Multimodal circulating tumor DNA (ctDNA) colorectal neoplasia detection assay for asymptomatic and early-stage colorectal cancer

Hee Cheol Kim¹, Seung Tae Kim², Yupeng He³, Paul Sample³, Yoshiaki Nakamura⁴, Victoria M. Raymond³, Ariel Jaimovich³, AmirAli Talasaz³, and Jeeyun Lee² ¹Department of Surgery, Samsung Medical Center Sungkyunkwan University School of Medicine, Seoul South Korea, ²Division of Hematology-Oncology, Department of Medicine, Samsung Medical Center Sungkyunkwan University School of Medicine Seoul South Korea, ³Guardant Health, Redwood City, CA USA, ⁴Department of Gastroenterology and Gastrointestinal Oncology, National Cancer Center Hospital East, Chiba, Japan

Background

- National colorectal cancer (CRC) screening rates have remained largely unchanged and well below rates recommended by leading public health organizations, despite numerous available screening options.
- CRC screening options that address patient and provider reported compliance barriers (e.g., time and convenience) are needed to improve screening compliance.
- We evaluated the LUNAR-2 blood-based colorectal neoplasia detection test (Figure 1) in a large cohort of patients with newly diagnosed CRC

Figure 1: The LUNAR-2 test is a multimodal blood-based colorectal neoplasia detection assay incorporating ctDNA (circulating tumor DNA) assessment of **somatic mutations** and tumor derived methylation and fragmentomic patterns, aimed to maximize sensitivity for early-stage CRC detection



Methods

- 434 individuals diagnosed with stage I, II, or III CRC between 2013 and 2016 consented to provide blood samples prior to surgical resection (Table 1)
 - Those treated with neoadjuvant chemotherapy were excluded
 - Blood was collected in EDTA tubes and processed to plasma (median 3mL)
- Control samples were collected from age-matched CRC–free individuals (Figure 2)
- Isolated plasma samples were analyzed with LUNAR-2 (Guardant Health, USA)
- A held-out training set of samples from 614 individuals (CRC or CRC-free) was used to generate a model with calling thresholds targeting 90% specificity
- Final "ctDNA detected" or "ctDNA not detected" results were generated using this separately trained model
- ctDNA results and clinical characteristics were correlated







- Overall CRC sensitivity was **91%** with high sensitivity across stages I, II, and III CRC and 94% specificity (blue bars)
- No difference in sensitivity was observed when excluding those with early (<45 years) or late (>84 years) onset CRC (p=0.95; orange bars)

There was no difference in sensitivity between asymptomatic as compared to symptomatic presentation of CRC (Panel A; p=0.4). However, lower cell-free DNA tumor fractions were observed in the asymptomatic cohort especially in stage I and stage II CRC (Panel B)

88%

Conclusions

In this large early-stage CRC cohort, multimodal ctDNA assessment has *high sensitivity for CRC* detection with high specificity.

Equivalent sensitivity in the asymptomatic cohort suggests this test will have *clinically meaningful* performance in an average-risk screening population.

No difference in CRC sensitivity was observed when focusing on the *aged* 45 – 84 cohort.

A prospective registrational study is ongoing to evaluate the test in an average-risk CRC screening cohort.

SAMSUNG

SAMSUNG MEDICAL CENTER



Results





At 94% specificity, sensitivity was maintained across a multi-cohort analysis