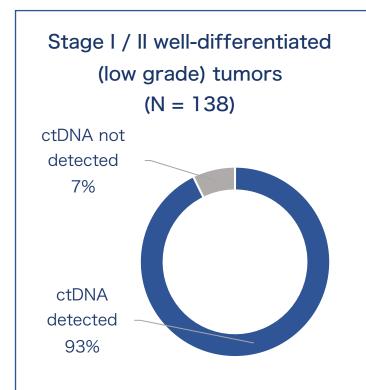
clinical parameters

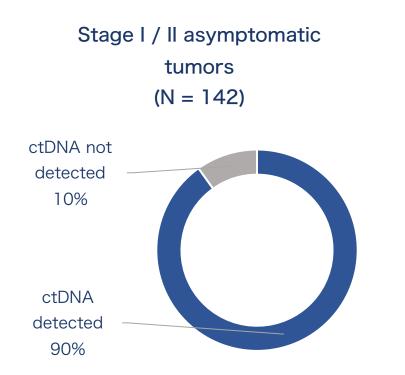
- The assay demonstrated clinically meaningful sensitivity across
  - Cancer presentation
  - Primary tumor location
  - Histology
  - MSI status
  - Tumor grade
  - Degree of tumor invasion
- While sensitivity was greater than 90% in each group, significant performance differences were observed in
  - CRC presentation
  - Presence of perineural invasion (prognostic indicator)

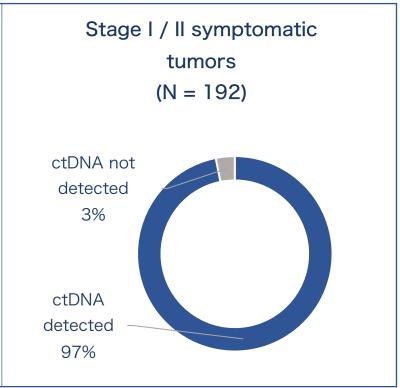
	Clinical Variable		Number of individuals	% ctDNA detected	
	Symptomatic CRC presentation	Asymptomatic Symptomatic Unknown	256 350 93	93 97 99	p< 0.05*
	Primary Tumor Location	Right Colon Left Colon Transverse Colon	151 511 37	96 96 95	ns <sup>+</sup>
	Histology	Adenocarcinoma Mucinous adenocarcinoma	666 32	96 100	ns*
		Mucinous adenocarcinoma with signet ring features	1	100	
	MSI Status	MSI-High MSI-Low / MSS Unknown	55 628 16	91 97 94	ns*
		Well-Differentiated (low grade)	175	93	
	Tumor Grade	Moderately Differentiated (intermediate grade)	462	97	ns <sup>+</sup>
		Poorly Differentiated (high grade) Unknown	32 30	100 100	
	Lymphatic Invasion	Present Not Identified	67 632	100 96	ns*
	Venous Invasion	Present Not identified	251 448	97 96	ns*
	Perineural invasion	Present Not identified	317 382	98 94	p < 0.05*

# Potential application to CRC screening

Given this is a cohort of individuals with known CRC, completed additional analysis in those with Stage I / II CRC to understand applicability to a screening population







# Strengths and Limitations

#### Strengths:

- Large cohort with extensive clinical data
- Allows for informative subanalyses across tumor subtypes, especially in earlystage cancers
  - 59% of cohort had Stage I or II disease
  - 37% presented with asymptomatic disease

#### Limitations:

- Cohorts not reflective of intended use screening population
  - Cohort bias may lead to differential assay performance
  - Ethnicity was not matched for cases and controls in neither training nor testing cohorts which may impact assay results
  - Ongoing studies to evaluate performance in an average risk screening population
- Retrospective analysis of banked plasma samples collected > 5 years ago

#### Conclusions

- This multimodal ctDNA CRC screening test has <u>clinically</u> <u>meaningful sensitivity</u> across multiple clinical parameters, most notably in those with <u>early-stage asymptomatic disease and early-stage low-grade tumors</u>
- The data suggest this test may have <u>clinically meaningful</u>
   <u>performance</u> in an <u>average risk screening population</u> presenting
   with varying cancer stages and tumor histologic features
- Future studies aim to <u>validate this test</u> in a screen-relevant population
- A <u>prospective registrational study</u> to evaluate the test in an average risk, screening cohort is ongoing (NCT04136002)

# Thank you

 Patients, Healthy Individuals, and their families who participated in this research

Clinical and Research Teams at Samsung Medical Center

Guardant Health Collaborators

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