

Researchers advance role of circulating tumor DNA to detect early melanoma growth

November 26 2018



Evidence of disease activity detected in circulating tumor DNA (ctDNA) predicts radiographic melanoma progression. Credit: Johns Hopkins Kimmel Cancer Center

Researchers at the Johns Hopkins Kimmel Cancer Center say they have added to evidence that measuring and monitoring tumor DNA that naturally circulates in the blood of melanoma patients can not only reliably help reveal the early stages of cancer growth and spread but also

uncover new treatment options that tumor genetic analysis alone may not.

"For some patients in our study, ctDNA (circulating tumor DNA) levels measured in a relatively simple blood test revealed tumor [mutations](#) that could be potentially targeted with current or new drugs that inhibit tumor growth mutations that are not revealed by genetic profiling of the tumor itself," says Evan Lipson, M.D., an associate professor of oncology at the Kimmel Cancer Center and a member of The Bloomberg~Kimmel Institute for Cancer Immunotherapy.

"For other patients, ctDNA levels accurately predicted [disease progression](#) as seen on CT scans, further demonstrating ctDNA's role as a blood-based biomarker of disease activity in melanoma patients."

Melanoma currently lacks a consistently and wholly reliable predictive blood-based biomarker of disease progression. Having one, Lipson says, would not only improve treatment outcomes but also reduce unnecessary or ineffective therapies.

For the study, published in the October 2018 issue of *Molecular Oncology*, the Kimmel Cancer Center scientists analyzed blood-based ctDNA from 119 advanced-melanoma Johns Hopkins patients. The patients were divided into three groups based on the type of tumor they had and potential mutations that were common and reoccurring in those tumors.

The first [group](#) of 60 people were patients with radiographically measurable metastatic melanoma, regardless of tumor mutation status. In this group, ctDNA testing revealed a targetable mutation in 38 of the 60 patients. In 33 of those patients, the mutations found using ctDNA matched the mutations found in tumor specimens. In two patients, ctDNA testing revealed a mutation that tumor testing had not.

In the second group, there were 29 patients with surgically removed high-risk (stage IIB-IV) melanoma whose tumor tissue revealed any of the seven common mutations. In this group, none of the patients whose melanoma tumors were surgically removed had evidence of disease before the study. However, five of 29 patients were discovered to have recurrent melanoma during the study, and in two of those cases, ctDNA was detected.

The third group had 30 patients who were receiving or had received therapy for advanced melanoma and had any of the seven common mutations. Of the 30, 17 experienced partial or complete response to therapy, which was confirmed using CT scans over 8, 14, 25 and 38 week periods, and no ctDNA was found in those patients after an initial CT scan evaluation. In the remaining 13 patients, ctDNA was detected during their treatment. In four of those 13, the disease was detected simultaneously by a CT scan and ctDNA results. In four others, ctDNA results predicted disease progression that was confirmed by using CT scan.

"When genetic testing of the tumor alone was used for some of the patients, it did not reveal any option for targeted therapy," Lipson says. "It turned out that when we looked in the bloodstream, lo and behold, we found ctDNA that uncovered options for therapy that provided benefit for patients and that otherwise were not going to be used."

Lipson notes that in general, melanoma patients receiving treatment typically have CT scans performed months apart over time to compare the growth or regression of the tumors. By using repeated ctDNA blood test results that reflect [tumor](#) activity in conjunction with those scans, the researchers say the biomarker was predictive of eventual disease progression seen on CT scans. These findings add to evidence that ctDNA testing may help radiologists and oncologists better interpret results of tests and treatments in patients with advanced melanoma.

According to the National Cancer Institute, there were 91,270 new cases of melanoma in 2018, making up more than 5 percent of all new cancer cases. There were 9,320 estimated deaths in 2018, and 91.8 percent of patients survive five years or longer.

Lipson said while larger trials will be needed to further investigate and confirm the findings, evidence is growing that using ctDNA can refine therapeutic outcomes and uncover additional avenues for therapy for some patients with melanoma.

"Our findings may serve as a blueprint for future, randomized investigations designed to further evaluate the clinical utility of incorporating ctDNA analysis among larger groups of [patients with melanoma](#)."

Provided by Johns Hopkins University School of Medicine

Citation: Researchers advance role of circulating tumor DNA to detect early melanoma growth (2018, November 26) retrieved 27 April 2024 from <https://medicalxpress.com/news/2018-11-advance-role-circulating-tumor-dna.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.
