

Sensitive Detection of 5% Somatic ctDNA Variants Using MagPurix® cfDNA Extraction in a Liquid Biopsy NGS workflow

Executive Summary

This study evaluated a complete liquid biopsy pipeline for detecting somatic variants at 5% variant allele frequency (VAF) in circulating tumor DNA (ctDNA) using plasma cfDNA. A key component of the workflow was automated extraction with the **MagPurix® 12 System** and **MagPurix® CFC DNA Extraction Kit (ZP02017)**, starting from plasma collected in Streck Cell-Free DNA BCT tubes.

MagPurix® provided sufficient cfDNA yield and the expected fragment size profile (120–220 bp peak) to support an optimized NGS protocol (Twist cfDNA library prep + Illumina MiSeq) and a dedicated bioinformatics pipeline. Using 0% and 5% reference standards, the combined NGS + confirmatory SNaPshot workflow achieved 94.12% sensitivity and 99% specificity for detecting clinically relevant somatic variants at 5% VAF, demonstrating that MagPurix®-extracted cfDNA is highly suitable for sensitive liquid biopsy applications.

Key Findings

- **cfDNA yield compatible with demanding NGS workflows**
 - Duplicate MagPurix® extractions from patient samples yielded, for example, ~46–56 ng cfDNA in 45 µL for one case (RX84.2023) and ~10–12 ng for a lower-yield case (RX87.2023), both adequate for Twist cfDNA library preparation (\geq 10–30 ng input).
- **High-quality cfDNA fragment profile for ctDNA detection**
 - TapeStation analysis of MagPurix®-extracted cfDNA showed the expected cfDNA fragment pattern with a main peak between 120–220 bp and an additional peak around 300 bp, matching the theoretical cfDNA distribution.
- **Enabling sensitive detection of 5% somatic variants**
 - Using MagPurix®-extracted cfDNA, the optimized NGS pipeline (Twist library prep with UMIs, 300 \times target coverage, and custom “Unbalance” variant-calling) successfully detected somatic variants present at 5% VAF in a 5% cfDNA reference standard.
 - When two independent somatic runs were combined, the method showed only 3 false negatives out of 51 variants, corresponding to 94.12% sensitivity.
- **High diagnostic reliability when coupled with orthogonal confirmation**
 - Initial NGS analysis displayed good but imperfect specificity; however, by confirming clinically relevant variants with a targeted SNaPshot assay, the authors report an overall specificity of 99% for the workflow.

Conclusion

Within this validated liquid biopsy workflow, the MagPurix® 12 System and MagPurix® CFC DNA Extraction Kit provided high-quality cfDNA from plasma, with yields and fragment profiles fully compatible with advanced NGS and ctDNA variant detection. The complete process, built

on MagPurix® extraction, enabled detection of 5% VAF somatic variants with 94.12% sensitivity and 99% specificity after orthogonal confirmation.

For clinical and research laboratories developing liquid biopsy assays, MagPurix® offers a reliable, automated, and sample-efficient extraction platform that supports sensitive detection of low-frequency tumor variants and helps ensure confidence in downstream molecular findings.

Reference

Mareso, C., Crosta, L., De Vita, M. G., Cristofoli, F., Tanzi, B., Benedetti, S., et al. (2024). Assessing the efficacy of an innovative diagnostic method for identifying 5% variants in somatic ctDNA. *Gene*, 928, 148771. <https://doi.org/10.1016/j.gene.2024.148771>