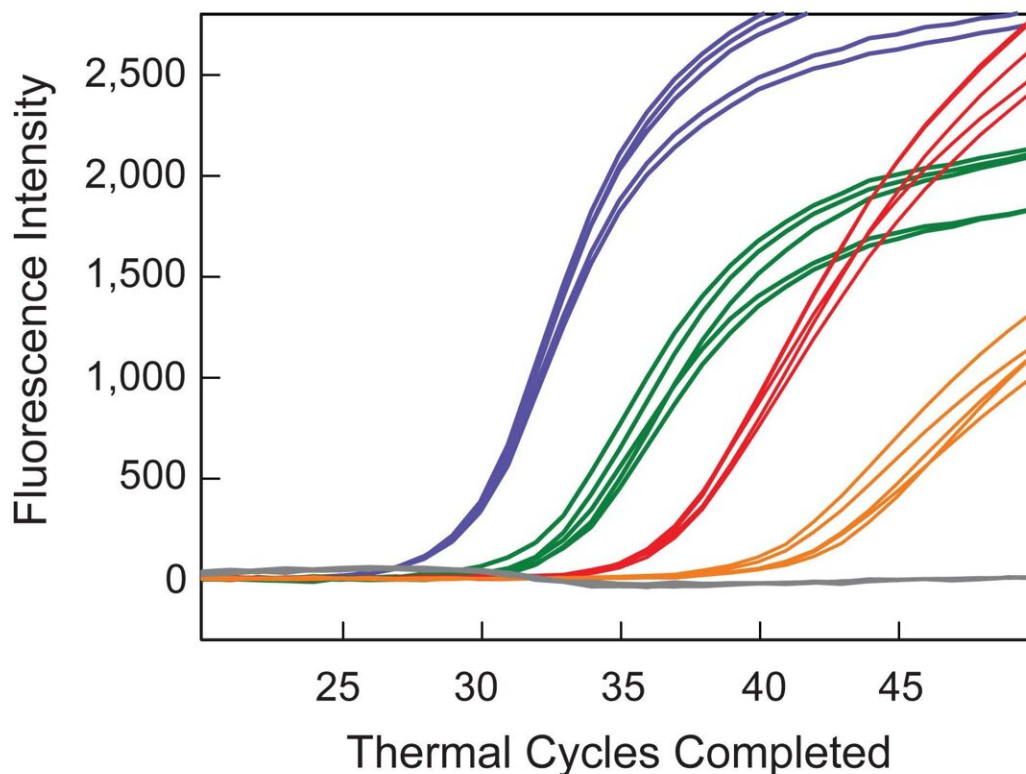


New assay shows promise for advancing personalized cancer treatment

February 11 2022



Demonstration of the sensitivity, selectivity, and repeatability of SuperSelective PCR assays that contain DNA fragments from 10,000 copies of human genomic DNA, in which β -actin serves as the wild-type reference gene (blue), and that also contain the following DNA fragments: 1,000 EGFR T790M point mutations (green); 100 EGFR Δ E745-A750 deletion mutations (red); 10 EGFR G719C point mutations (orange); and no EGFR L858R point mutations (gray). Credit: *Journal of Molecular Diagnostics*

Using a routine blood sample, investigators have developed an extremely sensitive assay to detect and quantify DNA fragments containing rare genetic mutations in a patient's cancer cells. This assay uses "SuperSelective" PCR primers that virtually ignore abundant closely related non-mutant DNA fragments derived from patients' normal cells. This new assay is faster and considerably cheaper than current methods and can be performed on widely available instruments. The investigators demonstrate the ease of utility of this approach in a new report in the *Journal of Molecular Diagnostics*.

"The long-term goal of this research is to develop highly multiplex assays that enable both the early detection of [mutations](#) relevant to cancer before any symptoms have occurred and the application of an effective early treatment. These assays will use noninvasive liquid biopsies (routine blood samples) taken during a person's annual medical checkup, and the DNA fragments transiently present in the [blood sample](#) (originating from cells that have died) will then be analyzed by highly multiplex real-time or digital PCR techniques," explained lead investigators Diana Yaneth Vargas, MS, and Fred Russell Kramer, Ph.D., from the Public Health Research Institute of the New Jersey Medical School of Rutgers University, Newark, NJ, U.S.

Current techniques for analyzing liquid biopsies involve either next-generation sequence analysis, which is expensive, lengthy, and has limited sensitivity; or use a PCR technique in which both the mutant DNA fragments and the closely related wild-type DNA fragments are amplified, limiting sensitivity when the mutant DNA fragments are rare. The unique design of SuperSelective PCR primers, on the other hand, enables the amplification of the mutant DNA fragments, while effectively preventing the amplification of the abundant closely related wild-type DNA fragments. Consequently, the SuperSelective PCR assays

described in this report are very sensitive.

Multiplex SuperSelective PCR assays, utilizing frequent non-invasive liquid biopsies, enable the choice and subsequent modification of an effective targeted therapy for the treatment of an individual's cancer. Furthermore, after treatment, these assays may help monitor the patient to detect the presence of minimal residual disease, enabling further intervention if necessary.

To demonstrate the clinical potential of their approach, six DNA samples from the plasma of patients with cancer were assayed. The samples from five of the patients were blinded. Multiplex SuperSelective PCR assays were performed to confirm the presence or absence of particular human epidermal growth factor receptor (EGFR) mutations in each [sample](#). These mutations were chosen because their presence determines which targeted therapy is most likely to be effective against non-small-cell lung cancer. Four samples were positive for different EGFR target mutations and two samples did not give signals for EGFR target mutations. After the blinded samples were unblinded, it was revealed that the two negative samples were from patients who did not have EGFR mutations, and those samples were purposely provided as a control. The assays identified the previously known mutations in the other samples. Significantly, the [assay](#) also identified a very rare resistance mutation in one blinded sample that was not previously known to be present, suggesting that the patient's targeted therapy needed to be changed.

"The only difference between these assays and conventional multiplex [real-time](#) PCR is the substitution of SuperSelective primers for conventional primers," the investigators said. "These assays can be performed rapidly at points of care, and the results can immediately be used to inform subsequent clinical decisions."

The authors note that this technology might have other medical applications, such as the identification of rare antibiotic-resistant bacteria and the identification of rare mutant genes found in fetal cells circulating in a mother's plasma. Thus, the availability of sensitive multiplex SuperSelective PCR assays is likely to have wide clinical utility."

More information: Diana Y. Vargas et al, Multiplex SuperSelective PCR Assays for the Detection and Quantitation of Rare Somatic Mutations in Liquid Biopsies, *The Journal of Molecular Diagnostics* (2021). [DOI: 10.1016/j.jmoldx.2021.11.006](https://doi.org/10.1016/j.jmoldx.2021.11.006)

Provided by Elsevier

Citation: New assay shows promise for advancing personalized cancer treatment (2022, February 11) retrieved 4 May 2024 from <https://medicalxpress.com/news/2022-02-assay-advancing-personalized-cancer-treatment.html>

This document is subject to copyright. Apart from any fair dealing for the purpose of private study or research, no part may be reproduced without the written permission. The content is provided for information purposes only.
