

The non-invasive Tempus xF liquid biopsy assay detects cell-free DNA (cfDNA) in blood specimens of advanced solid tumor patients. The assay is capable of detecting mutations in two variant classes in ~105 genes, including: Single Nucleotide Variants (SNVs) and insertions and deletions (indels), as well as Copy Number Amplifications (CNAs) in 6 genes, Copy Number Deletions (CNDs) in 2 genes\*, and gene rearrangements (translocations) in 7 genes spanning ~0.3 Mb of genomic space. The assay spans clinically relevant coding exons for 35 genes and covers recurrent hotspot mutations in 70 genes. Insertions and deletions will be reported down to the lower limit of detection (LLOD) in clinically relevant regions in 97 genes (list available upon request). Microsatellite Instability High (MSI-H) status is also reported when detected. The panel is designed to provide clinical decision support for patients with solid tumors and is focused on the identification of oncologic, including resistance, mutations.

CAP/CLIA validation of the Tempus xF panel focused on the detection of actionable oncologic and resistance variants in blood plasma. The assay requires two 8.5 mL Streck tubes of peripheral blood. Clinical sequencing is performed to ~20,000x coverage (at least 5,000x deduplicated reads). Performance specifications are listed in Table 1 below. These results establish, as shown in the table, high sensitivity and specificity for the Tempus xF assay.

Not indicated for:

- Hematologic malignancies
- Early stage (stage I/II) cancers
- Brain cancers
- Sarcomas

**TABLE 1: PERFORMANCE SPECIFICATIONS**

Variant Class	Variant Allele Fraction (VAF) (%)	Sensitivity (%)	Specificity (%)
Single Nucleotide Variants (SNVs)	>0.50	>99.9	>99.9
	0.50	>99.9	
	0.25	97.0	
	0.10	70.4	
Insertions and Deletions	>0.50	98.8	>99.9
	0.50	96.0	
	0.25	81.0	
Copy Number Amplifications (CNAs)	>0.50	>99.9	96.2
	0.50	>99.9	
Rearrangements/Fusions	>0.50	97.4	>99.9
	0.50	70.8	
Microsatellite Instability High (MSI-H) Status	N/A	37.5	>99.9

\*BRCA1 and BRCA2 copy number loss are reported when detected