

# Beyond the microscope: Identifying specific cancers using molecular analysis

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Researchers from Huntsman Cancer Institute (HCI) at the University of Utah report they have discovered a method to identify cancer-causing rearrangements of genetic material called chromosomal translocations quickly, accurately, and inexpensively. A description of the method and the research results appear online in this month's issue of the *EMBO Molecular Medicine* journal.

Many cancers result from chromosomal translocations in [tumor cells](#). Hundreds of cancer-causing translocations have been discovered, but current methods for detecting them have significant shortcomings.

The technique, developed in the lab of Stephen Lessnick, M.D., Ph.D., director of the Center for Children's [Cancer Research](#) at HCI, combines microarray technology, which can look for thousands of translocations in a single test, with a novel antibody that is used to detect the presence of the translocation.

"We're moving past the age when a pathologist looking through the microscope at a tumor sample is the best way to diagnose what type of cancer it is," said Lessnick. "The [molecular tests](#) currently available are slow, inefficient, and expensive, and one of the biggest issues is that you need high-quality tumor samples, not always available in the clinical setting, to do them." According to Lessnick, his method tolerates real-life specimens much better than the current standard techniques.

"Originally, this method was used in HCI's Cairns lab (headed by

Bradley R. Cairns, Ph.D.) to study RNA in yeast. We took their method and applied it to our study of chromosomal translocations in [human tissue](#)," Lessnick said. He said the next task is to find a commercial partner to develop this research from a 'proof of principle' into a [diagnostic test](#) that doctors can use to help their patients.

"With this method, there's potential to develop a single array that could test for every known cancer-causing translocation simultaneously. Currently, a clinician has to decide beforehand which specific cancer to test," he said.

The research used Ewing's sarcoma (a rare childhood cancer) as the case study for developing the method, but Lessnick maintains that the technology can be easily applied to any type of cancer caused by a translocation.

Funding for this project came from the National Institutes of Health's Innovative Molecular Analysis Technology program. The program focuses on rapid movement of new ideas from basic science labs (such as the Cairns lab) out into the clinical realm. "They were willing to fund this idea without a lot of preliminary data because it showed good potential to move toward clinical uses," said Lessnick.

Provided by University of Utah Health Sciences

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