Longitudinal clinical performance of a novel tumor-naïve minimal residual disease assay in resected stage II and III colorectal cancer patients: A subset analysis from the GALAXY study in CIRCULATE-Japan

Yoshiaki Nakamura¹, Kristiyana Kaneva², Christine Lo², Farahnaz Islam², Seung Won Hyun², Daisuke Kotani¹, Kate Sasser², Halla Nimeiri², Eiji Okl³, Takayuki Yoshino¹ ¹Department of Gastroenterology and Gastrointestinal Oncology, National Cancer Center Hospital East, Kashiwa Japan; ²Tempus AI, Inc., Chicago, IL; ³Department of Surgery and Science, Graduate School of Medical Sciences, Kyushu University, Fukuoka, Japan Published Abstract Number: 3618

INTRODUCTION

The presence of circulating tumor DNA (ctDNA) can identify patients at higher risk of disease recurrence. Longitudinal surveillance of ctDNA may enable early identification of patients who are likely to relapse and presents a window of opportunity for early interventions to improve outcomes. A tumor-naïve plasma-only approach for minimal residual disease (MRD) assessment accelerates turnaround time, enabling rapid adjuvant chemotherapy (ACT) treatment decisions and ongoing molecular surveillance.

METHODS

Patients with pre-specified eligibility criteria (n=80) were randomly selected from GALAXY in the CIRCULATE-Japan study. Patients were selected for recurrence to 50% while maintaining the stage II:III recurrent/non-recurrent ratio observed in GALAXY. There were 69 evaluable patients for landmark timepoint (LMT) analysis and 74 for longitudinal analysis. LMT was defined as 4 weeks after surgery in pathological stage II or III colorectal cancer (CRC).

Residual plasma samples were analyzed with the Tempus xM MRD assay (xM), a tumor-naïve ctDNA assay for MRD that integrates methylation and genomic variant data to deliver a binary MRD call. Calls were blinded to clinical outcomes.

Longitudinal sensitivity and specificity were assessed based on the LMT sample and all the evaluable longitudinal samples (every 3 months after surgery until recurrence, death, or 24 months follow-up was reached, whichever occurred first).

All blood samples, including those at LMT, were analyzed using an improved analytical pipeline performance relative to previously presented work*. Longitudinal samples (post LMT) were analyzed using the methylation pipeline only.

Figure 1. Methyl and variant pipelines



Figure 1. Dual workflow was used at LMT. The methylation workflow alone was used for all subsequent timepoints.

SUMMARY

•xM is a rapid, tumor-naïve MRD assay that demonstrates robust clinical surveillance performance with longitudinal clinical sensitivity of 83.3% (increased from 61% at LMT) and longitudinal specificity of 89.5%. •xM ctDNA status is a stronger prognostic biomarker to DFS compared to standard of care CEA (Adj. HR 9.69 vs. 2.13). •xM has an overall mean lead time of 4.66 months & a lead time of 5.62 months for surgery-only patients prior to recurrence.

RESULTS



Figure 2. Swimmer plot of recurrent patients (n=41) showing follow-up time, recurrence status, timing of ACT and xM MRD status at baseline, LMT, and longitudinal timepoints. Recurrence includes death events.

Table 1. Distribution of lead time (time from first MRD+ call to date of recurrence or death) for TP (n=30) patients. Overall mean lead time defined from first MRD+ to recurrence is 4.66 months. For patients with surgery only treatment, the mean lead time is 5.62 months.

Figure 3. Clinical sensitivity and specificity 2/ ____ 50% **b** 25%



True Negative (TN) False Positive (FP) Invalid assav

Figure 3. Sensitivity at landmark (61.1%) and longitudinal (83.3%) timepoints in recurrent patients (left). Specificity at landmark (87.9%) and longitudinal (89.5%) timepoints in non-recurrent patients (right). Arrows show patients who switched MRD status between landmark and longitudinal

Table 1. First MRD+ lead time to recurrence or death

	Total (n=30)	Surgery Only (n=22)
Minimum	0.00	0.02
1st Quantile	1.00	2.14
Median	4.77	5.30
Mean	4.66	5.62
3rd Quantile	6.22	6.45
Maximum	22.97	22.97

Table 2. Clinical performance by pathological stage

Landmark Sensitivity Specificity Adj PPV* Adj NPV* Adj HR* (MRD+/MRD-)

Longitudinal

Sensitivity Specificity Adj PPV* Adj NPV* Adj HR* (MRD+/MRD-)

Table 2. Clinical landmark performance (top). Clinical longitudinal performance (bottom). Adj PPV*, Adj NPV*, and Adj HR* are the estimates based on the anticipated true recurrence rate of 24% observed in GALAXY.

Figure 4. 12-week CEA & MRD status & Disease-Free Survival



Figure 4. Adjusted hazard ratio (HR) for xM MRD is nearly 5-fold higher compared to carcinoembryonic antigen (CEA) testing at 12 weeks post surgery. Adjusted HR* is the hazard ratio adjusted by anticipated true recurrence rate (24%). The adjusted median DFS time for MRD+ is 25.1 weeks (6.3 months) vs. not reached within 72 weeks (18 months) for MRD-.

Overall	Stage II	Stage III
61.1% (43.5%, 76.9%)	64.3% (35.1%, 87.2%)	59.1% (36.4%, 79.3%)
87.9% (71.8%, 96.6%)	93.3% (68.1%, 99.8%)	83.3% (58.6%, 96.4%)
61.4%	75.3%	52.8%
87.7%	89.2%	86.6%
7.28	12.25	5.21
Overall	Stage II	Stage III
Overall 83.3% (67.2%, 93.6%)	Stage II 91.7% (61.5%, 99.8%)	Stage III 79.2% (57.8%, 92.9%)
Overall 83.3% (67.2%, 93.6%) 89.5% (75.2%, 97.1%)	Stage II 91.7% (61.5%, 99.8%) 88.2% (63.6%, 98.5%)	Stage III 79.2% (57.8%, 92.9%) 90.5% (69.6%, 98.8%)
Overall 83.3% (67.2%, 93.6%) 89.5% (75.2%, 97.1%) 71.4%	Stage II 91.7% (61.5%, 99.8%) 88.2% (63.6%, 98.5%) 71.1%	Stage III 79.2% (57.8%, 92.9%) 90.5% (69.6%, 98.8%) 72.4%
Overall 83.3% (67.2%, 93.6%) 89.5% (75.2%, 97.1%) 71.4% 94.4%	Stage II 91.7% (61.5%, 99.8%) 88.2% (63.6%, 98.5%) 71.1% 97.1%	Stage III 79.2% (57.8%, 92.9%) 90.5% (69.6%, 98.8%) 72.4% 93.2%

Correspondence Yoshiaki Nakamura: yoshinak@east.ncc.go.jp Kristiyana Kaneva: kristiyana.kaneva@tempus.com