

MicroRNA can help predict which breast cancer patients are more likely to see their cancer come back

November 4 2022



Credit: Pixabay/CC0 Public Domain

MicroRNA (miRNA) can be used as a biomarker to predict which patients are likely to face breast cancer recurrence and mortality,

according to study results published online ahead of print in the *Journal of the American College of Surgeons (JACS)*.

While long-term outcomes have improved for [patients](#) with [breast cancer](#), the most common cancer diagnosed in women, 20% to 30% of these patients will see their breast cancer come back. The process of identifying which patients are more likely to have a recurrence has been a challenge. Therefore, a research team in Galway, Republic of Ireland, set out to determine whether miRNAs—small, non-coding molecules that modulate [genetic expression](#) and affect cancer development—predict which breast cancer patients are more likely to have a recurrence of, and die from, breast cancer.

The researchers discovered that patients with an increased expression of a certain type of miRNA, miR-145, are unlikely to have a recurrence of breast cancer. In other words, miR-145 inhibited the development and progression of cancer, explained lead author Matthew G. Davey, MRCSI, Ph.D., discipline of surgery, University of Galway, Republic of Ireland.

"We showed that increased expression of this biomarker, which was measured in patients' [blood samples](#) during chemotherapy, actually predicted their long-term oncological outcome," Dr. Davey said. "We can predict those who are likely to suffer recurrence and those who will be free of recurrence."

The authors believe that miR-145 could help identify patients who could benefit from closer breast cancer surveillance and a tailored treatment strategy in the postoperative phase of treatment. Likewise, it could also identify patients who are at lower risk of recurrence and may not need systemic treatments, which often can have harmful side effects.

"This biomarker will help us give the right treatment to the right

patients," said study coauthor Michael J. Kerin, MCh, FRCSI, FRCS.Ed., FRCS Gen., chair of surgery, University of Galway.

Study details

This prospective, multicenter trial recruited 124 patients who were treated with standard-of-care [neoadjuvant chemotherapy](#) for localized breast cancer in eight independent treatment sites across Ireland. The type of chemotherapy varied according to the judgment of the patient's healthcare team.

Blood samples were collected from the patients over a three-year period (May 2011-April 2014) at two timepoints—at diagnosis and halfway through chemotherapy treatment. To establish their roles in predicting whether the patient would be free of recurrence or disease and their overall survival, miRNA expression levels were evaluated at each timepoint.

Key findings

The study found that increased miR-145 expression correlated with improved outcomes at almost nine years follow-up.

- Increased miR-145 independently predicted improved recurrence-free survival (HR: 0.00, 95% CI: 0.00—0.99, P=0.050) and trended towards improved disease-free survival (HR: 0.00, 95% CI: 0.00—1.42, P=0.067) when statistical analyses were performed.
- Increased miR-145 expression levels did not predict overall survival.

Future research opportunities

For the current study, the researchers looked at patients with any of the five subtypes of breast cancer, though the study was not designed or powered to determine how effective miR-145 was at predicting outcomes for any particular subtype of breast cancer. They plan further [clinical trials](#) to help answer that question.

There is also research underway to see if the expression of miRNA could be increased in [breast cancer patients](#), according to Professor Kerin. For instance, "one trial is testing miRNA replacement therapies in mice, but the research is in its early stages and it is unclear if the therapy could be transferred to humans," he said.

Dr. Davey said their group plans to conduct a similar trial to the one they've just published in *JACS* in which researchers focus on a specific subtype of [breast cancer](#) (HER2-positive patients).

Study coauthors are Andrew McGuire, Maire Caitlin Casey, Ronan M. Waldron, Maxwell Paganga, Emma Holian, John Newell, Helen M. Heneghan, Ailbhe M. McDermott, Aoife J. Lowery, and Nicola Miller, all of the University of Galway; and Maccon M. Keane of the Galway University Hospital.

More information: valuating the Role of Circulating MicroRNAs in Predicting Long-Term Survival Outcomes in Breast Cancer: A Prospective, Multicenter Clinical Trial, *Journal of the American College of Surgeons* (2022). [DOI: 10.1097/XCS.0000000000000465](https://doi.org/10.1097/XCS.0000000000000465)

Provided by American College of Surgeons

Citation: MicroRNA can help predict which breast cancer patients are more likely to see their cancer come back (2022, November 4) retrieved 6 May 2024 from

<https://medicalxpress.com/news/2022-11-micrna-breast-cancer-patients.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.